

Package: ckbplotr (via r-universe)

October 13, 2024

Title Create CKB Plots

Description ckbplotr provides functions to help create and style plots in R. It is being developed by, and primarily for, China Kadoorie Biobank researchers.

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LazyData true

Depends ggplot2 (>= 3.4.0), R (>= 2.10)

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Suggests patchwork, testthat

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URL <https://neilstats.github.io/ckbplotr/>,
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Contents

ckb_style	2
forest_plot	3
geom_text_move	8
ggpreview	9
plot_like_ckb	10
prepare_figure	10
save_figure	11
shape_plot	12
theme_ckb	15

Index	16
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ckb_style	<i>Make a ggplot into CKB style</i>
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Description

Make a ggplot into CKB style

Usage

```
ckb_style(
  xlims = NULL,
  ylims = NULL,
  gap = c(0.025, 0.025),
  ext = c(0, 0),
  ratio = 1.5,
  width = NULL,
  height = NULL,
  base_size = 11,
  base_line_size = base_size/22,
  colour = "black",
  plot.margin = margin(0.5, 1.5, 0.5, 0.5, "lines"),
  axes = "both"
)
```

Arguments

xlims	A numeric vector of length two. The limits of the x-axis.
ylims	A numeric vector of length two. The limits of the y-axis.
gap	A numeric vector of length two. The gap between plotting area and axis to the left and bottom of the plot, as a proportion of the x-axis length. (Default: c(0.025, 0.025))
ext	A numeric vector of length two. The extensions to add to the right and top of the plot, as a proportion of the x-axis length. (Default: c(0, 0))

ratio	The ratio (y-axis:x-axis) to use for the plot. Ignored if both width and height are set. (Default: 1.5)
width	A <code>grid::unit</code> object to set the width of the plot (not including the gap or extension).
height	A <code>grid::unit</code> object to set the height of the plot (not including the gap or extension).
base_size	base font size, given in pts.
base_line_size	base size for line elements
colour	Colour for non-data aspects of the plot. (Default: "black")
plot.margin	Margin around entire plot (Default: <code>margin(0.5, 0, 0.5, 0, "lines")</code>)
axes	Choice of axis lines to add to the plot, one of "both", "x" or "y". (Default: "both")

forest_plot

Make a forest plot with ggplot2

Description

Creates a forest plot with ggplot

Usage

```
forest_plot(
  panels,
  row.labels = NULL,
  row.labels.levels = NULL,
  rows = NULL,
  row.labels.heading = NULL,
  row.labels.space = c(0, 1, 0, 0),
  exponentiate = TRUE,
  logscale = exponentiate,
  panel.names = NULL,
  panel.headings = NULL,
  col.key = "key",
  col.estimate = c("estimate", "est", "beta", "loghr"),
  col.stderr = c("stderr", "std.err", "se"),
  col.lci = NULL,
  col.uci = NULL,
  col.left = NULL,
  col.right = NULL,
  col.right.parse = FALSE,
  col.left.heading = "",
  col.right.heading = as.list(xlab),
  col.left.pos = NULL,
  col.right.pos = NULL,
```

```
col.left.hjust = 1,
col.right.hjust = 0,
col.left.gap = c("I", "W"),
col.right.gap = c("I", "W"),
col.heading.space = 0,
estcolumn = TRUE,
col.keep = NULL,
ci.delim = ", ",
digits = 2,
title = "",
xlab = "HR (95% CI)",
xlim = NULL,
xticks = NULL,
nullval = NULL,
col.diamond = NULL,
diamond = NULL,
col.bold = NULL,
bold.labels = NULL,
scalepoints = FALSE,
minse = NULL,
pointsize = 3,
shape = 15,
plotcolour = "black",
colour = plotcolour,
cicolour = colour,
fill = colour,
ciunder = NULL,
addtext = NULL,
bottom.space = 0.7,
left.space = NULL,
right.space = NULL,
mid.space = unit(5, "mm"),
plot.margin = margin(8, 8, 8, 8, "mm"),
panel.width = NULL,
panel.height = NULL,
base_size = 11,
base_line_size = base_size/22,
stroke = 0,
diamonds.linewidth = base_line_size,
quiet = FALSE,
printplot = !quiet,
showcode = !quiet,
data.function = NULL,
addaes = NULL,
addarg = NULL,
add = NULL,
envir = NULL,
blankrows = NULL
```

)

Arguments

panels	A list of data frames. These should include columns or point estimates, and standard errors or confidence interval limits. If you specify a row.labels data frame, then they must also all contain a key column with the same name (which can be specified by col.key).
row.labels	A data frame that contains the labels to be used for the rows of the plot. Use NA if a lower level heading is not required for a given row.
row.labels.levels	A character vector. The names of columns in row.labels to use as headings/subheadings/labels for labelling rows.
rows	If set, then only rows matching these labels (at the first level) will be included.
row.labels.heading	Title to be placed above row labels.
row.labels.space	A numeric vector specifying the space after a row label heading, at the end of a row label heading 'section'. (Default: c(0, 1, 0, 0))
exponentiate	Exponentiate estimates (and CIs) before plotting. (Default: TRUE)
logscale	Use log scale on the axis, and add a line at null effect. (Default: exponentiate)
panel.names	A character vector. The names to be used for each forest plot panel. If none provided, then they will be numbered 1, 2, 3 ...
panel.headings	Titles to be placed above each forest plot.
col.key	Name of column that links the results given in each data frame provided in panels and the labels given in row.labels. If row.labels data frame is not given, then this column will be used as row labels. (Default: "key")
col.estimate, col.stderr, col.lci, col.uci	Names of columns for: point estimates, standard errors, lower and upper limits of confidence intervals.
col.left, col.right	Names of columns to be printed to the left/right of the plot.
col.right.parse	A logical vector, the same length as col.right (+ 1 if estcolumn = TRUE). Should the contents of the columns be parsed into expressions. (Default: FALSE)
col.left.heading, col.right.heading	Headings for columns.
col.left.pos, col.right.pos	A unit vector to position col.left/col.right columns.
col.left.hjust, col.right.hjust	A numeric vector. The horizontal justification of col.left/col.right columns. (Default: 1)
col.left.gap, col.right.gap	A character vector of length two. The two characters control the gaps between the first text column and the panel, and successive text columns. (Default: c("I", "W"))

<code>col.heading.space</code>	Position of the titles given by <code>col.left.heading</code> and <code>col.right.heading</code> . Increase to move them up. (Default: 0)
<code>estcolumn</code>	Include column of estimates and confidence intervals to the right of each plot. (Default: TRUE)
<code>col.keep</code>	Names of additional columns to be kept in returned data frame.
<code>ci.delim</code>	Character string to separate lower and upper limits of confidence interval. (Default: ", ")
<code>digits</code>	Number of digits after decimal point to show for estimates and confidence intervals. (Default: 2)
<code>title</code>	Title to appear at the top of the plot.
<code>xlab</code>	Label to appear below the x-axis. (Default: "HR (95% CI)")
<code>xlim</code>	A numeric vector. The limits of the x axis.
<code>xticks</code>	A numeric vector. The tick points of the x axis.
<code>nullval</code>	Add a vertical reference line at this value. (If <code>logscale == TRUE</code> then by default it will be added at 1, but use NA not to plot this line.)
<code>col.diamond</code>	Plot estimates and CIs as diamonds. Name of a column of logical values.
<code>diamond</code>	Alternative to <code>col.diamond</code> . A character vectors identify the rows (using the key values) for which the estimate and CI should be plotted using a diamond.
<code>col.bold</code>	Plot text as bold. Name of a column of logical values.
<code>bold.labels</code>	A character vector identifying row labels (using key values) which should additionally be bold. (Default: NULL)
<code>scalepoints</code>	Should the points be scaled by inverse of the standard error? (Default: FALSE)
<code>minse</code>	Minimum standard error to use when scaling point size. (Default will use minimum in the data.)
<code>pointsize</code>	The (largest) size of box to use for plotting point estimates. (Default: 3)
<code>shape</code>	Shape of points. An integer, or name of a column of integers. (Default: 15 (square))
<code>plotcolour</code>	Colour for all parts of the plot. (Default: "black")
<code>colour</code>	Colour of points. Name of a colour, or name of a column of colour names. (Default will use <code>plotcolour</code> .)
<code>cicolour</code>	Colour of CI lines. Colour of CI lines. Name of a colour, or name of a column of colour names. (Default will use <code>colour</code> .)
<code>fill</code>	Fill colour of points. Name of a colour, or name of a column of colour names. (Default will use <code>colour</code> .)
<code>ciunder</code>	Plot CI lines before points. A logical value, or name of a column of logical values. (Default will plot CI lines after points.)
<code>addtext</code>	A list of data frames. List must be the same length as panels. Data frames should contain a column with the name specified in <code>col.key</code> , and one or more of: <ol style="list-style-type: none"> 1. a column named 'text' containing character strings 2. columns named 'het_dof', 'het_stat', and 'het_p' containing character strings

	3. columns names 'trend_stat' and 'trend_p' containing character strings
	The character strings, heterogeneity test, and trend test results will be plotted in the column of estimates and CIs, below the row with the key given in the col.key column.
bottom.space	Space between bottom row and axis. (Default: 0.7)
left.space, right.space, mid.space	Space to the left/right/between panels. (Default mid.space: unit(5, "mm"))
plot.margin	Plot margin, given as margin(top, right, bottom, left, units). (Default: margin(8, 8, 8, 8, "mm"))
panel.width, panel.height	Set width/height of panels. A grid::unit object, if a numeric is given assumed to be in mm. If panel.width is used, will also apply different formatting to narrow CIs.
base_size	base font size, given in pts.
base_line_size	base size for line elements
stroke	Size of outline of shapes. (Default: 0)
diamonds.linewidth	Line width for diamonds. (Default: base_line_size)
quiet	Set to TRUE to not print the plot nor show generated code in the RStudio 'Viewer' pane. (Default: FALSE)
printplot	Print the plot. (Default: !quiet)
showcode	Show the ggplot2 code to generate the plot in RStudio 'Viewer' pane. (Default: !quiet)
data.function	Name of a function to apply to data frame before plotting.
addaes, addarg, add	Methods for customising the plot. See documentation for details.
envir	Environment in which to evaluate the plot code. May be useful when calling this function inside another function.
blankrows	DEPRECATED

Details

The function returns the plot and ggplot2 code to create the plot. In RStudio, the ggplot2 code will be shown in the viewer.

Value

A list:

plot the plot

code ggplot2 code to generate the plot

geom_text_move	<i>Text that can be moved</i>
----------------	-------------------------------

Description

This geom adds a fixed horizontal and/or vertical move to `ggplot2::geom_text()`

Usage

```
geom_text_move(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,
  parse = FALSE,
  nudge_x = 0,
  nudge_y = 0,
  move_x = unit(0, "pt"),
  move_y = unit(0, "pt"),
  check_overlap = FALSE,
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE
)
```

Arguments

mapping	Set of aesthetic mappings created by <code>aes()</code> . If specified and <code>inherit.aes = TRUE</code> (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
data	The data to be displayed in this layer. There are three options: If <code>NULL</code> , the default, the data is inherited from the plot data as specified in the call to <code>ggplot()</code> . A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(., 10)</code>).
stat	The statistical transformation to use on the data for this layer, either as a <code>ggproto</code> <code>Geom</code> subclass or as a string naming the stat stripped of the <code>stat_</code> prefix (e.g. "count" rather than "stat_count")
position	Position adjustment, either as a string, or the result of a call to a position adjustment function. Cannot be jointly specified with <code>nudge_x</code> or <code>nudge_y</code> .

...	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.
<code>parse</code>	If TRUE, the labels will be parsed into expressions and displayed as described in <code>?plotmath</code> .
<code>nudge_x</code> , <code>nudge_y</code>	Horizontal and vertical adjustment to nudge labels by. Useful for offsetting text from points, particularly on discrete scales. Cannot be jointly specified with <code>position</code> .
<code>move_x</code>	Unit value to move text horizontally (Default: <code>unit(0, "pt")</code>)
<code>move_y</code>	Unit value to move text vertically (Default: <code>unit(0, "pt")</code>)
<code>check_overlap</code>	If TRUE, text that overlaps previous text in the same layer will not be plotted. <code>check_overlap</code> happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling <code>geom_text()</code> . Note that this argument is not supported by <code>geom_label()</code> .
<code>na.rm</code>	If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.
<code>show.legend</code>	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.
<code>inherit.aes</code>	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. <code>borders()</code> .

Aesthetics

`geom_text_move()` understands the same aesthetics as `ggplot2::geom_text()`

ggpreview

Create a plot preview and display it in the Viewer pane.

Description

This function saves a `ggplot2` plot to a temporary PNG file and then embeds it in an HTML page, which is opened in the Viewer pane.

Usage

```
ggpreview(...)
```

Arguments

... Arguments passed to `ggsave` and the PNG device function.

Device

The plot is saved using ggsave with the png device, regardless of what is specified in the call, so any arguments not used by ggsave or png are ignored.

plot_like_ckb	<i>Make a ggplot into CKB style</i>
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Description

Make a ggplot into CKB style

Usage

```
plot_like_ckb(plot, ...)
```

Arguments

plot	A ggplot2 plot
...	Arguments passed to ckb_style()

Value

A ggplot2 plot.

prepare_figure	<i>Prepare figure for saving</i>
----------------	----------------------------------

Description

Prepare figure for saving

Usage

```
prepare_figure(
  figure,
  title = "",
  title.pos = grid::unit.c(unit(1.27/2, "cm"), unit(1, "npc") - unit(1.27/2, "cm")),
  title.just = c(0, 1),
  title.gpar = list(fontsize = 12, fontface = "bold"),
  footer = "",
  footer.pos = grid::unit.c(unit(1.27/2, "cm"), unit(1.27/3, "cm")),
  footer.just = c(0, 0),
  footer.gpar = list(fontsize = 9),
  margin = unit(c(2.27, 1.27, 1.27, 1.27), units = "cm"),
  size = NULL,
```

```

    valign = 0.5,
    halign = 0.5,
    pagesize = c("A4", "A5"),
    landscape = FALSE,
    pagedim = NULL
)

```

Arguments

figure	Plot (or graphical object).
title	Title to be added to the page. (Default: "")
title.pos	Position of the title text. Default is 1/4 inch from top left of page. (Default: <code>unit.c(unit(1.27/2, "cm"), unit(1, "npc") - unit(1.27/2, "cm"))</code>)
title.just	Justification of the title text. (Default: <code>c(0, 1)</code>)
title.gpar	Graphical parameters for title. (Default: <code>list(fontsize = 12, fontface = "bold")</code>)
footer	Footer to be added to the page. (Default: "")
footer.pos	Position of the footer text. Default is 1/6 inch from bottom and 1/4 inch from left of page. (Default: <code>unit.c(unit(1.27/2, "cm"), unit(1.27/3, "cm"))</code>)
footer.just	Justification of the footer text. (Default: <code>c(0, 0)</code>)
footer.gpar	Graphical parameters for footer. (Default: <code>list(fontsize = 9)</code>)
margin	Margin to be placed around the plot. Default is 2.27cm top, 1.27cm (1/2 inch) other sides. (Default: <code>unit(c(2.27, 1.27, 1.27, 1.27), units = "cm")</code>)
size	A unit vector of length two (width, height). Size of plot (a width/height larger than page weight/height minus margins will be ignored), centred within margins. By default, plot will fill the space within margins.
valign	If size is set, where to place figure within margins. 1 = top, 0.5 = middle, 0 = bottom. (Default: 0.5)
halign	If size is set, where to place figure within margins. 1 = right, 0.5 = middle, 0 = left (Default: 0.5)
pagesize	Page size of output: "A4" or "A5". (Default: "A4")
landscape	Landscape page orientation? (Default: False)
pagedim	Dimensions (width, height) of output. Overrides pagesize and landscape arguments if used.

 save_figure

Output plots as files

Description

Output plots as files

Usage

```
save_figure(
  figure,
  filename,
  cropped = NULL,
  args = NULL,
  args_cropped = NULL,
  preview = FALSE,
  ...
)
```

Arguments

figure	Plot (or graphical object).
filename	Name of file to create.
cropped	Name of second output file of the figure without margins or title.
args	List of arguments passed to <code>ggplot2::ggsave()</code> for the main figure.
args_cropped	List of arguments passed to <code>ggplot2::ggsave()</code> for the cropped figure.
preview	Preview the output in the RStudio Viewer pane. (Default: False)
...	Other arguments passed to prepare_figure .

shape_plot

Make a shape plot with ggplot2

Description

Make a shape plot with ggplot2

Usage

```
shape_plot(
  data,
  col.x = "x",
  col.estimate = c("estimate", "est", "beta", "loghr"),
  col.stderr = c("stderr", "std.err", "se"),
  col.lci = NULL,
  col.uci = NULL,
  col.n = NULL,
  exponentiate = FALSE,
  logscale = exponentiate,
  scalepoints = FALSE,
  digits = 2,
  minse = NA,
  pointsize = 3,
  col.group = NULL,
```

```

shape = 15,
plotcolour = "black",
colour = plotcolour,
cicolour = colour,
fill = colour,
ciunder = NULL,
lines = FALSE,
xlims,
ylims,
height = NULL,
width = NULL,
gap = c(0.025, 0.025),
ext = c(0.025, 0.025),
ratio = 1.5,
base_size = 11,
base_line_size = base_size/22,
stroke = base_size/22,
xbreaks = NULL,
ybreaks = NULL,
xlab = "Risk factor",
ylab = "Estimate (95% CI)",
legend.name = "",
legend.position = "top",
title = NULL,
quiet = FALSE,
printplot = !quiet,
showcode = !quiet,
addaes = NULL,
addarg = NULL,
add = NULL,
envir = NULL
)

```

Arguments

<code>data</code>	The data frame containing estimates to be plotted.
<code>col.x</code>	Name of column that provides the x-axis value (e.g. exposure, risk factor, dependent variable). (Default: "x")
<code>col.estimate</code>	Name of column that provides point estimates. (Default: "estimate")
<code>col.stderr</code>	Name of column that provides standard errors. (Default: "stderr")
<code>col.lci</code>	Name of column that provides lower limit of confidence intervals.
<code>col.uci</code>	Name of column that provides upper limit of confidence intervals.
<code>col.n</code>	Name of column that provides number to be plotted below CIs.
<code>exponentiate</code>	Exponentiate estimates (and CIs) before plotting, use log scale on the axis. (Default: FALSE)
<code>logscale</code>	Use log scale for vertical axis. (Default: exponentiate)

scalepoints	Should the points be scaled by inverse of the standard error? (Default: FALSE)
digits	Number of digits to use in text of estimates.
minse	Minimum standard error to use when scaling point size. (Default will use minimum in the data.)
pointsize	The (largest) size of box to use for plotting point estimates. (Default: 3)
col.group	Name of column that groups the estimates. (Default: NULL)
shape	Shape of points. An integer, or name of a column of integers. (Default: 15)
plotcolour	Colour for non-data aspects of the plot. (Default: "black")
colour	Colour of points. Name of a colour, or name of a column of colour names. (Default will use plotcolour)
cicolour	Colour of CI lines. Colour of CI lines. Name of a colour, or name of a column of colour names. (Default will use plotcolour)
fill	Fill colour of points. Fill colour of points. Name of a colour, or name of a column of colour names. (Default will use plotcolour)
ciunder	Plot CI lines before points. A logical value, or name of a column of logical values. (Default will plot CI lines after points.)
lines	Plot lines (linear fit through estimates, weighted by inverse variance). (Default: FALSE)
xlims	A numeric vector of length two. The limits of the x-axis.
ylims	A numeric vector of length two. The limits of the y-axis.
height	Panel height to use and apply different formatting to short CIs. A grid::unit() object, or if numeric is assumed to be in mm.
width	Panel width. A grid::unit() object, or if numeric is assumed to be in mm.
gap	A numeric vector of length two. The gap between plotting area and axis to the left and bottom of the plot, as a proportion of the x-axis length. (Default: c(0.025, 0.025))
ext	A numeric vector of length two. The extensions to add to the right and top of the plot, as a proportion of the x-axis length. (Default: c(0.025, 0.025))
ratio	The ratio (y-axis:x-axis) to use for the plot. (Default: 1.5)
base_size	base font size, given in pts.
base_line_size	base size for line elements
stroke	Size of outline of shapes. (Default: base_size/22)
xbreaks	Breaks for the x axis. Passed to ggplots::scale_x_continuous. (Default: NULL)
ybreaks	Breaks for the y axis. Passed to ggplots::scale_y_continuous. (Default: NULL)
xlab	Label for x-axis. (Default: "Risk factor")
ylab	Label for y-axis. (Default: "Estimate (95% CI)")
legend.name	The name of the colour scale/legend for groups. (Default: "")
legend.position	Position of the legend for groups ("none", "left", "right", "bottom", "top", or two-element numeric vector). (Default: "top")

title	Plot title. (Default: NULL)
quiet	Set to TRUE to not print the plot nor show generated code in the RStudio 'Viewer' pane. (Default: FALSE)
printplot	Print the plot. (Default: !quiet)
showcode	Show the ggplot2 code to generate the plot in RStudio 'Viewer' pane. (Default: !quiet)
addaes, addarg, add	Methods for customising the plot. See documentation for details.
envir	Environment in which to evaluate the plot code. May be useful when calling this function inside another function.

Value

A list:

plot the plot

code ggplot2 code to generate the plot

theme_ckb	<i>CKB ggplot theme</i>
-----------	-------------------------

Description

Based on theme_bw

Usage

```
theme_ckb(
  base_size = 11,
  base_line_size = base_size/22,
  colour = "black",
  plot.margin = margin(0.5, 1.5, 0.5, 0.5, "lines")
)
```

Arguments

base_size	base font size, given in pts.
base_line_size	base size for line elements
colour	Colour for non-data aspects of the plot. (Default: "black")
plot.margin	Margin around entire plot (Default: margin(0.5, 0, 0.5, 0, "lines"))

Index

* datasets

- geom_text_move, 8
- aes(), 8
- borders(), 9
- ckb_style, 2
- forest_plot, 3
- fortify(), 8
- geom_text_move, 8
- GeomTextMove (geom_text_move), 8
- ggplot(), 8
- ggpreview, 9
- layer(), 9
- plot_like_ckb, 10
- prepare_figure, 10, 12
- save_figure, 11
- shape_plot, 12
- theme_ckb, 15