

Package ‘spiralize’

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Type Package

Title Visualize Data on Spirals

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Suggests ComplexHeatmap, knitr, rmarkdown, grImport, grImport2, jpeg, png, tiff, ape, cranlogs, cowplot, dendextend, bezier, magick

Description It visualizes data along an Archimedean spiral <https://en.wikipedia.org/wiki/Archimedean_spiral>, makes so-called spiral graph or spiral chart.
It has two major advantages for visualization: 1. It is able to visualize data with very long axis with high resolution. 2. It is efficient for time series data to reveal periodic patterns.

VignetteBuilder knitr

URL <https://github.com/jokergoo/spiralize>

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NeedsCompilation no

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cartesian_to_xy	<i>Convert canvas coordinates to the data coordinates</i>
-----------------	---

Description

Convert canvas coordinates to the data coordinates

Usage

```
cartesian_to_xy(x, y, track_index = current_track_index())
```

Arguments

x	X-locations of the data points in canvas coordinates.
y	Y-locations of the data points in canvas coordinates.
track_index	Index of the track.

Details

The data points are assigned to the nearest inner loops. Denote the a data point has a coordinate (r, theta) in the polar coordinate system, r_k and r_{k+1} are the radius of the two loops at $\theta + 2\pi \cdot a$ and $\theta + 2\pi \cdot (a+1)$ that below and above the data point, the data point is assigned to the loop k.

Value

A data frame with two columns: x and y.

Examples

```
x = runif(100, -5, 5)
y = runif(100, -5, 5)
spiral_initialize()
spiral_track()
df = cartesian_to_xy(x, y)
# directly draw in the viewport
grid.points(x, y, default.units="native")
# check whether the converted xy are correct (should overlap to the previous points)
spiral_points(df$x, df$y, pch = 16)
```

current_spiral	<i>Get current spiral object</i>
----------------	----------------------------------

Description

Get current spiral object

Usage

```
current_spiral()
```

Details

The returned value is an object of spiral reference class. The following methods might be useful:

`$curve()`: It returns the radius for given angles (in radians).

`$spiral_length()`: It returns the length of the spiral (from the origin) for a given angle (in radians), thus if you want to get the length of a spiral segment, it will be `spiral$spiral_length(theta2) - spiral$spiral_length(theta1)` where `spiral` is the spiral object.

Also there are the following meta-data for the current spiral (assume the object is named `s`):

`s$xlim`: Data range.

`s$xrange`: `s$xlim[2] - s$xlim[1]`

`s$theta_lim`: The corresponding range of theta

`s$theta_range`: `s$theta_lim[2] - s$theta_lim[1]`

`s$spiral_length_lim`: The corresponding range of spiral length

`s$spiral_length_range`: `s$spiral_length_lim[2] - s$spiral_length_lim[1]`

`s$max_radius`: Radius at `s$theta_lim[2]`

Value

A spiral object.

Examples

```
spiral_initialize()
s = current_spiral()
s$curve(2*pi*2)
s$spiral_length(2*pi*2)
```

current_spiral_vp *Viewport name of current spiral*

Description

Viewport name of current spiral

Usage

`current_spiral_vp()`

Value

A string of the viewport name.

Examples

```
# There is no example  
NULL
```

current_track_index *Current track index*

Description

Current track index

Usage

`current_track_index()`

Value

An integer of the index of the current track.

Examples

```
# There is no example  
NULL
```

get_track_data	<i>Get meta-data of a track</i>
----------------	---------------------------------

Description

Get meta-data of a track

Usage

```
get_track_data(field, track_index = current_track_index())
```

Arguments

field	Name, see Details section.
track_index	Which track?

Details

There are following fields that can be retrieved for a given track:

- ymin Lower boundary of the data.
- ymax Upper boundary of the data.
- ycenter $(ymin + ymax)/2$
- ylim $c(ylim, ymax)$
- yrange $ymax - ymin$
- height Height of the track, measured as the fraction of the distance between two neighbouring circles.

It is more suggested to directly use [TRACK_META](#) to retrieve meta data for the current track.

Value

A numeric value of the corresponding field.

Examples

```
# There is no example  
NULL
```

horizon_legend	<i>Legend for the horizon chart</i>
----------------	-------------------------------------

Description

Legend for the horizon chart

Usage

```
horizon_legend(lt, title = "", format = "%.2f",  
              template = "[{x1}, {x2}]", ...)
```

Arguments

lt	The object returned by spiral_horizon .
title	Title of the legend.
format	Number format of the legend labels.
template	Template to construct the labels.
...	Pass to Legend .

Value

A [Legend](#) object.

Examples

```
# There is no example  
NULL
```

is_in_track	<i>Test whether points are in a track</i>
-------------	---

Description

Test whether points are in a track

Usage

```
is_in_track(x, y, track_index = current_track_index())
```

Arguments

x	X-location of data points.
y	Y-location of data points.
track_index	Index of track.

Value

A logical vector.

Examples

```
# There is no example
NULL
```

names.TRACK_META	<i>Names of all supported meta data</i>
------------------	---

Description

Names of all supported meta data

Usage

```
## S3 method for class 'TRACK_META'
names(x)
```

Arguments

x Always use TRACK_META.

Value

A vector of characters.

Examples

```
names(TRACK_META)
```

n_tracks	<i>Number of tracks</i>
----------	-------------------------

Description

Number of tracks

Usage

```
n_tracks()
```

Value

An integer of the number of available tracks.

Examples

```
# There is no example
NULL
```

phylo_to_dendrogram *Convert a phylo object to a dendrogram object*

Description

Convert a phylo object to a dendrogram object

Usage

```
phylo_to_dendrogram(obj, log = FALSE)
```

Arguments

obj	A phylo object.
log	Whether the height of the phylogenetic tree should be log-transformed ($\log_{10}(x + 1)$).

Details

The motivation is that phylogenetic tree may contain polytomies, which means at a certain node, there are more than two children branches. Available tools that do the conversion only support binary trees.

The returned dendrogram object is not in its standard format which means it can not be properly drawn by the `plot.dendrogram` function. However, you can still apply `dendextend::cutree` to the returned dendrogram object with no problem and the dendrogram can be properly drawn with the `ComplexHeatmap` package.

Value

A dendrogram object.

Examples

```
require(ape)
data(bird.families)
d = phylo_to_dendrogram(bird.families)

require(ComplexHeatmap)
grid.dendrogram(d, test = TRUE)
```

`polar_to_cartesian` *Convert polar coordinates to catersian coordinates*

Description

Convert polar coordinates to catersian coordinates

Usage

```
polar_to_cartesian(theta, r)
```

Arguments

<code>theta</code>	Angles, in radians.
<code>r</code>	Radius.

Value

A data frame with two columns: x abd y.

Examples

```
# There is no example  
NULL
```

`print.TRACK_META` *Print TRACK_META*

Description

Print TRACK_META

Usage

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

Arguments

<code>x</code>	The TRACK_META object.
<code>...</code>	Additional parameters.

Value

No value is returned.

Examples

```
# There is no example  
NULL
```

set_current_track	<i>Set current track</i>
-------------------	--------------------------

Description

Set current track

Usage

```
set_current_track(track_index)
```

Arguments

track_index The index of the track.

Value

No value is returned.

Examples

```
# There is no example  
NULL
```

solve_theta_from_spiral_length	<i>Get theta from given spiral lengths</i>
--------------------------------	--

Description

Get theta from given spiral lengths

Usage

```
solve_theta_from_spiral_length(len, interval = NULL, offset = 0)
```

Arguments

len	A vector of spiral lengths.
interval	Interval to search for the solution.
offset	Offset of the spiral. In the general form: $r = a + r \cdot \theta$, offset is the value of a .

Details

The length of the spiral has a complicated form, see https://downloads.imagej.net/fiji/snapshots/arc_length.pdf. Let's say the form is $l = f(\theta)$, `solve_theta_from_spiral_length` tries to find θ by a known l . It uses `uniroot` to search solutions.

Value

The θ value.

Examples

```
spiral_initialize()
s = current_spiral()
theta = pi*seq(2, 3, length = 10)
len = s$spiral_length(theta)
solve_theta_from_spiral_length(len) # should be very similar as theta
```

spiral_arrow

Draw arrows in the spiral direction

Description

Draw arrows in the spiral direction

Usage

```
spiral_arrow(
  x1, x2,
  y = get_track_data("ycenter", track_index),
  width = get_track_data("yrange", track_index)/3,
  arrow_head_length = unit(4, "mm"),
  arrow_head_width = width*2,
  arrow_position = c("end", "start"),
  tail = c("normal", "point"),
  gp = gpar(),
  track_index = current_track_index())
```

Arguments

x1	Start of the arrow.
x2	End of the arrow.
y	Y-location of the arrow.
width	Width of the arrow. The value can be the one measured in the data coordinates or a <code>unit</code> object.
arrow_head_length	Length of the arrow head.
arrow_head_width	Width of the arrow head.
arrow_position	Position of the arrow. If the value is "end", then the arrow head is drawn at x = x2. If the value is "start", then the arrow head is drawn at x = x1.
tail	The shape of the arrow tail.
gp	Graphics parameters.
track_index	Index of the track.

Value

No value is returned.

See Also

Note `spiral_segments` also supports drawing line-based arrows.

Examples

```
spiral_initialize()
spiral_track()
spiral_arrow(0.3, 0.6, gp = gpar(fill = "red"))
spiral_arrow(0.8, 0.9, gp = gpar(fill = "blue"), tail = "point", arrow_position = "start")
```

spiral_axis

Draw axis along the spiral

Description

Draw axis along the spiral

Usage

```
spiral_axis(h = c("top", "bottom"), at = NULL, major_at = at,
  labels = TRUE, curved_labels = FALSE, minor_ticks = 4,
  major_ticks_length = unit(4, "bigpts"), minor_ticks_length = unit(2, "bigpts"),
  ticks_gp = gpar(), labels_gp = gpar(fontsize = 6),
  track_index = current_track_index())
```

Arguments

h	Position of the axis. The value can be a character of "top" or "bottom".
at	Breaks points on axis.
major_at	Breaks points on axis. It is the same as at.
labels	The corresponding labels for the break points.
curved_labels	Whether are the labels are curved?
minor_ticks	Number of minor ticks.
major_ticks_length	Length of the major ticks. The value should be a <code>unit</code> object.
minor_ticks_length	Length of the minor ticks. The value should be a <code>unit</code> object.
ticks_gp	Graphics parameters for the ticks.
labels_gp	Graphics parameters for the labels.
track_index	Index of the track.

Value

No value is returned.

Examples

```

spiral_initialize(); spiral_track()
spiral_axis()

# if the spiral is intepolated by the curve length
spiral_initialize(scale_by = "curve_length"); spiral_track()
spiral_axis()

spiral_initialize(xlim = c(0, 360*4), start = 360, end = 360*5); spiral_track()
spiral_axis(major_at = seq(0, 360*4, by = 30))

spiral_initialize(xlim = c(0, 12*4), start = 360, end = 360*5); spiral_track()
spiral_axis(major_at = seq(0, 12*4, by = 1), labels = c("", rep(month.name, 4)))

```

spiral_bars

Add bars to a track

Description

Add bars to a track

Usage

```

spiral_bars(pos, value, baseline = get_track_data("ymin", track_index),
            bar_width = min(diff(pos)), gp = gpar(), track_index = current_track_index())

```

Arguments

pos	X-locations of the center of bars.
value	Height of bars. The value can be a simple numeric vector, or a matrix.
baseline	Baseline of the bars. Note it only works when value is a simple vector.
bar_width	Width of bars.
gp	Graphical parameters.
track_index	Index of the track.

Value

No value is returned.

Examples

```
x = seq(1, 1000, by = 1) - 0.5
y = runif(1000)
spiral_initialize(xlim = c(0, 1000))
spiral_track(height = 0.8)
spiral_bars(x, y)

# a three-column matrix
y = matrix(runif(3*1000), ncol = 3)
y = y/rowSums(y)
spiral_initialize(xlim = c(0, 1000))
spiral_track(height = 0.8)
spiral_bars(x, y, gp = gpar(fill = 2:4, col = NA))
```

spiral_clear	<i>Clear the spiral curve</i>
--------------	-------------------------------

Description

Clear the spiral curve

Usage

```
spiral_clear(check_vp = TRUE)
```

Arguments

check_vp	Whether to check the viewport.
----------	--------------------------------

Details

It basically sets the internally spiral object to NULL, and reset all the global options.

Value

No value is returned.

Examples

```
# There is no example
NULL
```

spiral_dendrogram	<i>Draw dendrogram</i>
-------------------	------------------------

Description

Draw dendrogram

Usage

```
spiral_dendrogram(dend, gp = gpar(), track_index = current_track_index())
```

Arguments

dend	A stats::dendrogram object.
gp	Graphics parameters of the dendrogram edges.
track_index	Index of the track.

Details

Note the dendrogram edges can be rendered with the [dendextend](#) package.

Value

Height of the dendrogram.

Examples

```
k = 500
dend = as.dendrogram(hclust(dist(runif(k))))
spiral_initialize(xlim = c(0, k), start = 360, end = 360*3)
spiral_track(height = 0.8, background_gp = gpar(fill = "#EEEEEE", col = NA))

require(dendextend)
dend = color_branches(dend, k = 4)
spiral_initialize(xlim = c(0, k), start = 360, end = 360*3)
spiral_track(height = 0.8, background_gp = gpar(fill = "#EEEEEE", col = NA))
```

spiral_highlight	<i>Highlight a section of the spiral</i>
------------------	--

Description

Highlight a section of the spiral

Usage

```
spiral_highlight(x1, x2, type = c("rect", "line"), padding = unit(1, "mm"),
  line_side = c("inside", "outside"), line_width = unit(1, "pt"),
  gp = gpar(fill = "red"), track_index = current_track_index())
```

Arguments

x1	Start location of the highlighted section.
x2	End location of the highlighted section.
type	Type of the highlighting. "rect" means drawing transparent rectangles covering the whole track. "line" means drawing annotation lines on top of the track or at the bottom of it.
padding	When the highlight type is "rect", it controls the padding of the highlighted region. The value should be a unit object or a numeric value which is the fraction of the length of the highlighted section. The length can be one or two. Note it only extends in the radial direction.
line_side	If the highlight type is "line", it controls which side of the track to draw the lines.
line_width	Width of the annotation line. Value should be a unit object.
gp	Graphics parameters.
track_index	Index of the track.

Value

No value is returned.

Examples

```
spiral_initialize(); spiral_track()
spiral_highlight(0.4, 0.6)
spiral_highlight(0.1, 0.2, type = "line", gp = gpar(col = "blue"))
spiral_highlight(0.7, 0.8, type = "line", line_side = "outside")
```

spiral_highlight_by_sector
Highlight a sector

Description

Highlight a sector

Usage

```
spiral_highlight_by_sector(x1, x2, x3 = NULL, x4 = NULL, padding = unit(1, "mm"),
  gp = gpar(fill = "red"))
```

Arguments

x1	Start location which determines the start of the sector.
x2	End location which determines the end of the sector. Note x2 should be larger than x1 and the angular difference between x1 and x2 should be smaller than a circle.
x3	Start location which determines the start of the sector on the upper border.
x4	End location which determines the end of the sector on the upper border.
padding	It controls the radial extension of the sector. The value should be a <code>unit</code> object with length one or two.
gp	Graphics parameters.

Details

x1 and x2 determine the position of the highlighted sector. If x3 and x4 are not set, the sector extends until the most outside loop. If x3 and x4 are set, they determine the outer border of the sector. In this case, if x3 and x4 are set, x3 should be larger than x2.

Value

No value is returned.

Examples

```
spiral_initialize(xlim = c(0, 360*4), start = 360, end = 360*5)
spiral_track()
spiral_axis()
spiral_highlight_by_sector(36, 72)
spiral_highlight_by_sector(648, 684)
spiral_highlight_by_sector(216, 252, 936, 972, gp = gpar(fill = "blue"))
```

spiral_horizon *Draw horizon chart along the spiral*

Description

Draw horizon chart along the spiral

Usage

```
spiral_horizon(x, y, y_max = max(abs(y)), n_slices = 4, slice_size,
               pos_fill = "#D73027", neg_fill = "#313695",
               use_bars = FALSE, bar_width = min(diff(x)),
               negative_from_top = FALSE, track_index = current_track_index())
```

Arguments

x	X-locations of the data points.
y	Y-locations of the data points.
y_max	Maximal absolute value on y-axis.
n_slices	Number of slices.
slice_size	Size of the slices. The final number of sizes is <code>ceiling(max(abs(y))/slice_size)</code> .
pos_fill	Colors for positive values.
neg_fill	Colors for negative values.
use_bars	Whether to use bars?
bar_width	Width of bars.
negative_from_top	Should negative distribution be drawn from the top?
track_index	Index of the track.

Details

Since the track height is very small in the spiral, horizon chart visualization is an efficient way to visualize distribution-like graphics.

Value

A list of the following objects:

- a color mapping function for colors.
- a vector of intervals that split the data.

Examples

```
df = readRDS(system.file("extdata", "global_temperature.rds", package = "spiralize"))
df = df[df$Source == "GCAG", ]
spiral_initialize_by_time(xlim = range(df$Date), unit_on_axis = "months", period = "year",
  period_per_loop = 20, polar_lines_by = 360/20,
  vp_param = list(x = unit(0, "npc"), just = "left"))
spiral_track()
spiral_horizon(df$Date, df$Mean, use_bar = TRUE)
```

spiral_info	<i>Information of the current spiral</i>
-------------	--

Description

Information of the current spiral

Usage

```
spiral_info()
```

Details

It prints information of the current spiral.

Value

No value is returned.

Examples

```
# There is no example
NULL
```

spiral_initialize	<i>Initialize the spiral</i>
-------------------	------------------------------

Description

Initialize the spiral

Usage

```
spiral_initialize(xlim = c(0, 1), start = 360, end = 360*5,
  scale_by = c("angle", "curve_length"), period = NULL,
  clockwise = FALSE, flip = c("none", "vertical", "horizontal", "both"),
  reverse = FALSE, polar_lines = scale_by == "angle", polar_lines_by = 30,
  polar_lines_gp = gpar(col = "#808080", lty = 3),
  padding = unit(5, "mm"), newpage = TRUE, vp_param = list())
```

Arguments

<code>xlim</code>	Range on x-locations.
<code>start</code>	Start of the spiral, in degree. <code>start</code> and <code>end</code> should be positive and <code>start</code> should be smaller than <code>end</code> .
<code>end</code>	End of the spiral, in degree.
<code>scale_by</code>	How scales on x-axis are equally interpolated? The values can be one of "angle" and "curve_length". If the value is "angle", equal angle difference corresponds to equal difference of data. In this case, in outer loops, the scales are longer than in the inner loops, although the difference on the data are the same. If the value is "curve_length", equal curve length difference corresponds to the equal difference of the data.
<code>period</code>	Under "angle" mode, the number of loops can also be controlled by argument <code>period</code> which controls the length of data a spiral loop corresponds to. Note in this case, argument <code>end</code> is ignored and the value for <code>end</code> is internally recalculated.
<code>clockwise</code>	Whether the curve is in a closewise direction. If it is set to TRUE, argument <code>flip</code> and <code>reverse</code> are ignored.
<code>flip</code>	How to flip the spiral? By default, the spiral starts from the origin of the coordinate and grows reverseclockwisely. The argument controls the growing direction of the spiral.
<code>reverse</code>	By default, the most inside of the spiral corresponds to the lower boundary of x-location. Setting the value to FALSE can reverse the direction.
<code>polar_lines</code>	Whether draw the polar guiding lines.
<code>polar_lines_by</code>	Increment of the polar lines. Measured in degree. The value can also be a vector that defines where to add polar lines.
<code>polar_lines_gp</code>	Graphics parameters for the polar lines.
<code>padding</code>	Padding of the plotting region. The value can be a unit of length of one to two.
<code>newpage</code>	Whether to apply grid.newpage before making the plot?
<code>vp_param</code>	A list of parameters sent to viewport .

Value

No value is returned.

Examples

```

spiral_initialize(); spiral_track()
spiral_initialize(start = 180, end = 360+180); spiral_track()
spiral_initialize(flip = "vertical"); spiral_track()
spiral_initialize(flip = "horizontal"); spiral_track()
spiral_initialize(flip = "both"); spiral_track()
spiral_initialize(); spiral_track(); spiral_axis()
spiral_initialize(scale_by = "curve_length"); spiral_track(); spiral_axis()

# the following example shows the difference of `scale_by` more clearly:
make_plot = function(scale_by) {
  n = 100
  require(circlize)
  col = circlize::colorRamp2(c(0, 0.5, 1), c("blue", "white", "red"))
  spiral_initialize(xlim = c(0, n), scale_by = scale_by)
  spiral_track(height = 0.9)

  x = runif(n)
  spiral_rect(1:n - 1, 0, 1:n, 1, gp = gpar(fill = col(x), col = NA))
}
make_plot("angle")
make_plot("curve_length")

```

```
spiral_initialize_by_gcoor
```

Initialize the spiral with genomic coordinates

Description

Initialize the spiral with genomic coordinates

Usage

```
spiral_initialize_by_gcoor(xlim, scale_by = "curve_length", ...)
```

Arguments

<code>xlim</code>	Range of the genomic coordinates.
<code>scale_by</code>	For genomic plot, axis is linearly scaled by the curve length.
<code>...</code>	All pass to spiral_initialize .

Details

It is basically the same as [spiral_initialize](#). The only difference is the axis labels are automatically formatted for genomic coordinates.

Value

No value is returned.

Examples

```
spiral_initialize_by_gcoor(c(0, 1000000000))
spiral_track()
spiral_axis()
```

```
spiral_initialize_by_time
```

Initialize the spiral from time objects

Description

Initialize the spiral from time objects

Usage

```
spiral_initialize_by_time(xlim, start = NULL, end = NULL,
  unit_on_axis = c("days", "months", "weeks", "hours", "mins", "secs"),
  period = c("years", "months", "weeks", "days", "hours", "mins"),
  normalize_year = FALSE, period_per_loop = 1, polar_lines_by = NULL,
  verbose = TRUE, ...)
```

Arguments

<code>xlim</code>	Range of the time. The value can be time object such as Date, POSIXlt or POSIXct. The value can also be characters and it is converted to time objects automatically.
<code>start</code>	Start of the spiral, in degrees. By default it is automatically calculated.
<code>end</code>	End of the spiral, in degrees. By default it is automatically calculated.
<code>unit_on_axis</code>	Units on the axis.
<code>period</code>	Which period to use?
<code>normalize_year</code>	Whether to enforce one loop to represent a complete year?
<code>period_per_loop</code>	How many periods to put in a loop?
<code>polar_lines_by</code>	By default different value of <code>polar_lines_by</code> is set for different period. E.g. 360/7 is set if period is "weeks" or 360/24 is set if period is set to "hours". When period is year and <code>unit_on_axis</code> is day, the proportion of sectors by polar lines corresponds to the proportion of month days in a year.
<code>verbose</code>	Whether to print messages?
<code>...</code>	All pass to spiral_initialize .

Details

"start" and "end" are automatically calculated for different "unit_on_axis" and "period". For example, if "unit_on_axis" is "days" and "period" is "years", then the first day of each each year is always put on $\theta = 0 + 2\pi k$ where k is the index of loops.

Value

No value is returned.

Examples

```
spiral_initialize_by_time(xlim = c("2014-01-01", "2021-06-17"))
spiral_track(height = 0.6)
spiral_axis()
```

```
spiral_initialize_by_time(xlim = c("2021-01-01 00:00:00", "2021-01-05 00:00:00"))
spiral_track(height = 0.6)
spiral_axis()
```

```
spiral_initialize_by_time(xlim = c("2021-01-01 00:00:00", "2021-01-01 00:10:00"),
  unit_on_axis = "secs", period = "mins")
spiral_track(height = 0.6)
spiral_axis()
```

 spiral_lines

Add lines to a track

Description

Add lines to a track

Usage

```
spiral_lines(x, y, type = "l", gp = gpar(),
  baseline = "bottom", area = FALSE, track_index = current_track_index())
```

Arguments

x	X-locations of the data points.
y	Y-locations of the data points.
type	Type of the line. Value should be one of "l" and "h". When the value is "h", vertical lines (or radial lines if you consider the polar coordinates) relative to the baseline will be drawn.
gp	Graphical parameters.
baseline	Baseline used when type is "l" or area is TRUE.
area	Whether to draw the area under the lines? Note <code>gpar(fill = ...)</code> controls the filled of the areas.
track_index	Index of the track.

Value

No value is returned.

Examples

```

x = sort(runif(1000))
y = runif(1000)
spiral_initialize()
spiral_track()
spiral_lines(x, y)

spiral_initialize()
spiral_track()
spiral_lines(x, y, type = "h")

spiral_initialize()
spiral_track()
spiral_lines(x, y, area = TRUE, gp = gpar(fill = "red", col = NA))

```

spiral_opt

Global options

Description

Global options

Usage

```
spiral_opt(..., RESET = FALSE, READ.ONLY = NULL, LOCAL = FALSE, ADD = FALSE)
```

Arguments

...	Arguments for the parameters, see "details" section.
RESET	Whether to reset to default values.
READ.ONLY	Please ignore.
LOCAL	Please ignore.
ADD	Please ignore.

Details

There are following global parameters:

`min_segment_len` Minimal length of the segment that partitions a curve.

To access the value of an option: `spiral_opt$name` where `name` is the name of the option. To set a new value for an option: `spiral_opt$name = new_value`.

Value

A list of options.

Examples

```
# There is no example
NULL
```

spiral_phylo	<i>Draw phylogenetic tree</i>
--------------	-------------------------------

Description

Draw phylogenetic tree

Usage

```
spiral_phylo(obj, gp = gpar(), log = FALSE, reverse = FALSE,
             group = NULL, group_col = NULL, track_index = current_track_index())
```

Arguments

obj	A phylo object.
gp	Graphics parameters of the tree edges.
log	Whether the height of the tree should be log-transformed ($\log_{10}(x + 1)$)?
reverse	Whether the tree should be reversed?
group	A categorical variable for splitting the tree.
group_col	A named vector which contains group colors.
track_index	Index of the track.

Value

Height of the phylogenetic tree.

Examples

```
require(ape)
data(bird.families)
n = length(bird.families$tip.label)
spiral_initialize(xlim = c(0, n), start = 360, end = 360*3)
spiral_track(height = 0.8)
spiral_phylo(bird.families)
```

spiral_points *Add points to a track*

Description

Add points to a track

Usage

```
spiral_points(x, y, pch = 1, size = unit(0.4, "char"), gp = gpar(),  
             track_index = current_track_index())
```

Arguments

x	X-locations of the data points.
y	Y-locations of the data points.
pch	Point type.
size	Size of the points. Value should be a unit object.
gp	Graphical parameters.
track_index	Index of the track.

Value

No value is returned.

Examples

```
spiral_initialize()  
spiral_track()  
spiral_points(x = runif(1000), y = runif(1000))
```

spiral_polygon *Add polygons to a track*

Description

Add polygons to a track

Usage

```
spiral_polygon(x, y, id = NULL, gp = gpar(), track_index = current_track_index())
```

Arguments

x	X-locations of the data points.
y	Y-locations of the data points.
id	A numeric vector used to separate locations in x and y into multiple polygons.
gp	Graphical parameters.
track_index	Index of the track.

Details

Note the polygon must be closed, which means, the last data point should overlap to the first one.

Value

No value is returned.

Examples

```
# There is no example
NULL
```

spiral_raster	<i>Add image to a track</i>
---------------	-----------------------------

Description

Add image to a track

Usage

```
spiral_raster(x, y, image, width = NULL, height = NULL,
  facing = c("downward", "inside", "outside", "curved_inside", "curved_outside"),
  nice_facing = FALSE, scaling = 1, track_index = current_track_index())
```

Arguments

x	X-locations of the center of the image.
y	Y-locations of the center of the image.
image	A vector of file paths of images. The format of the image is inferred from the suffix name of the image file. NA value or empty string means no image to drawn. Supported formats are png/svg/pdf/eps/jpeg/jpg/tiff.
width	Width of the image. See Details.
height	Height of the image. See Details.
facing	Facing of the image.
nice_facing	Whether to adjust the facing.
scaling	Scaling factor when facing is set to "curved_inside" or "curved_outside".
track_index	Index of the track.

Details

When facing is set to one of "downward", "inside" and "outside", both of width and height should be `unit` objects. It is suggested to only set one of width and height, the other dimension will be automatically calculated from the aspect ratio of the image.

When facing is set to one of "curved_inside" and "curved_outside", the value can also be numeric, which are the values measured in the data coordinates. Note when the segment in the spiral that corresponds to width is very long, drawing the curved image will be very slow because each pixel is actually treated as a single rectangle.

Value

No value is returned.

Examples

```
image = system.file("extdata", "Rlogo.png", package = "circlize")
x = seq(0.1, 0.9, length = 10)

spiral_initialize()
spiral_track()
spiral_raster(x, 0.5, image)

spiral_initialize()
spiral_track()
spiral_raster(x, 0.5, image, facing = "inside")
```

<code>spiral_rect</code>	<i>Add rectangles to a track</i>
--------------------------	----------------------------------

Description

Add rectangles to a track

Usage

```
spiral_rect(xleft, ybottom, xright, ytop, gp = gpar(),
            track_index = current_track_index())
```

Arguments

<code>xleft</code>	X-locations of the left bottom of the rectangles.
<code>ybottom</code>	Y-locations of the left bottom of the rectangles.
<code>xright</code>	X-locations of the right top of the rectangles.
<code>ytop</code>	Y-locations of the right top of the rectangles.
<code>gp</code>	Graphical parameters.
<code>track_index</code>	Index of the track.

Value

No value is returned.

Examples

```
# to simulate heatmap
n = 1000
require(circlize)
col = circlize::colorRamp2(c(0, 0.5, 1), c("blue", "white", "red"))
spiral_initialize(xlim = c(0, n))
spiral_track(height = 0.9)

x1 = runif(n)
spiral_rect(1:n - 1, 0, 1:n, 0.5, gp = gpar(fill = col(x1), col = NA))
x2 = runif(n)
spiral_rect(1:n - 1, 0.5, 1:n, 1, gp = gpar(fill = col(x2), col = NA))
```

spiral_segments	<i>Add segments to a track</i>
-----------------	--------------------------------

Description

Add segments to a track

Usage

```
spiral_segments(x0, y0, x1, y1, gp = gpar(), arrow = NULL,
  track_index = current_track_index(), buffer = 10000)
```

Arguments

x0	X-locations of the start points of the segments.
y0	Y-locations of the start points of the segments.
x1	X-locations of the end points of the segments.
y1	Y-locations of the end points of the segments.
gp	Graphical parameters.
arrow	A arrow object.
track_index	Index of the track.
buffer	Number of segments to buffer.

Details

The segments on spiral are not straight lines while are more like curves. This means a spiral segment is formed by a list of real straight segments. If there are n_1 spiral segments, then there will be n_2 straight segments where n_2 is normally much larger than n_1 . To speed up drawing the spiral segments, the locations of the "real" segments are filled to a temporary data frame with buffer rows, when the number of rows exceeds buffer, [grid.segments](#) is called to draw all the buffered segments.

Value

No value is returned.

Examples

```
n = 1000
x0 = runif(n)
y0 = runif(n)
x1 = x0 + runif(n, min = -0.01, max = 0.01)
y1 = 1 - y0
```

```
spiral_initialize(xlim = range(c(x0, x1)))
spiral_track()
spiral_segments(x0, y0, x1, y1, gp = gpar(col = circlize::rand_color(n)))
```

```
n = 100
x0 = runif(n)
y0 = runif(n)
x1 = x0 + runif(n, min = -0.01, max = 0.01)
y1 = 1 - y0
```

```
spiral_initialize(xlim = range(c(x0, x1)))
spiral_track()
spiral_segments(x0, y0, x1, y1, arrow = arrow(length = unit(2, "mm")),
  gp = gpar(col = circlize::rand_color(n, luminosity = "bright"), lwd = runif(n, 0.5, 3)))
```

spiral_text

Add texts to a track

Description

Add texts to a track

Usage

```
spiral_text(x, y, text, offset = NULL, gp = gpar(),
  facing = c("downward", "inside", "outside", "clockwise", "reverse_clockwise",
  "curved_inside", "curved_outside"),
  letter_spacing = 0,
  nice_facing = FALSE, just = "centre", hjust = NULL, vjust = NULL,
  track_index = current_track_index(), ...)
```

Arguments

x	X-locations of the texts.
y	Y-locations of the texts.
text	A vector of texts.
offset	Radial offset of the text. The value should be a unit object.

gp	Graphical parameters.
facing	Facing of the text.
letter_spacing	Space between two letters. The value is a fraction of the width of current letter. It only works for curved texts.
nice_facing	If it is true, the facing will be automatically adjusted for texts which locate at different positions of the spiral. Note <code>hjust</code> and <code>vjust</code> will also be adjusted.
just	The justification of the text relative to (x, y). The same setting as in grid.text .
hjust	Horizontal justification. Value should be numeric. 0 means the left of the text and 1 means the right of the text.
vjust	Vertical justification. Value should be numeric. 0 means the bottom of the text and 1 means the top of the text.
track_index	Index of the track.
...	Pass to grid.text .

Details

For the curved text, it only supports one-line text.

Value

No value is returned.

Examples

```
x = seq(0.1, 0.9, length = 26)
text = strrep(letters, 6)
spiral_initialize(); spiral_track()
spiral_text(x, 0.5, text)

spiral_initialize(); spiral_track()
spiral_text(x, 0.5, text, facing = "inside")

spiral_initialize(); spiral_track()
spiral_text(x, 0.5, text, facing = "outside")

x = seq(0.1, 0.9, length = 10)
text = strrep(letters[1:10], 20)
spiral_initialize(); spiral_track()
spiral_text(x, 0.5, text, facing = "curved_inside")

spiral_initialize(); spiral_track()
spiral_text(x, 0.5, text, facing = "curved_outside")
```

spiral_track	<i>Add a new track or go to an existed track</i>
--------------	--

Description

Add a new track or go to an existed track

Usage

```
spiral_track(ylim = c(0, 1), height = 0.8, background = TRUE,  
             background_gp = gpar(col = NA, fill = "#EEEEEE"), reverse_y = FALSE,  
             track_index = current_track_index() + 1)
```

Arguments

ylim	Data range of the y-locations.
height	Height of the track. The value can be the fraction of the distance of the two neighbour loops. The value can also be a unit object.
background	Whether to draw the background of the track, i.e. border and filled color of background.
background_gp	Graphics parameters of the background.
reverse_y	Whether reverse the direction of y-axis.
track_index	Index of the track.

Details

If the track is already existed, the function simply mark the track as the current track and does nothing else.

Value

No value is returned.

Examples

```
spiral_initialize()  
spiral_track(height = 0.8)  
  
spiral_initialize()  
spiral_track(height = 0.4, background_gp = gpar(fill = "red"))  
spiral_track(height = 0.2, background_gp = gpar(fill = "green"))  
spiral_track(height = 0.1, background_gp = gpar(fill = "blue"))
```

spiral_xaxis	<i>Draw axis along the spiral</i>
--------------	-----------------------------------

Description

Draw axis along the spiral

Usage

```
spiral_xaxis(...)
```

Arguments

... All pass to [spiral_axis](#).

Value

No value is returned.

Examples

```
# There is no example  
NULL
```

spiral_yaxis	<i>Draw y-axis</i>
--------------	--------------------

Description

Draw y-axis

Usage

```
spiral_yaxis(side = c("both", "start", "end"), at = NULL, labels = TRUE,  
             ticks_length = unit(2, "bigpts"),  
             ticks_gp = gpar(), labels_gp = gpar(fontsize = 6),  
             track_index = current_track_index())
```

Arguments

side	On which side of the spiral the y-axis is drawn? "start" means the inside of the spiral and "end" means the outside of the spiral. Note if reverse was set to TRUE, then "start" corresponds to the most outside of the spiral.
at	Break points.
labels	Corresponding labels for the break points.
ticks_length	Length of the tick. Value should be a unit object.
ticks_gp	Graphics parameters for ticks.
labels_gp	Graphics parameters for labels.
track_index	Index of the track.

Value

No value is returned.

Examples

```
spiral_initialize(); spiral_track(height = 0.8)
spiral_yaxis("start")
spiral_yaxis("end", at = c(0, 0.25, 0.5, 0.75, 1), labels = letters[1:5])
```

TRACK_META

Get meta data in the current track

Description

Get meta data in the current track

Usage

```
TRACK_META
```

Details

The variable TRACK_META can only be used to get meta data from the "current" track. If the current track is not the one you want, you can first use [set_current_track](#) to set the current track.

Don't directly use TRACK_META. The value of TRACK_META itself is meaningless. Always use in form of TRACK_META\$name.

There are following meta data for the current track:

xlim: Data range on x-axis.

xmin: xlim[1].

xmax: xlim[2].

xrange: xlim[2] - xlim[1].

`xcenter`: `mean(xlim)`.
`theta_lim`: Range of the angles on the spiral, measured in radians.
`theta_min`: `theta_lim[1]`.
`theta_max`: `theta_lim[2]`.
`theta_range`: `theta_lim[2] - theta_lim[1]`.
`theta_center`: `mean(theta_lim)`.
`ylim`: Data range on y-axis.
`ymin`: `ylim[1]`.
`ymax`: `ylim[2]`.
`yrange`: `ylim[2] - ylim[1]`.
`ycenter`: `mean(ylim)`.
`rel_height`: Fraction of height of the track to the distance between two neighbouring loops.
`abs_height`: The height of the track, which is `rel_height` multiplied by the distance between two neighbouring loops.
`track_index`: Current track index.

Examples

```
# There is no example
NULL
```

<code>xy_to_cartesian</code>	<i>Convert data coordinates to the canvas coordinates</i>
------------------------------	---

Description

Convert data coordinates to the canvas coordinates

Usage

```
xy_to_cartesian(x, y, track_index = current_track_index())
```

Arguments

<code>x</code>	X-locations of the data points.
<code>y</code>	Y-locations of the data points.
<code>track_index</code>	Index of the track.

Details

The canvas coordinates correspond to the "native" coordinates of the viewport where the graphics are to be drawn.

Note different settings of `flip` and `reverse` in `spiral_initialize` affect the conversion.

Value

A data frame with two columns: x and y.

Examples

```
# There is no example  
NULL
```

xy_to_polar	<i>Convert data coordinates to polar coordinates</i>
-------------	--

Description

Convert data coordinates to polar coordinates

Usage

```
xy_to_polar(x, y, track_index = current_track_index(), flip = TRUE)
```

Arguments

x	X-locations of the data points.
y	Y-locations of the data points.
track_index	Index of the track.
flip	If it is FALSE, it returns theta for the original spiral (before flipping).

Details

Note different settings of flip and reverse in [spiral_initialize](#) affect the conversion.

Value

A data frame with two columns: theta (in radians) and r (the radius).

Examples

```
# There is no example  
NULL
```

\$.TRACK_META	<i>Get meta data in the current track</i>
---------------	---

Description

Get meta data in the current track

Usage

```
## S3 method for class 'TRACK_META'
x$name
```

Arguments

x	Always use TRACK_META.
name	Name of the meta name. For all supported names, type names(TRACK_META).

Details

The variable TRACK_META can only be used to get meta data from the "current" track. If the current track is not the one you want, you can first use [set_current_track](#) to set the current track.

There are following meta data for the current track:

xlim: Data range on x-axis.

xmin: xlim[1].

xmax: xlim[2].

xrange: xlim[2] - xlim[1].

xcenter: mean(xlim).

theta_lim: Range of the angles on the spiral, measured in radians.

theta_min: theta_lim[1].

theta_max: theta_lim[2].

theta_range: theta_lim[2] - theta_lim[1].

theta_center: mean(theta_lim).

ylim: Data range on y-axis.

ymin: ylim[1].

ymax: ylim[2].

yrange: ylim[2] - ylim[1].

ycenter: mean(ylim).

rel_height: Fraction of height of the track to the distance between two neighbouring loops.

abs_height: The height of the track, which is rel_height multiplied by the distance between two neighbouring loops.

track_index: Current track index.

Value

The corresponding value.

Examples

```
# There is no example  
NULL
```

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