Package ‘plotly’

April 10, 2019

Title Create Interactive Web Graphics via 'plotly.js'
Version 4.9.0
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Description
Create interactive web graphics from 'ggplot2' graphs and/or a custom interface to the (MIT-licensed) JavaScript library 'plotly.js' inspired by the grammar of graphics.
https://plot.ly/r
BugReports https://github.com/ropensci/plotly/issues
Depends R (>= 3.2.0), ggplot2 (>= 3.0.0)
Imports tools, scales, htrr, jsonlite (>= 1.6), magrittr, digest,
  viridisLite, base64enc, htmltools (>= 0.3.6), htmlwidgets (>= 1.3), tidyr, hexbin, RColorBrewer, dplyr, tibble, lazyeval (>= 0.2.0), rlang, crosstalk, purrr, data.table, promises
Suggests MASS, maps, ggthemes, GGally, testthat, knitr, devtools, shiny (>= 1.1.0), shinystan (>= 1.3.0), curl, rmarkdown, vdiffr, Cairo, broom, webshot, listviewer, dendextend, sf, maptools, rgeos, png, IRdisplay, processx, plotlyGeoAssets, forcats
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Author Carson Sievert [aut, cre] (<https://orcid.org/0000-0002-4958-2844>),
  Chris Parmer [aut],
  Toby Hocking [aut],
  Scott Chamberlain [aut],
  Karthik Ram [aut],
  Marianne Corvellec [aut] (<https://orcid.org/0000-0002-1994-3581>),
  Pedro Despouy [aut],
  Plotly Technologies Inc. [cph]
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add_annotations

Add an annotation(s) to a plot

Description

Add an annotation(s) to a plot

Usage

add_annotations(p, text = NULL, ..., data = NULL, inherit = TRUE)

Arguments

p a plotly object
text annotation text (required).
... these arguments are documented at https://github.com/plotly/plotly.js/blob/master/src/components/annotations/attributes.js
add_data

Add data to a plotly visualization

Description

Add data to a plotly visualization

Usage

add_data(p, data = NULL)

Arguments

p a plotly visualization
data a data frame.

Examples

plot_ly() %>% add_data(economics) %>% add_trace(x = ~date, y = ~pce)

add_fun

Apply function to plot, without modifying data

Description

Useful when you need two or more layers that apply a summary statistic to the original data.

Usage

add_fun(p, fun, ...)

Arguments

p a plotly object.
fun a function. Should take a plotly object as input and return a modified plotly object.
... arguments passed to fun.
add_trace

Add trace(s) to a plotly visualization

Description

Add trace(s) to a plotly visualization

Usage

add_trace(p, ..., data = NULL, inherit = TRUE)

add_markers(p, x = NULL, y = NULL, z = NULL, ..., data = NULL, inherit = TRUE)

add_text(p, x = NULL, y = NULL, z = NULL, text = NULL, ..., data = NULL, inherit = TRUE)

add_paths(p, x = NULL, y = NULL, z = NULL, ..., data = NULL, inherit = TRUE)

add_lines(p, x = NULL, y = NULL, z = NULL, ..., data = NULL, inherit = TRUE)

add_segments(p, x = NULL, y = NULL, xend = NULL, yend = NULL, ..., data = NULL, inherit = TRUE)

add_polygons(p, x = NULL, y = NULL, ..., data = NULL, inherit = TRUE)

add_sf(p, ..., x = ~x, y = ~y, data = NULL, inherit = TRUE)

add_table(p, ..., rownames = TRUE, data = NULL, inherit = TRUE)

add_ribbons(p, x = NULL, ymin = NULL, ymax = NULL, ..., data = NULL, inherit = TRUE)

add_area(p, r = NULL, t = NULL, ..., data = NULL, inherit = TRUE)

add_pie(p, values = NULL, labels = NULL, ..., data = NULL, inherit = TRUE)

add_bars(p, x = NULL, y = NULL, ..., data = NULL, inherit = TRUE)

add_histogram(p, x = NULL, y = NULL, ..., data = NULL, inherit = TRUE)

add_histogram2d(p, x = NULL, y = NULL, z = NULL, ..., data = NULL,
add_trace

add_histogram2dcontour(p, x = NULL, y = NULL, z = NULL, ...
  data = NULL, inherit = TRUE)

add_heatmap(p, x = NULL, y = NULL, z = NULL, ...
  data = NULL, inherit = TRUE)

add_contour(p, z = NULL, ...
  data = NULL, inherit = TRUE)

add_boxplot(p, x = NULL, y = NULL, ...
  data = NULL, inherit = TRUE)

add_surface(p, z = NULL, ...
  data = NULL, inherit = TRUE)

add_mesh(p, x = NULL, y = NULL, z = NULL, ...
  data = NULL, inherit = TRUE)

add_scattergeo(p, ...)

add_choropleth(p, z = NULL, ...
  data = NULL, inherit = TRUE)

Arguments

p a plotly object

... Arguments (i.e., attributes) passed along to the trace type. See schema() for a
  list of acceptable attributes for a given trace type (by going to traces -> type ->
  attributes). Note that attributes provided at this level may override other argu-
  ments (e.g. plot_ly(x = 1:10, y = 1:10, color = I("red"), marker = list(color = "blue"))).

data A data frame (optional) or crosstalk::SharedData object.

inherit inherit attributes from plot_ly()? 

x the x variable.

y the y variable.

z a numeric matrix

text textual labels.

xend "final" x position (in this context, x represents "start")

yend "final" y position (in this context, y represents "start")

rownames whether or not to display the rownames of data.

ymin a variable used to define the lower boundary of a polygon.

ymax a variable used to define the upper boundary of a polygon.

r For polar chart only. Sets the radial coordinates.

t For polar chart only. Sets the radial coordinates.

values the value to associated with each slice of the pie.

labels the labels (categories) corresponding to values.
**add_trace**

**Author(s)**
Carson Sievert

**References**

- https://plotly-r.com/overview.html
- https://plotly/r

**See Also**

plot_ly()

**Examples**

```r
# the `plot_ly()` function initiates an object, and if no trace type
# is specified, it sets a sensible default
p <- plot_ly(economics, x = ~date, y = ~uempmed)
p

# some `add_*()` functions are a specific case of a trace type
# for example, `add_markers()` is a scatter trace with mode of markers
add_markers(p)

# scatter trace with mode of text
add_text(p, text = "%")

# scatter trace with mode of lines
add_paths(p)

# like `add_paths()`, but ensures points are connected according to `\x`
add_lines(p)

# if you prefer to work with plotly.js more directly, can always
# use `add_trace()` and specify the type yourself
add_trace(p, type = "scatter", mode = "markers+lines")

# mappings provided to `plot_ly()` are "global", but can be overwritten
plot_ly(economics, x = ~date, y = ~uempmed, color = I("red"), showlegend = FALSE) %>%
  add_lines() %>%
  add_markers(color = "pop")

# a number of `add_*()` functions are special cases of the scatter trace
plot_ly(economics, x = ~date) %>%
  add_ribbons(ymin = ~pce - 1e3, ymax = ~pce + 1e3)

# use `group_by()` (or `group2NA()`) to apply visual mapping
# once per group (e.g. one line per group)
txhousing %>%
  group_by(city) %>
```
```
plot_ly(x = ~date, y = ~median) %>%
  add_lines(color = "black")

# Not run:
# use `add_sf()` or `add_polygons()` to create geo-spatial maps
if (requireNamespace("sf", quietly = TRUE)) {
  nc <- sf::st_read(system.file("shape/nc.shp", package = "sf"), quiet = TRUE)
  plot_ly() %>% add_sf(data = nc)
}

# univariate summary statistics
plot_ly(mtcars, x = ~factor(vs), y = ~mpg) %>%
  add_boxplot()
plot_ly(mtcars, x = ~factor(vs), y = ~mpg) %>%
  add_trace(type = "violin")

# `add_histogram()` does binning for you...
mtcars %>%
  plot_ly(x = ~factor(vs)) %>%
  add_histogram()

# ...but you can 'pre-compute' bar heights in R
mtcars %>%
  dplyr::count(vs) %>%
  plot_ly(x = ~vs, y = ~n) %>%
  add_bars()

# the 2d analogy of add_histogram() is add_histogram2d()/add_histogram2dcontour()
library(MASS)
(p <- plot_ly(geyser, x = ~waiting, y = ~duration))
add_histogram2d(p)
add_histogram2dcontour(p)

# the 2d analogy of add_bars() is add_heatmap()/add_contour()
# (i.e., bin counts must be pre-specified)
den <- kde2d(geyser$waiting, geyser$duration)
(p <- plot_ly(x = den$x, y = den$y, z = den$z))
add_heatmap(p)
add_contour(p)

# `add_table()` makes it easy to map a data frame to the table trace type
plot_ly(economics) %>%
  add_table()

# pie charts!
ds <- data.frame(labels = c("A", "B", "C"), values = c(10, 40, 60))
plot_ly(ds, labels = ~labels, values = ~values) %>%
  add_pie() %>%
  layout(title = "Basic Pie Chart using Plotly")
data(wind)
plot_ly(wind, r = ~r, t = ~t) %>%
```
animation_opts

```r
add_area(color = "nms")
layout(radialaxis = list(ticksuffix = ", "), orientation = 270)
```

```
# 3D chart types
plot_ly(z = ~volcano)
add_surface()
plot_ly(x = c(0, 0, 1), y = c(0, 1, 0), z = c(0, 0, 0))
add_mesh()
```

## End

---

**animation_opts**

**Animation configuration options**

### Description

Animations can be created by either using the `frame` argument in `plot_ly()` or the (unofficial) `frame` `ggplot2` aesthetic in `ggplotly()`. By default, animations populate a play button and slider component for controlling the state of the animation (to pause an animation, click on a relevant location on the slider bar). Both the play button and slider component transition between frames according rules specified by `animation_opts()`.

### Usage

```r
animation_opts(p, frame = 500, transition = frame, easing = "linear",
               redraw = TRUE, mode = "immediate")
```

```r
animation_slider(p, hide = FALSE, ...)
```

```r
animation_button(p, ..., label)
```

### Arguments

- **p**: a `plotly` object.
- **frame**: The amount of time between frames (in milliseconds). Note that this amount should include the transition.
- **transition**: The duration of the smooth transition between frames (in milliseconds).
- **easing**: The type of transition easing. See the list of options here [https://github.com/plotly/plotly.js/blob/master/src/plots/animation_attributes.js](https://github.com/plotly/plotly.js/blob/master/src/plots/animation_attributes.js)
- **redraw**: Trigger a redraw of the plot at completion of the transition? A redraw may significantly impact performance, but may be necessary to update graphical elements that can’t be transitioned.
mode

Describes how a new animate call interacts with currently-running animations. If immediate, current animations are interrupted and the new animation is started. If next, the current frame is allowed to complete, after which the new animation is started. If afterall all existing frames are animated to completion before the new animation is started.

hide

remove the animation slider?

... for animation_slider, attributes are passed to a special layout.sliders object tied to the animation frames. The definition of these attributes may be found here https://github.com/plotly/plotly.js/blob/master/src/components/sliders/attributes.js For animation_button, arguments are passed to a special layout.updatemenus button object tied to the animation https://github.com/plotly/plotly.js/blob/master/src/components/updatemenus/attributes.js

label

a character string used for the animation button’s label

Author(s)

Carson Sievert

Examples

def <- data.frame(
    x = c(1, 2, 2, 1, 1, 2),
    y = c(1, 2, 2, 1, 1, 2),
    z = c(1, 1, 2, 2, 3, 3)
)
plot_ly(df) %>%
    add_markers(x = 1.5, y = 1.5) %>%
    add_markers(x = ~x, y = ~y, frame = ~z)

# it's a good idea to remove smooth transitions when there is
# no relationship between objects in each view
plot_ly(mtcars, x = ~wt, y = ~mpg, frame = ~cyl) %>%
    animation_opts(transition = 0)

# works the same way with ggplotly
if (interactive()) {
    p <- ggplot(txhousing, aes(month, median)) +
        geom_line(aes(group = year), alpha = 0.3) +
        geom_smooth() +
        geom_line(aes(frame = year, ids = month), color = "red") +
        facet_wrap(~ city)

    ggplotly(p, width = 1200, height = 900) %>%
        animation_opts(1000)
}

# for more, see https://cpsievert.github.io/plotly_book/key-frame-animations.html
Description

Convenience functions for working with version 2 of plotly’s REST API. Upload R objects to a plotly account via `api_create()` and download plotly objects via `api_download_plot()`/`api_download_grid()`. For anything else, use `api()`.

Usage

```r
api_create(x = last_plot(), filename = NULL, fileopt = c("overwrite", "new"), sharing = c("public", "private", "secret"), ...)

## S3 method for class 'plotly'
api_create(x = last_plot(), filename = NULL,
    fileopt = "overwrite", sharing = "public", ...)

## S3 method for class 'ggplot'
api_create(x = last_plot(), filename = NULL,
    fileopt = "overwrite", sharing = "public", ...)

## S3 method for class 'data.frame'
api_create(x, filename = NULL,
    fileopt = "overwrite", sharing = "public", ...)

api_download_plot(id, username)

api_download_grid(id, username)

api(endpoint = "/", verb = "GET", body = NULL, ...)
```

Arguments

- `x` An R object to hosted on plotly's web platform. Can be a plotly/ggplot2 object or a `data.frame`.
- `filename` character vector naming file(s). If `x` is a plot, can be a vector of length 2 naming both the plot AND the underlying grid.
- `fileopt` character string describing whether to "overwrite" existing files or ensure "new" file(s) are always created.
- `sharing` If 'public', anyone can view this graph. It will appear in your profile and can appear in search engines. You do not need to be logged in to Plotly to view this chart. If 'private', only you can view this plot. It will not appear in the Plotly feed, your profile, or search engines. You must be logged in to Plotly to view this graph. You can privately share this graph with other Plotly users in your online Plotly account and they will need to be logged in to view this plot. If
'secret', anyone with this secret link can view this chart. It will not appear in the Plotly feed, your profile, or search engines. If it is embedded inside a webpage or an IPython notebook, anybody who is viewing that page will be able to view the graph. You do not need to be logged in to view this plot.

... For api(), these arguments are passed onto \texttt{httr::VERB()}. For \texttt{api_create()}, these arguments are included in the body of the HTTP request.

- \textbf{id} a filename id.
- \textbf{username} a plotly username.
- \textbf{endpoint} the endpoint (i.e., location) for the request. To see a list of all available endpoints, call api(). Any relevant query parameters should be included here (see examples).
- \textbf{verb} name of the HTTP verb to use (as in, \texttt{httr::VERB()}).
- \textbf{body} body of the HTTP request (as in, \texttt{httr::VERB()}). If this value is not already converted to JSON (via \texttt{jsonlite::toJSON()}), it uses the internal \texttt{to\_JSON()} to ensure values are "automatically unboxed" (i.e., vec.

\textbf{Author(s)}

Carson Sievert

\textbf{References}

\url{https://api.plot.ly/v2}

\textbf{See Also}

\texttt{signup()}

\textbf{Examples}

```
## Not run:

# api_create() makes it easy to upload ggplot2/plotly objects
# and/or data frames to your plotly account
# -----------------------------------------------

# A data frame creates a plotly "grid". Printing one will take you
# to the it's web address so you can start creating!
(m <- api_create(mtcars))

# A plotly/ggplot2 object create a plotly "plot".
p <- plot_ly(mtcars, x = ~factor(vs))
(r <- api_create(p))

# api_create() returns metadata about the remote "file". Here is
# one way you could use that metadata to download a plot for local use:
fileID <- strsplit(r$file$fid, ":")[[1]]
```
as.widget

```r
layout(
  api_download_plot(fileID[2], fileID[1]),
  title = sprintf("Local version of <a href='%s'>this</a> plot", r$file$web_url)
)
```

# The api() function provides a low-level interface for performing
# any action at any endpoint! It always returns a list.
# ---------------------------------------------------------------------

# list all the endpoints
api()

# search the entire platform!
# see https://api.plot.ly/v2/search
api("search?q=overdose")
api("search?q=plottype:pie trump fake")

# these examples will require a user account
usr <- Sys.getenv("plotly_username", NA)
if (!is.na(usr)) {
  # your account info https://api.plot.ly/v2/#users
  api(sprintf("users/%s", usr))
  # your folders/files https://api.plot.ly/v2/folders#user
  api(sprintf("folders/home?user=%s", usr))
}

# Retrieve a specific file https://api.plot.ly/v2/files#retrieve
api("files/cpsievert:14681")

# change the filename https://api.plot.ly/v2/files#update
# (note: this won't work unless you have proper credentials to the relevant account)
api("files/cpsievert:14681", "PATCH", list(filename = "toy file"))

# Copy a file https://api.plot.ly/v2/files#lookup
api("files/cpsievert:14681/copy", "POST")

# Create a folder https://api.plot.ly/v2/folders#create
api("folders", "POST", list(path = "/starts/at/root/and/ends/here"))

## End(Not run)

### as.widget

**Convert a plotly object to an htmlwidget object**

**Description**

This function was deprecated in 4.0.0, as plotly objects are now htmlwidget objects, so there is no need to convert them.
Usage

as.widget(x, ...)

Arguments

x  a plotly object.
...

other options passed onto htmlwidgets::createWidget

Description

Convert a list to a plotly htmlwidget object

Usage

as.widget(x, ...)

Arguments

x  a plotly object.
...

other options passed onto htmlwidgets::createWidget

Examples

trace <- list(x = 1, y = 1)
obj <- list(data = list(trace), layout = list(title = "my plot"))
as_widget(obj)

attrs_selected  Specify attributes of selection traces

Description

By default the name of the selection trace derives from the selected values.

Usage

attrs_selected(opacity = 1, ...)

bbox

**Arguments**

- `opacity` a number between 0 and 1 specifying the overall opacity of the selected trace
- ... 
  
  other trace attributes attached to the selection trace.

**Author(s)**

Carson Sievert

---

bbox 

*Estimate bounding box of a rotated string*

---

**Description**

Estimate bounding box of a rotated string

**Usage**

bbox(txt = "foo", angle = 0, size = 12)

**Arguments**

- `txt` a character string of length 1
- `angle` sets the angle of the tick labels with respect to the horizontal (e.g., `tickangle` of -90 draws the tick labels vertically)
- `size` vertical size of a character

**References**

https://www.dropbox.com/s/nc6968prgw8ne4w/bbox.pdf?dl=0

---

colorbar 

*Modify the colorbar*

---

**Description**

Modify the colorbar

**Usage**

colorbar(p, ..., limits = NULL, which = 1)
Arguments

p
a plotly object

... arguments are documented here https://plot.ly/r/reference/#scatter-marker-colorbar.

limits numeric vector of length 2. Set the extent of the colorbar scale.

which colorbar to modify? Should only be relevant for subplots with multiple colorbars.

Author(s)

Carson Sievert

Examples

p <- plot_ly(mtcars, x = ~wt, y = ~mpg, color = ~cyl)

# pass any colorbar attribute --
# https://plot.ly/r/reference/#scatter-marker-colorbar
colorbar(p, len = 0.5)

# Expand the limits of the colorbar
colorbar(p, limits = c(0, 20))
# values outside the colorbar limits are considered "missing"
colorbar(p, limits = c(5, 6))

# also works on colorbars generated via a z value
corr <- cor(diamonds[vapply(diamonds, is.numeric, logical(1))])
plot_ly(x = rownames(corr), y = colnames(corr), z = corr) %>%
  add_heatmap() %>%
  colorbar(limits = c(-1, 1))

config Set the default configuration for plotly

Description

Set the default configuration for plotly

Usage

config(p, ..., cloud = FALSE, showSendToCloud = cloud, locale = NULL,
  mathjax = NULL)
**Arguments**

- `p` argument: a plotly object.
  These arguments are documented at [https://github.com/plotly/plotly.js/blob/master/src/plot_api/plot_config.js](https://github.com/plotly/plotly.js/blob/master/src/plot_api/plot_config.js)

- `cloud` argument: deprecated. Use `showSendToCloud` instead.

- `showSendToCloud` argument: include the send data to cloud button?

- `locale` argument: locale to use. See [here](#) for more info.

- `mathjax` argument: add MathJax rendering support. If "cdn", mathjax is loaded externally (meaning an internet connection is needed for TeX rendering). If "local", the PLOTLY_MATHJAX_PATH environment variable must be set to the location (a local file path) of MathJax. IMPORTANT: `plotly` uses SVG-based mathjax rendering which doesn’t play nicely with HTML-based rendering (e.g., `rmarkdown` documents and `shiny` apps). To leverage both types of rendering, you must `<iframe>` your plotly graph(s) into the larger document (see [here](#) for an `rmarkdown` example and [here](#) for a `shiny` example).

**Author(s)**

Carson Sievert

**Examples**

```r
# remove the plotly logo and collaborate button from modebar
cfg <- plot_ly()$displaylogo = FALSE, collaborate = FALSE

# enable mathjax
# see more examples at https://plot.ly/r/latex/
plot_ly(x = c(1, 2, 3, 4), y = c(1, 4, 9, 16)) %>%
  layout(title = TeX("\text{Some mathjax: }\alpha+\beta x\text{ }")) %>%
  config(mathjax = "cdn")

# change the language used to render date axes and on-graph text
# (e.g., modebar buttons)
today <- Sys.Date()
x <- seq.Date(today, today + 360, by = "day")
p <- plot_ly(x = x, y = rnorm(length(x))) %>%
  add_lines()

# japanese
config(p, locale = "ja")
# german
config(p, locale = "de")
# spanish
config(p, locale = "es")
# chinese
config(p, locale = "zh-CN")
```
### embed_notebook

**Embed a plot as an iframe into a Jupyter Notebook**

**Description**

Embed a plot as an iframe into a Jupyter Notebook

**Usage**

```r
embed_notebook(x, width = NULL, height = NULL, file = NULL)
```

**Arguments**

- `x`: a plotly object
- `width`: attribute of the iframe. If NULL, the width in `plot_ly` is used. If that is also NULL, '100%' is the default.
- `height`: attribute of the iframe. If NULL, the height in `plot_ly` is used. If that is also NULL, '400px' is the default.
- `file`: deprecated.

**Author(s)**

Carson Sievert

### event_data

**Access plotly user input event data in shiny**

**Description**

This function must be called within a reactive shiny context.

**Usage**

```r
event_data(event = c("plotly_hover", "plotly_unhover", "plotly_click", 
  "plotly_doubleclick", "plotly_selected", "plotly_selecting", 
  "plotly_brushed", "plotly_brushing", "plotly_deselect", 
  "plotly_relayout", "plotly_restyle", "plotly_legendclick", 
  "plotly_legendedoubleclick", "plotly_clickannotation", 
  "plotly_afterplot"), source = "A", 
  session = shiny::getDefaultReactiveDomain(), priority = c("input", 
  "event"))
```
**event_register**

**Arguments**

- **event**: The type of plotly event. All supported events are listed in the function signature above (i.e., the usage section).
- **source**: a character string of length 1. Match the value of this string with the `source` argument in `plot_ly()` (or `ggplotly()`) to respond to events emitted from that specific plot.
- **session**: a shiny session object (the default should almost always be used).
- **priority**: the priority of the corresponding shiny input value. If equal to "event", then `event_data()` always triggers re-execution, instead of re-executing only when the relevant shiny input value changes (the default).

**Author(s)**

Carson Sievert

**References**


**See Also**

`event_register`, `event_unregister`

**Examples**

```r
## Not run:
plotly_example("shiny", "event_data")

## End(Not run)
```

---

**Description**

Register a shiny input value

**Usage**

```r
event_register(p, event = NULL)
```
Author(s)
Carson Sievert

See Also
event_data

---

**event_unregister**  
*Un-register a shiny input value*

**Description**
Un-register a shiny input value

**Usage**
```
event_unregister(p, event = NULL)
```

**Arguments**
- `p`: a plotly object.
- `event`: The type of plotly event. All supported events are listed in the function signature above (i.e., the usage section).

Author(s)
Carson Sievert

See Also
event_data

---

**export**  
*Export a plotly graph to a static file*

**Description**
This function is in the process of being deprecated (use orca instead).

**Usage**
```
export(p = last_plot(), file = "plotly.png", selenium = NULL, ...)
```
Arguments

- **p**: a plotly or ggplot object.
- **file**: a filename. The file type is inferred from the file extension. Valid extensions include 'jpeg' | 'png' | 'webp' | 'svg' | 'pdf'
- **selenium**: used only when `p` is a WebGL plot or the output format is 'webp' or 'svg'. Should be an object of class 'rsClientServer' returned by `RSelenium::rsDriver`.
- **...**: if `p` is non-WebGL and the output file format is jpeg/png/pdf arguments are passed along to `webshot::webshot()`. Otherwise, they are ignored.

Details

For SVG plots, a screenshot is taken via `webshot::webshot()`. Since phantomjs (and hence `webshot`) does not support WebGL, the RSelenium package is used for exporting WebGL plots.

Author(s)

Carson Sievert

---

### geom2trace

**Convert a "basic" geoms to a plotly.js trace.**

---

Description

This function makes it possible to convert ggplot2 geoms that are not included with ggplot2 itself. Users shouldn't need to use this function. It exists purely to allow other package authors to write their own conversion method(s).

Usage

`geom2trace(data, params, p)`

Arguments

- **data**: the data returned by `plotly::to_basic`.
- **params**: parameters for the geom, statistic, and 'constant' aesthetics
- **p**: a ggplot2 object (the conversion may depend on scales, for instance).
get_figure  
*Request a figure object*

**Description**

Deprecated: see `api_download_plot()`.

**Usage**

```r
get_figure(username, id)
```

**Arguments**

- **username**: corresponding username for the figure.
- **id**: of the Plotly figure.

---

gg2list  
*Convert a ggplot to a list.*

**Description**

Convert a ggplot to a list.

**Usage**

```r
gg2list(p, width = NULL, height = NULL, tooltip = "all",
dynamicTicks = FALSE, layerData = 1, originalData = TRUE,
source = "A", ...)
```

**Arguments**

- **p**: `ggplot2` plot.
- **width**: Width of the plot in pixels (optional, defaults to automatic sizing).
- **height**: Height of the plot in pixels (optional, defaults to automatic sizing).
- **tooltip**: a character vector specifying which aesthetic tooltips to show in the tooltip. The default, "all", means show all the aesthetic tooltips (including the unofficial "text" aesthetic).
- **dynamicTicks**: accepts the following values: FALSE, TRUE, "x", or "y". Dynamic ticks are useful for updating ticks in response to zoom/pan/filter interactions; however, there is no guarantee they reproduce axis tick text as they would appear in the static ggplot2 image.
- **layerData**: data from which layer should be returned?
- **originalData**: should the "original" or "scaled" data be returned?
source

a character string of length 1. Match the value of this string with the source argument in `event_data()` to retrieve the event data corresponding to a specific plot (shiny apps can have multiple plots).

... currently not used

Value

a 'built' plotly object (list with names "data" and "layout").

---

**ggplotly**

**Convert ggplot to plotly**

**Description**

This function converts a `ggplot2::ggplot()` object to a plotly object.

**Usage**

```r
ggplotly(p = ggplot2::last_plot(), width = NULL, height = NULL,
         tooltip = "all", dynamicTicks = FALSE, layerData = 1,
         originalData = TRUE, source = "A", ...)
```

**Arguments**

- `p` a ggplot object.
- `width` Width of the plot in pixels (optional, defaults to automatic sizing).
- `height` Height of the plot in pixels (optional, defaults to automatic sizing).
- `tooltip` a character vector specifying which aesthetic mappings to show in the tooltip. The default, "all", means show all the aesthetic mappings (including the unofficial "text" aesthetic). The order of variables here will also control the order they appear. For example, use tooltip = c("y", "x", "colour") if you want y first, x second, and colour last.
- `dynamicTicks` should `plotly.js` dynamically generate axis tick labels? Dynamic ticks are useful for updating ticks in response to zoom/pan interactions; however, they can not always reproduce labels as they would appear in the static ggplot2 image.
- `layerData` data from which layer should be returned?
- `originalData` should the "original" or "scaled" data be returned?
- `source` a character string of length 1. Match the value of this string with the source argument in `event_data()` to retrieve the event data corresponding to a specific plot (shiny apps can have multiple plots).
- `...` arguments passed onto methods.
Details

Conversion of relative sizes depends on the size of the current graphics device (if no device is open, width/height of a new (off-screen) device defaults to 640/480). In other words, height and width must be specified at runtime to ensure sizing is correct. For examples on how to specify the output container’s height/width in a shiny app, see plotly_example("shiny", "ggplotly_sizing").

Author(s)

Carson Sievert

References

https://plot.ly/ggplot2

See Also

plot_ly()

Examples

```r
## Not run:
# simple example
ggiris <- qplot(Petal.Width, Sepal.Length, data = iris, color = Species) # simple example
ggplotly(ggiris)

data(canada.cities, package = "maps")
viz <- ggplot(canada.cities, aes(long, lat)) +
  borders(regions = "canada") +
  coord_equal() +
  geom_point(aes(text = name, size = pop), colour = "red", alpha = 1/2)
ggplotly(viz, tooltip = c("text", "size"))

# linked scatterplot brushing
d <- highlight_key(mtcars)
qplot(data = d, x = mpg, y = wt) # simple example
  subplot(qplot(data = d, x = mpg, y = vs)) # simple example
  layout(title = "Click and drag to select points") # simple example
  highlight("plotly_selected")

# more brushing (i.e. highlighting) examples
demo("crosstalk-highlight-ggplotly", package = "plotly")

# client-side linked brushing in a scatterplot matrix
highlight_key(iris) # simple example
  GGally::ggpairs(aes(colour = Species), columns = 1:4) # simple example
  ggplotly(tooltip = c("x", "y", "colour")) # simple example
  highlight("plotly_selected")

## End(Not run)
```
group2NA

Separate groups with missing values

Description

This function is used internally by plotly, but may also be useful to some power users. The details section explains when and why this function is useful.

Usage

group2NA(data, groupNames = "group", nested = NULL, ordered = NULL, retrace.first = inherits(data, "GeomPolygon"))

Arguments

data: a data frame.
groupNames: character vector of grouping variable(s)
nested: other variables that group should be nested (i.e., ordered) within.
ordered: a variable to arrange by (within nested & groupNames). This is useful primarily for ordering by y
retrace.first: should the first row of each group be appended to the last row? This is useful for enclosing polygons with lines.

Details

If a group of scatter traces share the same non-positional characteristics (i.e., color, fill, etc), it is more efficient to draw them as a single trace with missing values that separate the groups (instead of multiple traces). In this case, one should also take care to make sure connectgaps is set to FALSE.

Value

a data.frame with rows ordered by: nested, then groupNames, then ordered. As long as groupNames contains valid variable names, new rows will also be inserted to separate the groups.

Examples

# note the insertion of new rows with missing values
group2NA(mtcars, "vs", "cyl")

# need to group lines by city somehow!
plot_ly(txhousing, x = ~date, y = ~median) %>% add_lines()

# instead of using group_by(), you could use group2NA()
tx <- group2NA(txhousing, "city")
plot_ly(tx, x = ~date, y = ~median) %>% add_lines()
# add_lines() will ensure paths are sorted by x, but this is equivalent
tx <- group2NA(txhousing, "city", ordered = "date")
plot_ly(tx, x = ~date, y = ~median) %>% add_paths()

---

hide_colorbar  
Hide color bar(s)

Description
Hide color bar(s)

Usage
hide_colorbar(p)

Arguments
p  
a plotly object.

See Also
hide_legend()

Examples

p <- plot_ly(mtcars, x = ~wt, y = ~cyl, color = ~cyl)
hide_colorbar(p)

---

hide_guides  
Hide guides (legends and colorbars)

Description
Hide guides (legends and colorbars)

Usage
hide_guides(p)

Arguments
p  
a plotly object.

See Also
hide_legend(), hide_colorbar()
**hide_legend**

**Description**
Hide legend

**Usage**
hide_legend(p)

**Arguments**

- `p` a plotly object.

**See Also**
hide_colorbar()

**Examples**

```r
p <- plot_ly(mtcars, x = ~wt, y = ~cyl, color = ~factor(cyl))
hide_legend(p)
```

**highlight**

**Query graphical elements in multiple linked views**

**Description**

This function sets a variety of options for brushing (i.e., highlighting) multiple plots. These options are primarily designed for linking multiple plotly graphs, and may not behave as expected when linking plotly to another htmlwidget package via crosstalk. In some cases, other htmlwidgets will respect these options, such as persistent selection in leaflet (see demo("highlight-leaflet", package = "plotly")).

**Usage**

```r
highlight(p, on = "plotly_click", off, persistent =getOption("persistent", FALSE), dynamic = FALSE, color = NULL, selectize = FALSE, defaultValues = NULL, opacityDim =getOption("opacityDim", .2), selected = attrs_selected(), debounce = 0, ...)
```
Arguments

on

- plotly_click
- plotly_hover
- plotly_selected: triggered through rectangular (layout.dragmode = 'select') or lasso (layout.dragmode = 'lasso') brush.

off

- plotly_doubleclick: triggered on a double mouse click while (layout.dragmode = 'zoom') or (layout.dragmode = 'pan')
- plotly_deselect: triggered on a double mouse click while (layout.dragmode = 'select') or (layout.dragmode = 'lasso')
- plotly_relayout: triggered whenever axes are rescaled (i.e., clicking the home button in the modebar) or whenever the height/width of the plot changes.

dynamic

should a widget for changing selection colors be included?

color

character string of color(s) to use for highlighting selections. See toRGB() for valid color specifications. If NULL (the default), the color of selected marks are not altered.

selectize

provide a selectize.js widget for selecting keys? Note that the label used for this widget derives from the groupName of the SharedData object.

defaultValues

a vector of values for setting a "default selection". These values should match the key attribute.

opacityDim

a number between 0 and 1 used to reduce the opacity of non-selected traces (by multiplying with the existing opacity).

selected

attributes of the selection, see attrs_selected().

debounce

amount of time to wait before firing an event (in milliseconds). The default of 0 means do not debounce at all. Debouncing is mainly useful when on = "plotly_hover" to avoid firing too many events when users clickly move the mouse over relevant graphical marks.

Author(s)

Carson Sievert

References

highlight_key

See Also
- attrs_selected()

Examples

# These examples are designed to show you how to highlight/brush a *single*
# view. For examples of multiple linked views, see `demo(package = "plotly")`

d <- highlight_key(txhousing, ~city)
p <- ggplot(d, aes(date, median, group = city)) + geom_line()
gg <- ggplotly(p, tooltip = "city")
highlight(gg, dynamic = TRUE)

# supply custom colors to the brush
cols <- toRGB(RColorBrewer::brewer_pal(3, "Dark2"), 0.5)
highlight(gg, on = "plotly_hover", color = cols, dynamic = TRUE)

# Use attrs_selected() for complete control over the selection appearance
# note any relevant colors you specify here should override the color argument
s <- attrs_selected(
  showlegend = TRUE,
  mode = "lines+markers",
  marker = list(symbol = "x")
)

highlight(layout(gg, showlegend = TRUE), selected = s)

highlight_key

Highlight/query data based on primary key

Description

This function simply creates an object of class crosstalk::SharedData. The reason it exists is to
make it easier to teach others how to leverage it’s functionality in plotly. It also makes it more
discoverable if one is already aware of highlight.

Usage

highlight_key(…)

Arguments

... arguments passed to crosstalk::SharedData$new()

Value

An object of class crosstalk::SharedData
**Author(s)**

Carson Sievert

**See Also**

highlight

---

**hobbs**

*Hobbs data*

---

**Description**

Description TBD.

**Usage**

hobbs

**Format**

A data frame with three variables: r, t, nms.

---

**knit_print.api_grid**

*Embed a plotly grid as an iframe in a knitr doc*

**Description**

Embed a plotly grid as an iframe in a knitr doc

**Usage**

`knit_print.api_grid(x, options, ...)`

**Arguments**

- `x` a plotly figure object
- `options` knitr options.
- `...` placeholder.

**References**

knit_print.api_grid_local

Embed a plotly grid as an iframe in a knitr doc

Description

Embed a plotly grid as an iframe in a knitr doc

Usage

knit_print.api_grid_local(x, options, ...)

Arguments

x  a plotly figure object
options  knitr options.
...  placeholder.

References


knit_print.api_plot

Embed a plotly figure as an iframe in a knitr doc

Description

Embed a plotly figure as an iframe in a knitr doc

Usage

knit_print.api_plot(x, options, ...)

Arguments

x  a plotly figure object
options  knitr options.
...  placeholder.

References

last_plot

*Retrieve the last plot to be modified or created.*

**Description**

Retrieve the last plot to be modified or created.

**Usage**

`last_plot()`

**See Also**

`ggplot2::last_plot()`

layout

*Modify the layout of a plotly visualization*

**Description**

Modify the layout of a plotly visualization

**Usage**

`layout(p, ..., data = NULL)`

**Arguments**

- `p` A plotly object.
- `...` Arguments to the layout object. For documentation, see [https://plot.ly/r/reference/#Layout_and_layout_style_objects](https://plot.ly/r/reference/#Layout_and_layout_style_objects)
- `data` A data frame to associate with this layout (optional). If not provided, arguments are evaluated using the data frame in `plot_ly()`.

**Author(s)**

Carson Sievert
mic

Mic data

Description
Description TBD.

Usage
mic

Format
A data frame with three variables: r, t, nms.

offline
Plotly Offline

Description
Deprecated in version 2.0 (offline plots are now the default)

Usage
offline(p, height, width, out_dir, open_browser)

Arguments
p a plotly object
height A valid CSS unit. (like "100%", "600px", "auto") or a number, which will be coerced to a string and have "px" appended.
width A valid CSS unit. (like "100%", "600px", "auto") or a number, which will be coerced to a string and have "px" appended.
out_dir a directory to place the visualization. If NULL, a temporary directory is used when the offline object is printed.
open_browser open the visualization after creating it?

Value
a plotly object of class "offline"

Author(s)
Carson Sievert
Static image exporting

Description

Export plotly objects to static images (e.g., pdf, png, jpeg, svg, etc) via the orca command-line utility.

Usage

orca(p, file = "plot.png", format = tools::file_ext(file),
    scale = NULL, width = NULL, height = NULL, mathjax = FALSE,
    parallel_limit = NULL, verbose = FALSE, debug = FALSE,
    safe = FALSE, more_args = NULL, ...)

orca_serve(port = 5151, mathjax = FALSE, safe = FALSE,
    request_limit = NULL, keep_alive = TRUE, window_max_number = NULL,
    quiet = FALSE, debug = FALSE, more_args = NULL, ...)

Arguments

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>p</code></td>
<td>a plotly object.</td>
</tr>
<tr>
<td><code>file</code></td>
<td>output filename.</td>
</tr>
<tr>
<td><code>format</code></td>
<td>the output format (png, jpeg, webp, svg, pdf, eps).</td>
</tr>
<tr>
<td><code>scale</code></td>
<td>Sets the image scale. Applies to all output images.</td>
</tr>
<tr>
<td><code>width</code></td>
<td>Sets the image width. If not set, defaults to <code>layout.width</code> value. Applies to all output images.</td>
</tr>
<tr>
<td><code>height</code></td>
<td>Sets the image height. If not set, defaults to <code>layout.height</code> value. Applies to all output images.</td>
</tr>
<tr>
<td><code>mathjax</code></td>
<td>whether or not to include MathJax (required to render TeX). If <code>TRUE</code>, the PLOTLY_MATHJAX_PATH environment variable must be set and point to the location of MathJax (this variable is also used to render TeX in interactive graphs, see <code>config</code>).</td>
</tr>
<tr>
<td><code>parallel_limit</code></td>
<td>Sets the limit of parallel tasks run.</td>
</tr>
<tr>
<td><code>verbose</code></td>
<td>Turn on verbose logging on stdout.</td>
</tr>
<tr>
<td><code>debug</code></td>
<td>Starts app in debug mode and turn on verbose logs on stdout.</td>
</tr>
<tr>
<td><code>safe</code></td>
<td>Turns on safe mode: where figures likely to make browser window hang during image generating are skipped.</td>
</tr>
<tr>
<td><code>more_args</code></td>
<td>additional arguments to pass along to system command. This is useful for specifying display and/or electron options, such as <code>--enable-webgl</code> or <code>--disable-gpu</code>.</td>
</tr>
<tr>
<td>...</td>
<td>for <code>orca()</code>, additional arguments passed along to <code>processx::run</code>. For <code>orca_serve()</code>, additional arguments passed along to <code>processx::process</code>.</td>
</tr>
<tr>
<td><code>port</code></td>
<td>Sets the server’s port number.</td>
</tr>
<tr>
<td><code>request_limit</code></td>
<td>Sets a request limit that makes orca exit when reached.</td>
</tr>
</tbody>
</table>
keep_alive  
Turn on keep alive mode where orca will (try to) relaunch server if process unexpectedly exits.

window_max_number  
Sets maximum number of browser windows the server can keep open at a given time.

quiet  
Suppress all logging info.

Details

The `orca()` function is designed for exporting one plotly graph whereas `orca_serve()` is meant for exporting many graphs at once. The former starts and stops an external (nodejs) process every time it is called whereas the latter starts up a process when called, then returns an export() method for exporting graphs as well as a close() method for stopping the external (background) process.

Methods

The `orca_serve()` function returns an object with two methods:

```r
export(p, file = "plot.png", format = tools::file_ext(file), scale = NULL, width = NULL, height = NULL)
```

Export a static image of a plotly graph. Arguments found here are the same as those found in `orca()`

```r
close()  
Close down the orca server and kill the underlying node process.
```

Fields

The `orca_serve()` function returns an object with two fields:

```r
port  
The port number that the server is listening to.
process  
An R6 class for controlling and querying the underlying node process.
```

Author(s)

Carson Sievert

Examples

```r
## Not run:
# NOTE: in a headless environment, you may need to set `more_args="--enable-webgl"`
# to export webgl correctly
p <- plot_ly(z = ~volcano) %>% add_surface()
orca(p, "surface-plot.svg")

#' # launch the server
server <- orca_serve()

# export as many graphs as you'd like
server$export(qplot(1:10), "test1.pdf")
server$export(plot_ly(x = 1:10, y = 1:10), "test2.pdf")
```
partial_bundle

# the underlying process is exposed as a field, so you
# have full control over the external process
server$process$is_alive()

# convenience method for closing down the server
server$close()

# remove the exported files from disk
unlink("test1.pdf")
unlink("test2.pdf")

## End (Not run)

### partial_bundle

**Use a partial bundle of plotly.js**

**Description**

Leveraging plotly.js' partial bundles can lead to smaller file sizes and faster rendering. The full list of available bundles, and the trace types that they support, are available [here](#).

**Usage**

```
partial_bundle(p, type = "auto", local = TRUE, minified = TRUE)
```

**Arguments**

- **p**: a plotly object.
- **type**: name of the (partial) bundle. The default, 'auto', attempts to find the smallest single bundle that can render `p`. If no single partial bundle can render `p`, then the full bundle is used.
- **local**: whether or not to download the partial bundle so that it can be viewed later without an internet connection.
- **minified**: whether or not to use a minified js file (non-minified file can be useful for debugging plotly.js)

**Details**

WARNING: use this function with caution when rendering multiple plotly graphs on a single website. That's because, if multiple plotly.js bundles are used, the most recent bundle will override the other bundles. See the examples section for an example.

**Author(s)**

Carson Sievert
plotly-shiny
Shiny bindings for plotly

Description

Output and render functions for using plotly within Shiny applications and interactive Rmd documents.
Usage

plotlyOutput(outputId, width = "100\%", height = "400px", inline = FALSE)

renderPlotly(expr, env = parent.frame(), quoted = FALSE)

Arguments

outputId: output variable to read from
width, height: Must be a valid CSS unit (like "100\%", "400px", "auto") or a number, which will be coerced to a string and have "px" appended. Note that, for height, using "auto" or "100 height is computed with HTML/CSS.
inline: use an inline (span()) or block container (div()) for the output
expr: An expression that generates a plotly
env: The environment in which to evaluate expr.
quoted: Is expr a quoted expression (with quote())? This is useful if you want to save an expression in a variable.

Description

Modify a plotly object inside a shiny app

Usage

plotlyProxy(outputId, session = shiny::getDefaultReactiveDomain(), deferUntilFlush = TRUE)

plotlyProxyInvoke(p, method, ...)

Arguments

outputId: single-element character vector indicating the output ID map to modify (if invoked from a Shiny module, the namespace will be added automatically)
session: the Shiny session object to which the map belongs; usually the default value will suffice.
deferUntilFlush: indicates whether actions performed against this instance should be carried out right away, or whether they should be held until after the next time all of the outputs are updated.
p: a plotly proxy object (created with plotlyProxy)
method: a plotlyjs method to invoke. For a list of options, visit the plotlyjs function reference
...: unnamed arguments passed onto the plotly.js method
Examples

```r
if (require("shiny") && interactive()) {
  plotly_example("shiny", "proxy_relayout")
  plotly_example("shiny", "proxy_mapbox")
}
```

---

**plotly_build**

"Build" (i.e., evaluate) a plotly object

**Description**

This generic function creates the list object sent to plotly.js for rendering. Using this function can be useful for overriding defaults provided by `ggplotly/plot_ly` or for debugging rendering errors.

**Usage**

```r
plotly_build(p, registerFrames = TRUE)
```

**Arguments**

- `p` a ggplot object, or a plotly object, or a list.
- `registerFrames` should a frame trace attribute be interpreted as frames in an animation?

**Examples**

```r
p <- plot_ly(economics, x = ~date, y = ~pce)
# the unevaluated plotly object
str(p)
# the evaluated data
str(plotly_build(p)$x$data)
```

---

**plotly_data**

Obtain data associated with a plotly graph

**Description**

`plotly_data()` returns data associated with a plotly visualization (if there are multiple data frames, by default, it returns the most recent one).
Usage

plotly_data(p, id = p$x$cur_data)

## S3 method for class 'plotly'
groups(x)

## S3 method for class 'plotly'
group_by_(x, ...)

group_by_(data, ..., .dots, add = FALSE)

## S3 method for class 'plotly'
summarise_(x, ...)

## S3 method for class 'plotly'
mutate_(x, ...)

dot_(data, ..., .dots)

## S3 method for class 'plotly'
arrange_(data, ..., .dots)

## S3 method for class 'plotly'
select_(data, ..., .dots)

## S3 method for class 'plotly'
filter_(data, ..., .dots)

## S3 method for class 'plotly'
distinct_(data, ..., .dots)

## S3 method for class 'plotly'
slice_(data, ..., .dots)

## S3 method for class 'plotly'
rename_(data, ..., .dots)

## S3 method for class 'plotly'
transmute_(data, ..., .dots)

Arguments

p a plotly visualization

id a character string or number referencing an "attribute layer".

x a plotly visualization
... stuff passed onto the relevant method
.data a plotly visualization
.dots Used to work around non-standard evaluation. See vignette("nse") for details
.add By default, when add = FALSE, group_by will override existing groups. To instead add to the existing groups, use add = TRUE

Examples

# use group_by() to define groups of visual markings
p <- txhousing %>%
    group_by(city) %>%
    plot_ly(x = ~date, y = ~sales)

# plotly objects preserve data groupings
p %>%
    group_by(city)
    plotly_data(p)

# dplyr verbs operate on plotly objects as if they were data frames
p <- economics %>%
    plot_ly(x = ~date, y = ~unemploy / pop) %>%
    add_lines() %>%
    mutate(rate = unemploy / pop) %>%
    filter(rate == max(rate))

p %>%
    group_by(city)
    plotly_data(p)

# mutate() + do() + subplot() for trellis displays
d <- group_by(mpg, drv)
plots <- do(d, p = plot_ly(.x = ~cty, name = ~drv))
subplot(plots[["p"]], nrow = 3, shareX = TRUE)

# arrange displays by their mean
means <- summarise(d, mn = mean(cty, na.rm = TRUE))
means %>%
    dplyr::left_join(plots) %>%
    arrange(mn) %>%
    subplot(nrows = NROW(.), shareX = TRUE)

# more dplyr verbs applied to plotly objects
p <- mtcars %>%
    plot_ly(x = ~wt, y = ~mpg, name = "scatter trace") %>%
    add_markers() %>%
    slice(1) %>%
    slice(1) %>%
    filter(cyl == 4) %>%
    plotly_data() %>%
    filter(cyl == 4) %>%
    add_markers(name = "four cylinders")
plotly_empty

Create a complete empty plotly graph.

Description

Useful when used with subplot()

Usage

plotly_empty(...)

Arguments

... arguments passed onto plot_ly()

plotly_example

Run a plotly example(s)

Description

Provides a unified interface for running demos, shiny apps, and Rmd documents which are bundled with the package.

Usage

plotly_example(type = c("demo", "shiny", "rmd"), name, edit = TRUE, ...

Arguments

type the type of example
name the name of the example (valid names depend on type).
edit whether to open the relevant source files using file.edit. Only relevant if type is "shiny" or "rmd".
... arguments passed onto the suitable method.

Author(s)

Carson Sievert
plotly_image

Create a static image

Description

The images endpoint turns a plot (which may be given in multiple forms) into an image of the desired format.

Usage

plotly_image(x, width = 1000, height = 500, format = "png", scale = 1, out_file, ...)

Arguments

- x: either a plotly object or a list.
- width: Image width in pixels
- height: Image height in pixels
- format: The desired image format 'png', 'jpeg', 'svg', 'pdf', 'eps', or 'webp'
- scale: Both png and jpeg formats will be scaled beyond the specified width and height by this number.
- out_file: A filename for writing the image to a file.
- ...: arguments passed onto httr::POST

Examples

### Not run:

p <- plot_ly(x = 1:10)
png <- plotly_image(p, out_file = "plotly-test-image.png")
jpeg <- plotly_image(p, format = "jpeg", out_file = "plotly-test-image.jpeg")
svg <- plotly_image(p, format = "svg", out_file = "plotly-test-image.svg")
pdf <- plotly_image(p, format = "pdf", out_file = "plotly-test-image.pdf")

### End(Not run)

plotly_json

Inspect JSON sent to plotly.js

Description

This function is useful for obtaining/viewing/debugging JSON sent to plotly.js.
Usage

plotly_json(p = last_plot(), jsonedit = interactive(), pretty = TRUE, ...)

Arguments

p a plotly or ggplot object.
jsonedit use listviewer::jsonedit to view the JSON?
pretty adds indentation whitespace to JSON output. Can be TRUE/FALSE or a number specifying the number of spaces to indent. See jsonlite::prettify.
... other options passed onto listviewer::jsonedit

Examples

plotly_json(plot_ly())
plotly_json(plot_ly(), FALSE)

plotly_POST Create/Modify plotly graphs

Description

Deprecated: see api_create().

Usage

plotly_POST(x = last_plot(), filename = NULL, fileopt = "overwrite", sharing = c("public", "private", "secret"), ...)

Arguments

x either a ggplot object, a plotly object, or a list.
filename character string describing the name of the plot in your plotly account. Use / to specify directories. If a directory path does not exist it will be created. If this argument is not specified and the title of the plot exists, that will be used for the filename.
fileopt character string describing whether to create a "new" plotly, "overwrite" an existing plotly, "append" data to existing plotly, or "extend" it.
sharing If 'public', anyone can view this graph. It will appear in your profile and can appear in search engines. You do not need to be logged in to Plotly to view this chart. If 'private', only you can view this plot. It will not appear in the Plotly feed, your profile, or search engines. You must be logged in to Plotly to view this graph. You can privately share this graph with other Plotly users in your online Plotly account and they will need to be logged in to view this plot. If
'secret', anyone with this secret link can view this chart. It will not appear in the
Plotly feed, your profile, or search engines. If it is embedded inside a webpage
or an IPython notebook, anybody who is viewing that page will be able to view
the graph. You do not need to be logged in to view this plot.

... not used

See Also

plot_ly(), signup()

plot_dendro

Plot an interactive dendrogram

Description

This function takes advantage of nested key selections to implement an interactive dendrogram.
Selecting a node selects all the labels (i.e. leafs) under that node.

Usage

plot_dendro(d, set = "A", xmin = -50, height = 500, width = 500,
...)

Arguments

d a dendrogram object
set defines a crosstalk group
xmin minimum of the range of the x-scale
height height
width width
... arguments supplied to subplot()

Author(s)

Carson Sievert

See Also

plot_ly(), plot_mapbox(), ggplotly()

Examples

hc <- hclust(dist(USArrests), "ave")
dend1 <- as.dendrogram(hc)
plot_dendro(dend1, height = 600) %>%
hide_legend() %>%
highlight(persistent = TRUE, dynamic = TRUE)
plot_geo

Initiate a plotly-geo object

Description

Use this function instead of plot_ly() to initialize a plotly-geo object. This enforces the entire plot so use the scattergeo trace type, and enables higher level geometries like add_polygons() to work.

Usage

plot_geo(data = data.frame(), ..., offline = FALSE)

Arguments

data  A data frame (optional).
...  arguments passed along to plot_ly().
offline  whether or not to include geo assets so that the map can be viewed with or without an internet connection. The plotlyGeoAssets package is required for this functionality.

Author(s)

Carson Sievert

See Also

plot_ly(), plot_mapbox(), ggplotly()

Examples

map_data("world", "canada") %>%
group_by(group) %>%
plot_geo(x = ~long, y = ~lat) %>%
add_markers(size = I(1)))
**plot_ly**

*Initiate a plotly visualization*

**Description**

This function maps R objects to plotly.js, an (MIT licensed) web-based interactive charting library. It provides abstractions for doing common things (e.g., mapping data values to fill colors (via `color`) or creating animations (via `frame`) and sets some different defaults to make the interface feel more 'R-like' (i.e., closer to `plot()` and `ggplot2::qplot()`).

**Usage**

```r
plot_ly(data = data.frame(), ..., type = NULL, name, color,
        colors = NULL, alpha = NULL, stroke, strokes = NULL,
        alpha_stroke = 1, size, sizes = c(10, 100), span, spans = c(1, 20),
        symbol, symbols = NULL, linetype, linetypes = NULL, split, frame,
        width = NULL, height = NULL, source = "A")
```

**Arguments**

- `data` A data frame (optional) or `crosstalk::SharedData` object.
- `...` Arguments (i.e., attributes) passed along to the trace type. See `schema()` for a list of acceptable attributes for a given trace type (by going to `traces` -> `type` -> `attributes`). Note that attributes provided at this level may override other arguments (e.g., `plot_ly(x = 1:10, y = 1:10, color = I("red"), marker = list(color = "blue")))
- `type` A character string specifying the trace type (e.g., "scatter", "bar", "box", etc). If specified, it *always* creates a trace, otherwise
- `name` Values mapped to the trace’s name attribute. Since a trace can only have one name, this argument acts very much like `split` in that it creates one trace for every unique value.
- `color` Values mapped to relevant 'fill-color' attribute(s) (e.g. `fillcolor`, `marker.color`, `textfont.color`, etc.). The mapping from data values to color codes may be controlled using `colors` and `alpha`, or avoided altogether via `I()` (e.g., `color = I("red")`). Any color understood by `grDevices::col2rgb()` may be used in this way.
- `colors` Either a colorbrewer2.org palette name (e.g. "YlOrRd" or "Blues"), or a vector of colors to interpolate in hexadecimal "#RRGGBB" format, or a color interpolation function like `colorRamp()`.  
- `alpha` A number between 0 and 1 specifying the alpha channel applied to `color`. Defaults to 0.5 when mapping to `fillcolor` and 1 otherwise.
- `stroke` Similar to `color`, but values are mapped to relevant 'stroke-color' attribute(s) (e.g., `marker.line.color` and `line.color` for filled polygons). If not specified, stroke inherits from `color`.
- `strokes` Similar to `colors`, but controls the stroke mapping.
- `alpha_stroke` Similar to `alpha`, but applied to `stroke`.
size  (Numeric) values mapped to relevant 'fill-size' attribute(s) (e.g., marker.size, textfont.size, and error_x.width). The mapping from data values to symbols may be controlled using sizes, or avoided altogether via I() (e.g., size = I(30)).

sizes  A numeric vector of length 2 used to scale size to pixels.

span  (Numeric) values mapped to relevant 'stroke-size' attribute(s) (e.g., marker.line.width, line.width for filled polygons, and error_x.thickness) The mapping from data values to symbols may be controlled using spans, or avoided altogether via I() (e.g., span = I(30)).

spans  A numeric vector of length 2 used to scale span to pixels.

symbol  (Discrete) values mapped to marker.symbol. The mapping from data values to symbols may be controlled using symbols, or avoided altogether via I() (e.g., symbol = I("pentagon")). Any pch value or symbol name may be used in this way.

symbols  A character vector of pch values or symbol names.

linetype  (Discrete) values mapped to line.dash. The mapping from data values to symbols may be controlled using linetypes, or avoided altogether via I() (e.g., linetype = I("dash")). Any lty (see par) value or dash name may be used in this way.

linetypes  A character vector of lty values or dash names.

split  (Discrete) values used to create multiple traces (one trace per value).

frame  (Discrete) values used to create animation frames.

width  Width in pixels (optional, defaults to automatic sizing).

height  Height in pixels (optional, defaults to automatic sizing).

source  a character string of length 1. Match the value of this string with the source argument in event_data() to retrieve the event data corresponding to a specific plot (shiny apps can have multiple plots).

Details

Unless type is specified, this function just initiates a plotly object with 'global' attributes that are passed onto downstream uses of add_trace() (or similar). A formula must always be used when referencing column name(s) in data (e.g. plot_ly(mtcars, x = ~wt)). Formulas are optional when supplying values directly, but they do help inform default axis/scale titles (e.g., plot_ly(x = mtcars$wt) vs plot_ly(x = ~mtcars$wt))

Author(s)

Carson Sievert

References

https://plotly-r.com/overview.html
See Also

- For initializing a plotly-geo object: `plot Geo()`
- For initializing a plotly-mapbox object: `plot_mapbox()`
- For translating a ggplot2 object to a plotly object: `ggplotly()`
- For modifying any plotly object: `layout().add_trace().style()`
- For linked brushing: `highlight()`
- For arranging multiple plots: `subplot().crosstalk::bscols()`
- For inspecting plotly objects: `plotly_json()`
- For quick, accurate, and searchable plotly.js reference: `schema()`

Examples

```r
## Not run:

# plot_ly() tries to create a sensible plot based on the information you
give it. If you don't provide a trace type, plot_ly() will infer one.
plot_ly(economics, x = ~pop)
plot_ly(economics, x = ~date, y = ~pop)
# plot_ly() doesn't require data frame(s), which allows one to take
# advantage of trace type(s) designed specifically for numeric matrices
plot_ly(z = ~volcano)
plot_ly(z = ~volcano, type = "surface")

# plotly has a functional interface: every plotly function takes a plotly
# object as its first input argument and returns a modified plotly object
add_lines(plot_ly(economics, x = ~date, y = ~unemploy/pop))

# To make code more readable, plotly imports the pipe operator from magrittr
economics %>% plot_ly(x = ~date, y = ~unemploy/pop) %>% add_lines()

# Attributes defined via plot_ly() set 'global' attributes that
# are carried onto subsequent traces, but those may be over-written
plot_ly(economics, x = ~date, color = I("black")) %>%
  add_lines(y = ~unempmed) %>%
  add_lines(y = ~psavert, color = I("red"))

# Attributes are documented in the figure reference -> https://plot.ly/r/reference
# You might notice plot_ly() has named arguments that aren't in this figure
# reference. These arguments make it easier to map abstract data values to
# visual attributes.
p <- plot_ly(iris, x = ~Sepal.Width, y = ~Sepal.Length)
add_markers(p, color = ~Petal.Length, size = ~Petal.Length)
add_markers(p, color = ~Species)
add_markers(p, color = ~Species, colors = "Set1")
add_markers(p, symbol = ~Species)
add_paths(p, linetype = ~Species)
```

## End(Not run)
plot_mapbox

Initiate a plotly-mapbox object

Description

Use this function instead of `plot_ly()` to initialize a plotly-mapbox object. This enforces the entire plot so use the scattermapbox trace type, and enables higher level geometries like `add_polygons()`

to work

Usage

```
plot_mapbox(data = data.frame(), ...)
```

Arguments

- `data` A data frame (optional).
- `...` arguments passed along to `plot_ly()`. They should be valid scattermapbox attributes - https://plot.ly/r/reference/#scattermapbox. Note that x/y can also be used in place of lat/lon.

Author(s)

Carson Sievert

See Also

`plot_ly()`, `plot_geo()`, `ggplotly()`

Examples

```
## Not run:

plot_mapbox(res_mn)
plot_mapbox(res_mn, color = ~INDRESNAME)

map_data("world", "canada") %>%
  group_by(group) %>%
  plot_mapbox(x = ~long, y = ~lat) %>%
  add_polygons() %>%
  layout(
    mapbox = list(
      center = list(lat = ~median(lat), lon = ~median(long))
    )
  )

## End(Not run)
```
**print.api**

*Print method for a 'generic' API response*

**Description**

Print method for a 'generic' API response

**Usage**

```r
## S3 method for class 'api'
print(x, ...)
```

**Arguments**

- `x` a list.
- `...` additional arguments (currently ignored)

---

**print.api_grid**

*Print a plotly grid object*

**Description**

Print a plotly grid object

**Usage**

```r
## S3 method for class 'api_grid'
print(x, ...)
```

**Arguments**

- `x` a plotly grid object
- `...` additional arguments (currently ignored)
print.api_grid_local  
Print a plotly grid object

Description

Print a plotly grid object

Usage

```r
## S3 method for class 'api_grid_local'
print(x, ...)
```

Arguments

- `x`: a plotly grid object
- `...`: additional arguments (currently ignored)

print.api_plot  
Print a plot on plotly's platform

Description

Print a plot on plotly's platform

Usage

```r
## S3 method for class 'api_plot'
print(x, ...)
```

Arguments

- `x`: a plotly figure object
- `...`: additional arguments (currently ignored)
rangeslider

Description

Add a range slider to the x-axis

Usage

rangeslider(p, start = NULL, end = NULL, ...)

Arguments

p plotly object.
start a start date/value.
end an end date/value.
...
these arguments are documented here https://plot.ly/r/reference/#layout-xaxis-rangeslider

Author(s)

Carson Sievert

Examples

plot_ly(x = time(USAccDeaths), y = USAccDeaths) %>%
  add_lines() %>%
  rangeslider()

d <- tibble::tibble(
  time = seq(as.Date("2016-01-01"), as.Date("2016-08-31"), by = "days"),
  y = rnorm(seq_along(time))
)

plot_ly(d, x = ~time, y = ~y) %>%
  add_lines() %>%
  rangeslider(d$time[5], d$time[50])
raster2uri

Convert a raster object to a data URI

Description


Usage

```r
raster2uri(r, ...)
```

Arguments

- `r` an object coercable to a raster object via `as.raster()`
- `...` arguments passed onto `as.raster()`.

Author(s)

Carson Sievert

Examples

```r
# a red gradient (from ?as.raster)
r <- as.raster(matrix(hcl(0, 80, seq(50, 80, 10)), nrow = 4, ncol = 5))
plot(r)

# embed the raster as an image
plot_ly(x = 1, y = 1) %>%
  layout(
    images = list(list(
      source = raster2uri(r),
      xref = "paper",
      yref = "paper",
      x = 0, y = 0,
      sizex = 0.5, sizey = 0.5,
      xanchor = "left", yanchor = "bottom"
    )
  )
)```
Description

By default, plotly.js’ TypedArray polyfill is included as a dependency, so printing "just works" in any context. Many users won’t need this polyfill, so this function may be used to remove it and thus reduce the size of the page.

Usage

remove_typedarray_polyfill(p)

Arguments

p a plotly object

Details

The polyfill seems to be only relevant for those rendering plots via phantomjs and RStudio on some Windows platforms.

Examples

```r
## Not run:
p1 <- plot_ly()
p2 <- remove_typedarray_polyfill(p1)
t1 <- tempfile(fileext = ".html")
htmlwidgets::saveWidget(p1, t1)
file.info(t1)$size
htmlwidgets::saveWidget(p2, t1)
file.info(t1)$size
## End(Not run)
```

---

res_mn Minnesotan Indian Reservation Lands

Description

Minnesotan Indian Reservation Lands

Usage

res_mn
Format

An sf data frame with 13 features and 5 fields

References

https://www.dot.state.mn.us/maps/gdma/gis-data.html

---

**schema**

Acquire (and optionally display) plotly’s plot schema

---

**Description**

The schema contains valid attributes names, their value type, default values (if any), and min/max values (if applicable).

**Usage**

```r
schema(jsonedit = interactive(), ...)
```

**Arguments**

- `jsonedit` use listviewer::jsonedit to view the JSON?
- `...` other options passed onto listviewer::jsonedit

**Examples**

```r
s <- schema()

# retrieve acceptable 'layout.mapbox.style' values
if (!is.na(Sys.getenv('MAPBOX_TOKEN', NA))) {
  styles <- s$layout$layoutAttributes$mapbox$style$values
  subplot(
    plot_mapbox() %>% layout(mapbox = list(style = styles[3])),
    plot_mapbox() %>% layout(mapbox = list(style = styles[5]))
  )
}
```
**showRGB**

*View colors already formatted by toRGB()*

**Description**

Useful for viewing colors after they've been converted to plotly.js' color format – "rgba(255, 255, 255, 1)"

**Usage**

```
showRGB(x, ...)
```

**Arguments**

- `x` character string specifying color(s).
- `...` arguments passed along to `scales::show_col`.

**Author(s)**

Carson Sievert

**Examples**

```
showRGB(toRGB(colors()), labels = FALSE)
```

**signup**

*Create a new plotly account.*

**Description**

A sign up interface to plotly through the R Console.

**Usage**

```
signup(username, email, save = TRUE)
```

**Arguments**

- `username` Desired username.
- `email` Desired email.
- `save` If request is successful, should the username & API key be automatically stored as an environment variable in a .Rprofile?
Value

- api_key key to use with the api
- tmp_pw temporary password to access your plotly account

References

https://plot.ly/rest/

Examples

```r
## Not run:
# You need a plotly username and API key to communicate with the plotly API.

# If you don't already have an API key, you can obtain one with a valid
# username and email via signup().
s <- signup('anna.lyst', 'anna.lyst@plot.ly')

# If you already have a username and API key, please create the following
# environment variables:
Sys.setenv("plotly_username" = "me")
Sys.setenv("plotly_api_key" = "mykey")
# You can also change the default domain if you have a plotly server.
Sys.setenv("plotly_domain" = "http://mydomain.com")

# If you want to automatically load these environment variables when you
# start R, you can put them inside your ~/.Rprofile
# (see help(.Rprofile) for more details)

## End(Not run)
```

---

**Description**

Modify trace(s) of an existing plotly visualization. Useful when used in conjunction with `get_figure()`.

**Usage**

```r
style(p, ..., traces = NULL)
```

**Arguments**

- `p` A plotly visualization.
- `...` Visual properties.
- `traces` numeric vector. Which traces should be modified? By default, attributes placed in ... will be applied to every trace.
Author(s)
Carson Sievert

See Also
api_download_plot()

Examples

# style() is especially useful in conjunction with ggplotly()
# It allows you to leverage the underlying plotly.js library to change
# the return result of ggplotly()
(p <- ggplotly(qplot(data = mtcars, wt, mpg, geom = c("point", "smooth"))))

# removes hoverinfo for the line/ribbon traces (use `plotly_json()` to verify!)
style(p, hoverinfo = "none", traces = c(2, 3))

# another example with plot_ly() instead of ggplotly()
marker <- list(
  color = "red",
  line = list(
    width = 20,
    color = "black"
  )
)
(p <- plot_ly(x = 1:10, y = 1:10, marker = marker))

# note how the entire (marker) object is replaced if a list is provided
style(p, marker = list(line = list(color = "blue")))

# similar to plotly.js, you can update a particular attribute like so
# https://github.com/plotly/plotly.js/issues/1866#issuecomment-314115744
style(p, marker.line.color = "blue")
# this clobbers the previously supplied marker.line.color
style(p, marker.line = list(width = 2.5), marker.size = 10)

 subplot View multiple plots in a single view

Description
View multiple plots in a single view

Usage
subnetplot(..., nrows = 1, widths = NULL, heights = NULL,
  margin = 0.02, shareX = FALSE, shareY = FALSE, titleX = shareX,
  titleY = shareY, which_layout = "merge")
Arguments

... One of the following

• any number of plotly/ggplot2 objects.
• a list of plotly/ggplot2 objects.
• a tibble with one list-column of plotly/ggplot2 objects.

nrows number of rows for laying out plots in a grid-like structure. Only used if no domain is already specified.

widths relative width of each column on a 0-1 scale. By default all columns have an equal relative width.

heights relative height of each row on a 0-1 scale. By default all rows have an equal relative height.

margin either a single value or four values (all between 0 and 1). If four values are provided, the first is used as the left margin, the second is used as the right margin, the third is used as the top margin, and the fourth is used as the bottom margin. If a single value is provided, it will be used as all four margins.

shareX should the x-axis be shared amongst the subplots?

shareY should the y-axis be shared amongst the subplots?

titleX should x-axis titles be retained?

titleY should y-axis titles be retained?

which_layout adopt the layout of which plot? If the default value of "merge" is used, layout options found later in the sequence of plots will override options found earlier in the sequence. This argument also accepts a numeric vector specifying which plots to consider when merging.

Value

A plotly object

Author(s)

Carson Sievert

Examples

# pass any number of plotly objects to subplot()
p1 <- plot_ly(economics, x = ~date, y = ~unempmed)
p2 <- plot_ly(economics, x = ~date, y = ~unemploy)
subplot(p1, p2, p1, p2, nrows = 2, margin = 0.05)

#' # anchor multiple traces on the same legend entry
p1 <- add_lines(p1, color = 1("black"), name = "1st", legendgroup = "1st")
p2 <- add_lines(p2, color = 1("red"), name = "2nd", legendgroup = "2nd")

subplot(
  p1, style(p1, showlegend = FALSE),
p2, style(p2, showlegend = FALSE),
nrows = 2, margin = 0.05
)

# or pass a list
economics_long %>%
  split(.variable) %>%
  lapply(function(d) plot_ly(d, x = ~date, y = ~value)) %>%
  subplot(nrows = NROW(.), shareX = TRUE)

# or pass a tibble with a list-column of plotly objects
economics_long %>%
  group_by(variable) %>%
  do(p = plot_ly(.x = ~date, y = ~value)) %>%
  subplot(nrows = NROW(.), shareX = TRUE)

# learn more at https://cpsievert.github.io/plotly_book/subplot.html

---

**TeX**

*Render TeX in a plotly graph using MathJax*

---

**Description**

This function makes it slightly easier to render TeX in a plotly graph – it ensures that MathJax is included with the final result and also ensures the provided string is surrounded with $ (this is what plotly.js uses to declare a string as TeX).

**Usage**

TeX(x)

**Arguments**

- **x** a character vector

**See Also**

config

**Examples**

```
plot_ly(x = c(1, 2, 3, 4), y = c(1, 4, 9, 16)) %>%
  layout(title = TeX("\\text{Some mathjax: }\\alpha+\\beta x")) %>%
  config(mathjax = "cdn")
```
toRGB

*Convert R colours to RGBA hexadecimal colour values*

**Description**

Convert R colours to RGBA hexadecimal colour values

**Usage**

```r
torgb(x, alpha = 1)
```

**Arguments**

- `x` see the `col` argument in `col2rgb` for valid specifications
- `alpha` alpha channel on 0-1 scale

**Value**

hexadecimal colour value (if `is.na(x)`, return "transparent" for compatibility with Plotly)

**See Also**

`showRGB()`

**Examples**

```r
torgb("steelblue")
# [1] "rgba(70,130,180,1)"

m <- list(
  color = toRGB("red"),
  line = list(
    color = toRGB("black"),
    width = 19
  )
)

plot_ly(x = 1, y = 1, marker = m)
```
**to WebGL**  
*Convert trace types to WebGL*

**Description**

Convert trace types to WebGL.

**Usage**

```r
to WebGL(p)
```

**Arguments**

- `p`: a plotly or ggplot object.

**Examples**

```r
# currently no bargl trace type
to WebGL(ggplot() + geom_bar(aes(1:10)))
to WebGL(qplot(1:10, 1:10))
```

---

**to_basic**  
*Convert a geom to a "basic" geom.*

**Description**

This function makes it possible to convert ggplot2 geoms that are not included with ggplot2 itself. Users shouldn’t need to use this function. It exists purely to allow other package authors to write their own conversion method(s).

**Usage**

```r
to basic(data, prestats_data, layout, params, p, ...)
```

**Arguments**

- `data`: the data returned by `ggplot2::ggplot_build()`.
- `prestats_data`: the data before statistics are computed.
- `layout`: the panel layout.
- `params`: parameters for the geom, statistic, and 'constant' aesthetics
- `p`: a ggplot2 object (the conversion may depend on scales, for instance).
- `...`: currently ignored
Description

Description TBD.

Usage

wind

Format

A data frame with three variables: r, t, nms.
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