

Package ‘deeptime’

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Title Plotting Tools for Anyone Working in Deep Time

Version 0.2.0

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Description Extends the functionality of other plotting packages like 'ggplot2' and 'lattice' to help facilitate the plotting of data over long time intervals, including, but not limited to, geological, evolutionary, and ecological data. The primary goal of 'deeptime' is to enable users to add highly customizable timescales to their visualizations. Other functions are also included to assist with other areas of deep time visualization.

URL <https://github.com/willgearty/deeptime>

BugReports <https://github.com/willgearty/deeptime/issues>

Depends R (>= 3.4)

License GPL (>= 2)

Encoding UTF-8

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LazyData true

biocViews

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R topics documented:

coord_geo	2
coord_trans_flip	5
coord_trans_xy	6
disparity_through_time	7
eons	9
epochs	10
eras	10
getScaleData	11
ggarrange2	12
gggeo_scale	13
gggeo_scale_old	17
gtable_frame2	20
panel.disparity	21
periods	22
stages	22

Index	24
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coord_geo	<i>Transformed coordinate system with geological timescale</i>
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Description

coord_geo behaves similarly to [coord_trans](#) in that it occurs after statistical transformation and will affect the visual appearance of geoms. The main difference is that it also adds a geological timescale to the specified side of the plot.

Usage

```
coord_geo(
  pos = "bottom",
  dat = "periods",
  xlim = NULL,
  ylim = NULL,
  xtrans = identity_trans(),
  ytrans = identity_trans(),
  clip = "on",
  expand = FALSE,
  fill = NULL,
  color = "black",
  alpha = 1,
  height = unit(2, "line"),
  lab = TRUE,
  rot = 0,
  abbrev = TRUE,
  skip = c("Quaternary", "Holocene", "Late Pleistocene"),
```

```

    size = 5,
    lwd = 0.25,
    neg = FALSE,
    bord = c("left", "right", "top", "bottom"),
    center_end_labels = FALSE,
    dat_is_discrete = FALSE,
    fittext_args = list()
  )

```

Arguments

pos	Which side to add the scale to (left, right, top, or bottom). First letter may also be used.
dat	Either A) a string indicating a built-in dataframe with interval data from the ICS ("periods", "epochs", "stages", "eons", or "eras"), B) a string indicating a timescale from macrostrat (see list here: https://macrostrat.org/api/defs/timescales?all), or C) a custom data.frame of time interval boundaries (see Details).
xlim, ylim	Limits for the x and y axes.
xtrans, ytrans	Transformers for the x and y axes. For more information see coord_trans .
clip	Should drawing be clipped to the extent of the plot panel? For more information see coord_trans .
expand	If 'FALSE', the default, limits are taken exactly from the data or 'xlim'/'ylim'. If 'TRUE', adds a small expansion factor to the limits to ensure that data and axes don't overlap.
fill	The fill color of the boxes. The default is to use the colors included in dat. If a custom dataset is provided with dat without color and without fill, a greyscale will be used. Custom fill colors can be provided with this option and will be recycled if/as necessary.
color	The outline color of the interval boxes.
alpha	The transparency of the fill colors.
height	The height (or width if pos is left or right) of the scale.
lab	Whether to include labels.
rot	The amount of counter-clockwise rotation to add to the labels (in degrees).
abbrv	If including labels, whether to use abbreviations instead of full interval names.
skip	A vector of interval names indicating which intervals should not be labeled. If abbrv is TRUE, this can also include interval abbreviations.
size	Label size. Either a number as you would specify in geom_text or "auto" to use geom_fit_text .
lwd	Line width.
neg	Set this to true if your x-axis is using negative values.
bord	A vector specifying on which sides of the scale to add borders (same options as pos).

center_end_labels	Should labels be centered within the visible range of intervals at the ends of the axis?
dat_is_discrete	Are the ages in dat already converted for a discrete scale?
fittext_args	A list of named arguments to provide to <code>geom_fit_text</code> . Only used if size is set to "auto".

Details

Transforming the side with the scale is not currently implemented. If a custom data.frame is provided (with dat), it should consist of at least 3 columns of data. See `data(periods)` for an example.

- The name column lists the names of each time interval. These will be used as labels if no abbreviations are provided.
- The max_age column lists the oldest boundary of each time interval.
- The min_age column lists the youngest boundary of each time interval.
- The abbr column is optional and lists abbreviations that may be used as labels.
- The color column is also optional and lists a hex color code (which can be obtained with `rgb()`) for each time interval.

If the axis of the time scale is discrete, max_age and min_age will automatically be converted to the discrete scale. In this case, the categories of the discrete axis should match the values in the name column. If the ages within dat are already discretized, you can set dat_is_discrete to TRUE to prevent this automatic conversion. This can be useful for adding a time scale where categories and time intervals are not 1:1.

pos may also be a list of sides (including duplicates) if multiple time scales should be added to the plot. In this case, dat, fill, color, alpha, height, lab, rot, abbrev, skip, size, lwd, neg, bord, center_end_labels, and dat_is_discrete can also be lists. If these lists are not as long as pos, the elements will be recycled. If individual values are used for these parameters, they will be applied to all time scales.

Examples

```
library(ggplot2)
#single scale on bottom
ggplot() +
  geom_point(aes(y = runif(1000, 0, 8), x = runif(1000, 0, 1000))) +
  scale_x_reverse() +
  coord_geo(xlim = c(1000, 0), ylim = c(0,8)) +
  theme_classic()

#stack multiple scales
ggplot() +
  geom_point(aes(y = runif(1000, 0, 8), x = runif(1000, 0, 100))) +
  scale_x_reverse() +
  coord_geo(xlim = c(100, 0), ylim = c(0,8), pos = as.list(rep("bottom", 3)),
  dat = list("stages", "epochs", "periods"),
  height = list(unit(4, "lines"), unit(4, "lines"), unit(2, "line"))),
```

```
rot = list(90, 90, 0), size = list(2.5, 2.5, 5), abbrev = FALSE) +  
theme_classic()
```

coord_trans_flip	<i>Transformed and flipped Cartesian coordinate system</i>
------------------	--

Description

coord_trans_flip behaves similarly to [coord_trans](#) in that it occurs after statistical transformation and will affect the visual appearance of geoms. The main difference is that it also flips the x and y coordinates like [coord_flip](#).

Usage

```
coord_trans_flip(  
  x = "identity",  
  y = "identity",  
  xlim = NULL,  
  ylim = NULL,  
  clip = "on",  
  expand = TRUE  
)
```

Arguments

x	Transformers for x and y axes or their names.
y	Transformers for x and y axes or their names.
xlim	Limits for the x and y axes.
ylim	Limits for the x and y axes.
clip	Should drawing be clipped to the extent of the plot panel? A setting of "on" (the default) means yes, and a setting of "off" means no. In most cases, the default of "on" should not be changed, as setting clip = "off" can cause unexpected results. It allows drawing of data points anywhere on the plot, including in the plot margins. If limits are set via xlim and ylim and some data points fall outside those limits, then those data points may show up in places such as the axes, the legend, the plot title, or the plot margins.
expand	If TRUE, the default, adds a small expansion factor to the limits to ensure that data and axes don't overlap. If FALSE, limits are taken exactly from the data or xlim/ylim.

Examples

```
library(ggplot2)  
ggplot(mtcars, aes(displ, wt)) +  
  geom_point() +  
  coord_trans_flip(x = "log10", y = "log10")
```

 coord_trans_xy

Transformed XY Cartesian coordinate system

Description

coord_trans_xy behaves similarly to [coord_trans](#) in that it occurs after statistical transformation and will affect the visual appearance of geoms. The main difference is that it takes a single transformer that is applied to the x and y axes simultaneously. Any transformers produced by [linear_trans](#) that have x and y arguments should work, but any other transformers produced using [trans_new](#) that take x and y arguments should also work. Axis limits will be adjusted to account for transformation unless limits are specified with 'xlim' or 'ylim'. This only works with geoms where all points are defined with x and y coordinates (e.g. [geom_point](#), [geom_polygon](#)). This does not currently work with geoms where point coordinates are extrapolated (e.g. [geom_rect](#)).

Usage

```
coord_trans_xy(
  trans = NULL,
  xlim = NULL,
  ylim = NULL,
  expand = TRUE,
  default = FALSE,
  clip = "on"
)
```

Arguments

trans	Transformer for x and y axes.
xlim, ylim	Limits for the x and y axes.
expand	If 'TRUE', the default, adds a small expansion factor to the limits to ensure that data and axes don't overlap. If 'FALSE', limits are taken exactly from the data or 'xlim'/'ylim'.
default	Is this the default coordinate system? If 'FALSE' (the default), then replacing this coordinate system with another one creates a message alerting the user that the coordinate system is being replaced. If 'TRUE', that warning is suppressed.
clip	Should drawing be clipped to the extent of the plot panel? A setting of "on" (the default) means yes, and a setting of "off" means no. In most cases, the default of "on" should not be changed, as setting 'clip = "off"' can cause unexpected results. It allows drawing of data points anywhere on the plot, including in the plot margins. If limits are set via 'xlim' and 'ylim' and some data points fall outside those limits, then those data points may show up in places such as the axes, the legend, the plot title, or the plot margins.

Examples

```

#make transformer
library(ggforce)
trans <- linear_trans(shear(2, 0), rotate(-pi / 3))

#set up data to be plotted
square <- data.frame(x = c(0, 0, 4, 4), y = c(0, 1, 1, 0))
points <- data.frame(x = runif(100, 0, 4), y = runif(100, 0, 1))

#plot data normally
library(ggplot2)
ggplot(data = points, aes(x = x, y = y)) +
  geom_polygon(data = square, fill = NA, color = "black") +
  geom_point(color = 'black') +
  coord_cartesian(expand = FALSE) +
  theme_classic()

#plot data with transformation
ggplot(data = points, aes(x = x, y = y)) +
  geom_polygon(data = square, fill = NA, color = "black") +
  geom_point(color = 'black') +
  coord_trans_xy(trans = trans, expand = FALSE) +
  theme_classic()

```

disparity_through_time

Disparity through time plot using lattice

Description

Plots points on 2-D surfaces within a 3-D framework. See [wireframe](#) and [panel.cloud](#) for customization options.

Usage

```

disparity_through_time(
  x,
  data,
  groups,
  pch = 16,
  col.point = c("blue"),
  scales = list(arrows = FALSE, distance = 1, col = "black", z = list(rot = 90)),
  colorkey = FALSE,
  screen = list(z = 90, x = 70, y = 180),
  aspect = c(1.5, 4),
  drape = TRUE,
  col.regions = c("white"),
  alpha.regions = c(1),

```

```

perspective = FALSE,
R.mat = matrix(c(1, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1, 0, 0, 0, 0, 1), 4, 4),
par.settings = list(axis.line = list(col = "transparent"), layout.heights =
  list(top.padding = 0, main.key.padding = 0, key.axis.padding = 0, axis.xlab.padding =
    0, xlab.key.padding = 0, key.sub.padding = 0, bottom.padding = 0), layout.widths =
  list(left.padding = 0, key.ylab.padding = 0, ylab.axis.padding = 0, axis.key.padding
    = 0, right.padding = 0)),
lattice.options = list(axis.padding = list(factor = 0)),
...
)

```

Arguments

<code>x</code>	a formula (most likely of the form $z \sim x * y$)
<code>data</code>	a data frame in which variables in the formula are to be evaluated
<code>groups</code>	a variable in data to be used as a grouping variable (this is probably the z variable)
<code>pch</code>	the point type
<code>col.point</code>	color(s) for points on surfaces
<code>scales</code>	a list specifying how the axes are drawn (see xyplot for details)
<code>colorkey</code>	logical, should a legend be drawn (or a list describing the legend; see levelplot for details)
<code>screen</code>	a list of the rotations that should be applied to each axis
<code>aspect</code>	a numeric vector of length 2, giving the relative aspects of the y-size/x-size and z-size/x-size of the enclosing cube
<code>drape</code>	logical, whether the surfaces should be colored based on <code>col.regions</code> and <code>alpha.regions</code>
<code>col.regions</code>	color(s) for surfaces
<code>alpha.regions</code>	alpha value(s) for surfaces
<code>perspective</code>	logical, whether to plot a perspective view
<code>R.mat</code>	a transformational matrix that is applied to the orientation of the axes
<code>par.settings</code>	plotting settings (see trellis.par.set)
<code>lattice.options</code>	lattice settings (see lattice.options)
<code>...</code>	Other arguments passed to wireframe

Value

An object of class "trellis", as output by [wireframe](#).

Examples

```
g <- data.frame(x = runif(100, 0, 60), y = runif(100,0,10),
               z = factor(rep( periods$name[1:5], each=20),
                           levels = periods$name[1:5]))
disparity_through_time(z~x*y, data = g, groups = z, aspect = c(1.5,2),
                       xlim = c(0,60), ylim = c(0,10), col.regions = "lightgreen",
                       col.point = c("red","blue"))
```

eons	<i>Eon data from the International Commission on Stratigraphy (v2013-1)</i>
------	---

Description

A dataset containing the boundary ages, abbreviations, and colors for the eons of the Geologic Time Scale. Based on The ICS International Chronostratigraphic Chart (2013), by Cohen, Finney, Gibbard, and Fan.

Usage

```
eons
```

Format

A data frame with 4 rows and 5 variables:

name eon name

max_age maximum age, in millions of years

min_age minimum age, in millions of years

abbr eon name abbreviations

color the colors for each eon, according to the Commission for the Geological Map of the World

Source

https://stratigraphy.org/ICSchart/Cohen2013_Episodes.pdf via https://paleobiodb.org/data1.2/intervals/list.txt?scale_id=1&scale_level=1

epochs	<i>Epoch data from the International Commission on Stratigraphy (v2013-1)</i>
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Description

A dataset containing the boundary ages, abbreviations, and colors for the epochs of the Geologic Time Scale. Based on The ICS International Chronostratigraphic Chart (2013), by Cohen, Finney, Gibbard, and Fan.

Usage

epochs

Format

A data frame with 34 rows and 5 variables:

name epoch name

max_age maximum age, in millions of years

min_age minimum age, in millions of years

abbr epoch name abbreviations

color the colors for each epoch, according to the Commission for the Geological Map of the World

Source

https://stratigraphy.org/ICSchart/Cohen2013_Episodes.pdf via https://paleobiodb.org/data1.2/intervals/list.txt?scale_id=1&scale_level=4

eras	<i>Era data from the International Commission on Stratigraphy (v2013-1)</i>
------	---

Description

A dataset containing the boundary ages, abbreviations, and colors for the eras of the Geologic Time Scale. Based on The ICS International Chronostratigraphic Chart (2013), by Cohen, Finney, Gibbard, and Fan.

Usage

eras

Format

A data frame with 10 rows and 5 variables:

name era name

max_age maximum age, in millions of years

min_age minimum age, in millions of years

abbr era name abbreviations

color the colors for each era, according to the Commission for the Geological Map of the World

Source

https://stratigraphy.org/ICSchart/Cohen2013_Episodes.pdf via https://paleobiodb.org/data1.2/intervals/list.txt?scale_id=1&scale_level=2

getScaleData

Get geological timescale data

Description

This function takes a name of a geological timescale and returns data for the timescale.

Usage

```
getScaleData(name)
```

Arguments

name The name of the desired timescale.

Details

Valid names include those of built-in dataframes ("periods", "epochs", "stages", "eons", or "eras") and those hosted by macrostrat (see list here: <https://macrostrat.org/api/defs/timescales?all>).

Value

A dataframe with the following columns:

name the names of the time intervals.

max_age the oldest boundaries of the time intervals, in millions of years.

min_age the youngest boundaries of the time intervals, in millions of years.

abbr either traditional abbreviations of the names of the time intervals (if they exist) or custom abbreviations created with R.

color hex color codes associated with the time intervals (if applicable).

 ggarrange2

ggarrange2

Description

Arrange multiple ggplot, grobified ggplot, or geo_scale objects on a page, aligning the plot panels, axes, and axis titles.

Usage

```
ggarrange2(
  ...,
  plots = list(...),
  layout = NULL,
  nrow = NULL,
  ncol = NULL,
  widths = NULL,
  heights = NULL,
  byrow = TRUE,
  top = NULL,
  bottom = NULL,
  left = NULL,
  right = NULL,
  padding = unit(0.5, "line"),
  margin = unit(0.5, "line"),
  clip = "on",
  draw = TRUE,
  newpage = TRUE,
  debug = FALSE,
  labels = NULL,
  label.args = list(gp = gpar(font = 4, cex = 1.2))
)
```

Arguments

...	ggplot, grobified ggplot (gtable), or geo_scale objects
plots	list of ggplot, gtable, or geo_scale objects
layout	a matrix of integers specifying where each plot should go, like mat in layout ; NA or a value less than 0 or greater than the number of plots indicates a blank plot; overrides nrow/ncol/byrow
nrow	number of rows
ncol	number of columns
widths	list of requested widths
heights	list of requested heights
byrow	logical, fill by rows

top	optional string, or grob
bottom	optional string, or grob
left	optional string, or grob
right	optional string, or grob
padding	unit of length one, margin around annotations
margin	vector of units of length 4: top, right, bottom, left (as in gtable_add_padding)
clip	argument of gtable
draw	logical: draw or return a grob
newpage	logical: draw on a new page
debug	logical, show layout with thin lines
labels	character labels used for annotation of subfigures (should be in the same order as plots)
label.args	label list of parameters for the formatting of labels

Value

gtable of aligned plots

Examples

```
library(ggplot2)
p1 <- ggplot(mtcars, aes(mpg, wt, colour = factor(cyl))) +
  geom_point()
p2 <- ggplot(mtcars, aes(mpg, wt, colour = factor(cyl))) +
  geom_point() + facet_wrap(~ cyl, ncol=2, scales = 'free') +
  guides(colour='none') +
  theme()
ggarrange2(p1, p2, widths = c(2,1), labels = c('a', 'b'))

p3 <- ggplot() +
  geom_point(aes(y = runif(1000, 0, 8), x = runif(1000, 0, 1000))) +
  scale_x_reverse() +
  coord_geo(xlim = c(1000, 0), ylim = c(0,8)) +
  theme_classic()
ggarrange2(ggarrange2(p1, p2, widths = c(2,1), draw = FALSE), p3, nrow = 2)
```

gggeo_scale

Add a geologic scale to ggplots

Description

This function takes a ggplot object and adds a geologic time scale at the specified side.

Usage

```
gggeo_scale(obj, ...)  
  
## S3 method for class 'gtable'  
gggeo_scale(  
  obj,  
  lims,  
  dat = "periods",  
  fill = NULL,  
  color = "black",  
  alpha = 1,  
  height = unit(2, "line"),  
  pos = "bottom",  
  lab = TRUE,  
  rot = 0,  
  abbrev = TRUE,  
  skip = c("Quaternary", "Holocene", "Late Pleistocene"),  
  size = 5,  
  lwd = 0.25,  
  margin = unit(0, "line"),  
  neg = FALSE,  
  bord = c("left", "right", "top", "bottom"),  
  center_end_labels = FALSE,  
  ...  
)  
  
## S3 method for class 'ggplot'  
gggeo_scale(  
  obj,  
  dat = "periods",  
  fill = NULL,  
  color = "black",  
  alpha = 1,  
  height = unit(2, "line"),  
  pos = "bottom",  
  lab = TRUE,  
  rot = 0,  
  abbrev = TRUE,  
  skip = c("Quaternary", "Holocene", "Late Pleistocene"),  
  size = 5,  
  lwd = 0.25,  
  margin = unit(0, "line"),  
  neg = FALSE,  
  bord = c("left", "right", "top", "bottom"),  
  center_end_labels = FALSE,  
  ...  
)
```

```

## S3 method for class 'geo_scale'
gggeo_scale(
  obj,
  dat = "periods",
  fill = NULL,
  color = "black",
  alpha = 1,
  height = unit(2, "line"),
  pos = "bottom",
  lab = TRUE,
  rot = 0,
  abbrev = TRUE,
  skip = c("Quaternary", "Holocene", "Late Pleistocene"),
  size = 5,
  lwd = 0.25,
  margin = unit(0, "line"),
  neg = FALSE,
  bord = c("left", "right", "top", "bottom"),
  center_end_labels = FALSE,
  ...
)

## S3 method for class 'geo_scale'
print(x, ...)

```

Arguments

obj	An object of class ggplot, gtable, or geo_scale (as produced by this function).
...	further arguments passed to <code>grid.draw</code> .
lims	The limits of the axis of the desired side of the plot. Only required if using a gtable object not created by this function.
dat	Either A) a string indicating a built-in dataframe with interval data from the ICS ("periods", "epochs", "stages", "eons", or "eras"), B) a string indicating a timescale from macrostrat (see list here: https://macrostrat.org/api/defs/timescales?all), or C) a custom dataframe of time interval boundaries (see Details).
fill	The fill color of the boxes. The default is to use the colors included in dat. If a custom dataset is provided with dat without color and without fill, a greyscale will be used. Custom fill colors can be provided with this option and will be recycled if/as necessary.
color	The outline color of the interval boxes.
alpha	The transparency of the fill colors.
height	The height (or width if pos is left or right) of the scale.
pos	Which side to add the scale to (left, right, top, or bottom). First letter may also be used.

lab	Whether to include labels.
rot	The amount of counter-clockwise rotation to add to the labels (in degrees).
abbrv	If including labels, whether to use abbreviations instead of full interval names.
skip	A vector of interval names indicating which intervals should not be labeled.
size	Label size.
lwd	Line width.
margin	The width of the margin around the returned object (can be a vector of length 4).
neg	Set this to true if your x-axis is using negative values.
bord	A vector specifying on Which sides of the scale to add borders (same options as pos).
center_end_labels	Should labels be centered within the visible range of intervals at the ends of the axis?
x	An object of class <code>geo_scale</code> .

Details

If custom data is provided (with `dat`), it should consist of at least 3 columns of data. See `data(periods)` for an example. The `name` column lists the names of each time interval. These will be used as labels if no abbreviations are provided. The `max_age` column lists the oldest boundary of each time interval. The `min_age` column lists the youngest boundary of each time interval. The `abbr` column is optional and lists abbreviations that may be used as labels. The `color` column is also optional and lists a hex color code (which can be obtained with `rgb()`) for each time interval.

Value

A `geo_scale` object. Basically a `gtable` object but with the axis limits included.

Examples

```
library(ggplot2)
# bottom scale by default
p <- ggplot() +
  geom_point(aes(y = runif(1000, 0, 8), x = runif(1000, 0, 1000))) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(1000, 0), ylim = c(0, 8), expand = FALSE) +
  theme_classic()
gggeo_scale(p)

# can specify any side of the plot
p <- ggplot() +
  geom_point(aes(x = runif(1000, 0, 8), y = runif(1000, 0, 1000))) +
  scale_y_reverse() +
  coord_cartesian(xlim = c(0, 8), ylim = c(1000, 0), expand = FALSE) +
  theme_classic()
gggeo_scale(p, pos = "left", rot = 90)
```



```

# can add multiple scales
p <- ggplot() +
  geom_point(aes(y = runif(1000, 0, 8), x = runif(1000, 0, 100))) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(100, 0), ylim = c(0, 8), expand = FALSE) +
  theme_classic()
p <- gggeo_scale(p, abbrv = FALSE)
p <- gggeo_scale(p, dat = "epochs", height = unit(4, "lines"), rot = 90, size = 2.5, abbrv = FALSE)
gggeo_scale(p, dat = "stages", height = unit(4, "lines"), rot = 90, size = 2.5, abbrv = FALSE)

# intervals on both sides for different timescales (ICS stages vs North American Land Mammal Ages)
p <- ggplot() +
  geom_point(aes(x = runif(1000, 0, 10), y = runif(1000, 0, 65))) +
  scale_y_reverse() +
  coord_cartesian(xlim = c(0, 10), ylim = c(65, 0), expand = FALSE) +
  theme_classic()
p <- gggeo_scale(p, dat = "stages", pos = "left", height = unit(4, "lines"), size = 2.5,
  abbrv = FALSE)
gggeo_scale(p, dat = "North American Land Mammal Ages", pos = "right", height = unit(4, "lines"),
  size = 2.5, abbrv = FALSE)

#can add scales to a faceted plot
#use gggeo_scale_old() if you have more than one column
df <- data.frame(x = runif(1000, 0, 541), y = runif(1000, 0, 8),
  z = sample(c(1, 2, 3, 4), 1000, TRUE))
p <- ggplot(df) +
  geom_point(aes(x, y)) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(541, 0), ylim = c(0, 8), expand = FALSE) +
  theme_classic() +
  facet_wrap(~z, ncol = 1)
gggeo_scale(p)

#can even add a scale to a phylogeny (using ggtree)

library(phytools)
library(ggtree)
tree <- pbtree(b = .03, d = .01, n=100)
p <- ggtree(tree) +
  coord_cartesian(xlim = c(-500, 0), ylim = c(-2, Ntip(tree)), expand = FALSE) +
  scale_x_continuous(breaks=seq(-500, 0, 100), labels=abs(seq(-500, 0, 100))) +
  theme_tree2()
p <- revts(p)
gggeo_scale(p, neg = TRUE)

```

gggeo_scale_old

Add a geologic scale to ggplots (old version)

Description

This function takes a ggplot object and adds a geologic time scale at the specified side.

Usage

```
gggeo_scale_old(
  gg,
  dat = "periods",
  fill = NULL,
  color = "black",
  alpha = 1,
  height = 0.05,
  gap = 0,
  pos = "bottom",
  lab = TRUE,
  rot = 0,
  abbrev = TRUE,
  skip = c("Quaternary", "Holocene", "Late Pleistocene"),
  size = 5,
  neg = FALSE
)
```

Arguments

gg	The ggplot object.
dat	Either A) a string indicating a built-in dataframe with interval data from the ICS ("periods", "epochs", "stages", "eons", or "eras"), B) a string indicating a timescale from macrostrat (see list here: https://macrostrat.org/api/defs/timescales?all), or C) a custom dataframe of time interval boundaries (see Details).
fill	The fill color of the boxes. The default is to use the colors included in dat. If a custom dataset is provided with dat without color and without fill, a greyscale will be used. Custom fill colors can be provided with this option and will be recycled if/as necessary.
color	The outline color of the interval boxes.
alpha	The transparency of the fill colors.
height	The proportional height (or width if pos is left or right) of the entire plot to use for the scale.
gap	The proportional height (or width) of the entire plot to use as a gap between the axis and the scale.
pos	Which side to add the scale to (left, right, top, or bottom). First letter may also be used.
lab	Whether to include labels.
rot	The amount of counter-clockwise rotation to add to the labels (in degrees).
abbrev	If including labels, whether to use abbreviations instead of full interval names.
skip	A vector of interval names indicating which intervals should not be labeled.
size	Label size.
neg	Set this to true if your x-axis is using negative values.

Details

If custom data is provided (with `dat`), it should consist of at least 3 columns of data. See `data(periods)` for an example. The `name` column lists the names of each time interval. These will be used as labels if no abbreviations are provided. The `max_age` column lists the oldest boundary of each time interval. The `min_age` column lists the youngest boundary of each time interval. The `abbr` column is optional and lists abbreviations that may be used as labels. The `color` column is also optional and lists a hex color code (which can be obtained with `rgb()`) for each time interval.

Value

A ggplot object.

Examples

```
library(ggplot2)
# bottom scale by default
p <- ggplot() +
  geom_point(aes(y = runif(1000, .5, 8), x = runif(1000, 0, 1000))) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(0, 1000), ylim = c(0,8), expand = FALSE) +
  theme_classic()
gggeo_scale_old(p)

# can specify any side of the plot
p <- ggplot() +
  geom_point(aes(x = runif(1000, .5, 8), y = runif(1000, 0, 1000))) +
  scale_y_reverse() +
  coord_cartesian(xlim = c(0, 8), ylim = c(0,1000), expand = FALSE) +
  theme_classic()
gggeo_scale_old(p, pos = "left", rot = 90)

# can add multiple scales
p <- ggplot() +
  geom_point(aes(y = runif(1000, 1, 8), x = runif(1000, 0, 1000))) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(0, 100), ylim = c(0,8), expand = FALSE) +
  theme_classic()
p <- gggeo_scale_old(p, height = .03, abbrv = FALSE)
p <- gggeo_scale_old(p, dat = "epochs", gap = .03, height = .1, rot = 90, size = 2.5, abbrv = FALSE)

# intervals on both sides for different timescales (ICS stages vs North American Land Mammal Ages)
p <- ggplot() +
  geom_point(aes(x = runif(1000, 1, 9), y = runif(1000, 0, 65))) +
  scale_y_reverse() +
  coord_cartesian(xlim = c(0, 10), ylim = c(0,65), expand = FALSE) +
  theme_classic()
p <- gggeo_scale_old(p, dat = "stages", pos = "left", height = .1, size = 2.5, abbrv = FALSE)
gggeo_scale_old(p, dat = "North American Land Mammal Ages", pos = "right", height = .1, size = 2.5,
  abbrv = FALSE)

#can add scales to a faceted plot
```

```

df <- data.frame(x = runif(1000,0,541), y = runif(1000,.5,8), z = sample(c(1,2,3,4), 1000, TRUE))
p <- ggplot(df) +
  geom_point(aes(x, y)) +
  scale_x_reverse() +
  coord_cartesian(xlim = c(0, 541), ylim = c(0,8), expand = FALSE) +
  theme_classic() +
  facet_wrap(~z, nrow = 2)
gggeo_scale_old(p)

#can even add a scale to a phylogeny (using ggtree)
library(phytools)
library(ggtree)
tree <- pbtree(b = .03, d = .01, n=100)
p <- ggtree(tree) +
  coord_cartesian(xlim = c(0,-500), ylim = c(-10,Ntip(tree)), expand = FALSE) +
  scale_x_continuous(breaks=seq(-500,0,100), labels=abs(seq(-500,0,100))) +
  theme_tree2()
p <- revts(p)
gggeo_scale_old(p, neg = TRUE)

```

gtable_frame2

gtable_frame2

Description

Reformat the gtable associated with a ggplot object into a 7x7 gtable where the central cell corresponds to the plot panel(s), the rectangle of cells around that corresponds to the axes, and the rectangle of cells around that corresponds to the axis titles.

Usage

```

gtable_frame2(
  g,
  width = unit(1, "null"),
  height = unit(1, "null"),
  debug = FALSE
)

```

Arguments

g	gtable
width	requested width
height	requested height
debug	logical draw gtable cells

Value

7x7 gtable wrapping the plot

Examples

```

library(grid)
library(gridExtra)
library(ggplot2)
p1 <- ggplot(mtcars, aes(mpg, wt, colour = factor(cyl))) +
  geom_point()

p2 <- ggplot(mtcars, aes(mpg, wt, colour = factor(cyl))) +
  geom_point() + facet_wrap(~ cyl, ncol=2, scales = 'free') +
  guides(colour='none') +
  theme()

p3 <- ggplot(mtcars, aes(mpg, wt, colour = factor(cyl))) +
  geom_point() + facet_grid(. ~ cyl, scales = 'free')

g1 <- ggplotGrob(p1);
g2 <- ggplotGrob(p2);
g3 <- ggplotGrob(p3);
fg1 <- gtable_frame2(g1)
fg2 <- gtable_frame2(g2)
fg12 <- gtable_frame2(gtable_rbind(fg1,fg2), width=unit(2,'null'), height=unit(1,'null'))
fg3 <- gtable_frame2(g3, width=unit(1,'null'), height=unit(1,'null'))
grid.newpage()
combined <- gtable_cbind(fg12, fg3)
grid.draw(combined)

```

panel.disparity

Combined wireframe and cloud panel

Description

Plots the provided data on 2-D surfaces within a 3-D framework. See [disparity_through_time](#).

Usage

```
panel.disparity(x, y, z, groups, subscripts, ...)
```

Arguments

x, y, z, groups, subscripts, ...
 Same as for [panel.cloud](#)

Value

No return value, plots the results of both [panel.cloud](#) and [panel.wireframe](#).

periods	<i>Period data from the International Commission on Stratigraphy (v2013-1)</i>
---------	--

Description

A dataset containing the boundary ages, abbreviations, and colors for the periods of the Geologic Time Scale. Based on The ICS International Chronostratigraphic Chart (2013), by Cohen, Finney, Gibbard, and Fan.

Usage

periods

Format

A data frame with 22 rows and 5 variables:

name period name

max_age maximum age, in millions of years

min_age minimum age, in millions of years

abbr period name abbreviations

color the colors for each period, according to the Commission for the Geological Map of the World

Source

https://stratigraphy.org/ICSchart/Cohen2013_Episodes.pdf via https://paleobiodb.org/data1.2/intervals/list.txt?scale_id=1&scale_level=3

stages	<i>Stage data from the International Commission on Stratigraphy (v2013-1)</i>
--------	---

Description

A dataset containing the boundary ages, abbreviations, and colors for the stages of the Geologic Time Scale. Based on The ICS International Chronostratigraphic Chart (2013), by Cohen, Finney, Gibbard, and Fan.

Usage

stages

Format

A data frame with 100 rows and 5 variables:

name stage name

max_age maximum age, in millions of years

min_age minimum age, in millions of years

abbr stage name abbreviations

color the colors for each stage, according to the Commission for the Geological Map of the World

Source

https://stratigraphy.org/ICSchart/Cohen2013_Episodes.pdf via https://paleobiodb.org/data1.2/intervals/list.txt?scale_id=1&scale_level=5

Index

* datasets

- coord_geo, [2](#)
- coord_trans_flip, [5](#)
- coord_trans_xy, [6](#)
- eons, [9](#)
- epochs, [10](#)
- eras, [10](#)
- periods, [22](#)
- stages, [22](#)

coord_flip, [5](#)

coord_geo, [2](#)

coord_trans, [2](#), [3](#), [5](#), [6](#)

coord_trans_flip, [5](#)

coord_trans_xy, [6](#)

CoordGeo (coord_geo), [2](#)

CoordTransFlip (coord_trans_flip), [5](#)

CoordTransXY (coord_trans_xy), [6](#)

disparity_through_time, [7](#), [21](#)

eons, [9](#)

epochs, [10](#)

eras, [10](#)

geom_fit_text, [3](#), [4](#)

geom_point, [6](#)

geom_polygon, [6](#)

geom_rect, [6](#)

geom_text, [3](#)

getScaleData, [11](#)

ggarrange2, [12](#)

gggeo_scale, [13](#)

gggeo_scale_old, [17](#)

grid.draw, [15](#)

gtable_add_padding, [13](#)

gtable_frame2, [20](#)

lattice.options, [8](#)

layout, [12](#)

levelplot, [8](#)

linear_trans, [6](#)

panel.cloud, [7](#), [21](#)

panel.disparity, [21](#)

panel.wireframe, [21](#)

periods, [22](#)

print.geo_scale (gggeo_scale), [13](#)

stages, [22](#)

trans_new, [6](#)

trellis.par.set, [8](#)

wireframe, [7](#), [8](#)

xyplot, [8](#)