

Package ‘spiralize’

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Type Package

Title Visualize Data on Spirals

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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>,
<https://jokergoo.github.io/spiralize/>

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<i>current_spiral</i>	<i>Get the current spiral object</i>
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Description

Get the current spiral object

Usage

```
current_spiral()
```

Details

The returned value is an object of `spiral` reference class. The following methods might be useful (assume the object is named `s`):

- `s$curve()`: It returns the radius for given angles (in radians).
- `s$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 `s$spiral_length(theta2) - s$spiral_length(theta1)`.

Also there are the following meta-data for the current spiral:

- `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 the current spiral

Description

Viewport name of the current spiral

Usage

```
current_spiral_vp()
```

Value

A string of the viewport name.

current_track_index *Helper functions for handling tracks*

Description

Helper functions for handling tracks

Usage

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

Arguments

track_index	The index of the track.
x	X-location of data points.
y	Y-location of data points.

Details

is_in_track() tests whether data points are inside a certain track.

Value

current_track_index() returns the index of the current track.
set_current_track() returns no value.
n_tracks() returns the number of available tracks.
is_in_track() returns a logical vector.

get_track_data *Meta-data of a track*

Description

Meta-data of a track

Usage

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

Arguments

field	Name of the field, see the Details section.
track_index	The index of the track.

Details

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

- `ymin`: Minimal value on the y-axis.
- `ymax`: Maximal value on the y-axis.
- `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 spiral loops.

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

Value

A numeric vector (of length one or two) for the corresponding field.

horizon_legend	<i>Legend for the horizon chart</i>
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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 ComplexHeatmap::Legend() .

Value

A `ComplexHeatmap::Legend` object.

Examples

```
# see examples in `spiral_horizon()` .
```

`solve_theta_from_spiral_length`

Get theta from given spiral lengths

Description

Get theta from given spiral lengths

Usage

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

Arguments

<code>len</code>	A vector of spiral lengths.
<code>interval</code>	Interval to search for the solution.
<code>offset</code>	Offset of the spiral. In the general form: $r = a + r*\theta$, 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 $l = f(\theta)$ where $f()$ is the complex equation for calculating l , `solve_theta_from_spiral_length()` tries to find θ with a known l . It uses `stats::uniroot()` to search for the solutions.

Value

The theta value.

Examples

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

spiral_arrow	<i>Draw arrows in the spiral direction</i>
--------------	--

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 grid::unit() 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	Graphical 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()
)
spiral_xaxis(...)
```

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>grid::unit()</code> object.
minor_ticks_length	Length of the minor ticks. The value should be a <code>grid::unit()</code> object.
ticks_gp	Graphical parameters for the ticks.
labels_gp	Graphical parameters for the labels.
track_index	Index of the track.
...	All pass to <code>spiral_axis()</code> .

Value

No value is returned.

Examples

```
spiral_initialize()
spiral_track()
spiral_axis()

# if the spiral is interpolated 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

<code>pos</code>	X-locations of the center of bars.
<code>value</code>	Height of bars. The value can be a simple numeric vector, or a matrix.
<code>baseline</code>	Baseline of the bars. Note it only works when <code>value</code> is a simple vector.
<code>bar_width</code>	Width of bars.
<code>gp</code>	Graphical parameters.
<code>track_index</code>	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*Clear the spiral curve***Description**

Clear the spiral curve

Usage

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

Arguments

<code>check_vp</code>	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.

spiral_dendrogram *Draw dendrogram*

Description

Draw dendrogram

Usage

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

Arguments

- | | |
|-------------|---|
| dend | A stats::dendrogram object. |
| gp | Graphical parameters of the dendrogram edges, mainly as a global setting. |
| track_index | Index of the track. |

Details

Graphical parameters for individual edges can be set via the `edgePar` attribute on each node in the dendrogram, see [stats::dendrogram](#) for how to set `edgePar`.

The dendrogram edges can also be rendered by [dendextend::color_branches\(\)](#).

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))
spiral_dendrogram(dend)

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_dendrogram(dend)
```

<code>spiral_git_commits</code>	<i>Visualize git commits</i>
---------------------------------	------------------------------

Description

Visualize git commits

Usage

```
spiral_git_commits(
  repo = ".",
  show_legend = TRUE,
  start = NULL,
  end = Sys.Date(),
  pt_range = c(2, 16),
  commits_range = c(1, ceiling(quantile(n[n > 0], 0.95))),
  type = c("points", "heatmap"),
  colors = c("#3288BD", "#99D594", "#E6F598", "#FFFFBF", "#FEE08B", "#FC8D59", "#D53E4F")
)
```

Arguments

<code>repo</code>	Path of the git repo. The value can be a single repo or a vector of repos.
<code>show_legend</code>	Whether to show the legend.
<code>start</code>	Start date. By default it is the first date of the commit. The value can be a string such as "2022-01-01" or a <code>base::Date</code> object.
<code>end</code>	End date. By default it is the current date. The value can be a string such as "2022-01-01" or a <code>base::Date</code> object.
<code>pt_range</code>	Range of the point sizes. The default is between 1 and the 90 percentile of daily commits.
<code>commits_range</code>	Range of the numbers of commits.
<code>type</code>	Type of the plot.
<code>colors</code>	If type is the heatmap, it controls the list of colors.

Examples

```
## Not run:
spiral_git_commits("~/project/development/ComplexHeatmap")
spiral_git_commits("~/project/development/ComplexHeatmap", type = "heatmap")

## End(Not run)
```

<code>spiral_highlight</code>	<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

<code>x1</code>	Start location of the highlighted section.
<code>x2</code>	End location of the highlighted section.
<code>type</code>	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.
<code>padding</code>	When the highlight type is "rect", it controls the padding of the highlighted region. The value should be a <code>grid::unit()</code> 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.
<code>line_side</code>	If the highlight type is "line", it controls which side of the track to draw the lines.
<code>line_width</code>	Width of the annotation line. Value should be a <code>grid::unit()</code> object.
<code>gp</code>	Graphical parameters.
<code>track_index</code>	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 grid::unit() object with length one or two.
gp	Graphical 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 ceiling(max(abs(y))/slice_size).
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.

See Also

[horizon_legend\(\)](#) for generating the legend.

Examples

```
df = readRDS(system.file("extdata", "global_temperature.rds", package = "spiralyze"))
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)
spiral_track()
spiral_horizon(df$date, df$Mean, use_bar = TRUE)

# with legend
require(ComplexHeatmap)
spiral_initialize_by_time(xlim = range(df$date), unit_on_axis = "months", period = "year",
    period_per_loop = 20, polar_lines_by = 360/20,
    vp_param = list(x = unit(0, "npc"), just = "left"))
spiral_track()
lt = spiral_horizon(df$date, df$Mean, use_bar = TRUE)
lgd = horizon_legend(lt, title = "Temperature difference")
draw(lgd, x = unit(1, "npc") + unit(2, "mm"), just = "left")
```

spiral_info

Information of the current spiral

Description

Information of the current spiral

Usage

`spiral_info()`

Details

It prints information of the current spiral.

Value

No value is returned.

Examples

```
spiral_initialize()
spiral_track(ylim = c(0, 1), height = 0.4)
spiral_track(ylim = c(-10, 10), height = 0.4)
spiral_info()
```

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

xlim	Range on x-locations.
start	Start of the spiral, in degree. start and end should be positive and start should be smaller than end.
end	End of the spiral, in degree.
scale_by	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 clockwise 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 <code>grid::unit()</code> of length of one to two.
<code>newpage</code>	Whether to apply <code>grid::grid.newpage()</code> before making the plot?
<code>vp_param</code>	A list of parameters sent to <code>grid::viewport()</code> .

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

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

Details

It is basically the same as `spiral_initialize()`. The only difference is the axis labels are automatically formated 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 base::Date , base::POSIXlt or base::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 spiral loop to represent a complete year?
<code>period_per_loop</code>	How many periods to put in a spiral 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 <code>period</code> is "weeks" or 360/24 is set if <code>period</code> is set to "hours". When <code>period</code> 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 year is always put on $\theta = 0 + 2\pi k$ where k is the index of spiral 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 gpar(fill = ...) controls the filled colors 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 the following global parameters:

- `min_segment_len` Minimal length of the segment that partitions a curve.
- `help` Whether to print the help messages?

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

```
spiral_opt
```

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

Arguments

obj	A stats::dendrogram object.
gp	Graphical parameters of the tree edges, mainly as a global setting.
log	Whether the height of the tree to be log-transformed $\log_{10}(x + 1)$?
reverse	Whether the tree to 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.

Details

[phylo_to_dendrogram\(\)](#) converts a phylo object to a dendrogram object.

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 `stats::plot.dendrogram()` function. However, you can still apply `stats::cutree()` to the returned dendrogram object with no problem and the dendrogram can be properly drawn with the **ComplexHeatmap** package (see examples).

Value

Height of the phylogenetic tree.
A `stats::dendrogram` object.

Examples

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

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

spiral_pkg_downloads Visualize package downloads

Description

Visualize package downloads

Usage

```
spiral_pkg_downloads(
  pkg,
  from = "2012-10-01",
  to = "last-day",
  show_legend = TRUE
)
```

Arguments

<code>pkg</code>	A single CRAN package name.
<code>from</code>	Starting date.
<code>to</code>	Ending date.
<code>show_legend</code>	Whether to show the legend.

Details

The **cranlogs** package is used to retrieve the download history from the Rstudio server.

Examples

```
spiral_pkg_downloads("ggplot2")
```

spiral_points	<i>Add points to a track</i>
---------------	------------------------------

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 grid::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

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

Value

No value is returned.

Examples

```
x = seq(0, 2*pi*10, length = 1000)
y = c(sin(x), cos(rev(x)))
x2 = c(x, rev(x))

# in the normal cartesian coordinate system
plot(NULL, xlim = range(x2), ylim = range(y))
polygon(x2, y, col = "red")

# in the spiral coordinate system
spiral_initialize(xlim = range(x2))
spiral_track(ylim = range(y))
spiral_polygon(x2, y, gp = gpar(fill = "red"))

# try a different scale
spiral_initialize(xlim = range(x2), scale_by = "curve_length")
spiral_track(ylim = range(y))
spiral_polygon(x2, y, gp = gpar(fill = "red"))
```

spiral_raster *Add image to a track*

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 `grid::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")
```

spiral_rect	<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 *Add segments to a track*

Description

Add segments to a track

Usage

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

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 grid::arrow() object.
track_index	Index of the track.

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()
col = circlize::rand_color(n, luminosity = "bright")
spiral_segments(x0, y0, x1, y1,
  arrow = arrow(length = unit(2, "mm")), gp = gpar(col = col))

# if the segments are short and you want the straight "real" segments
spiral_initialize(xlim = range(c(x0, x1)))
spiral_track()
df0 = xy_to_cartesian(x0, y0)
df1 = xy_to_cartesian(x1, y1)
grid.segments(df0$x, df0$y, df1$x, df1$y, default.units = "native",
  arrow = arrow(length = unit(2, "mm")), gp = gpar(col = col))

```

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 grid::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 hjust and vjust will also be adjusted.
just	The justification of the text relative to (x, y). The same setting as in grid::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::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*Add a new track or move to an existed track***Description**

Add a new track or move to an existed track

Usage

```

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

```

Arguments

<code>ylim</code>	Data range of the y-locations.
<code>height</code>	Height of the track. The value can be the fraction of the distance of the two neighbour spiral loops. The value can also be a <code>grid::unit()</code> object.
<code>background</code>	Whether to draw the background of the track, i.e. border and filled color of background.
<code>background_gp</code>	Graphical parameters of the background.
<code>reverse_y</code>	Whether reverse the direction of y-axis (i.e. pointing to the center of the spiral)?
<code>gradient</code>	Whether draw the background in gradient? The value can be a positive integer of the number of gradients from <code>background_gp\$fill</code> to white.
<code>track_index</code>	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_initialize()
spiral_track(height = 0.8, gradient = TRUE) # by default 10 gradients

spiral_initialize()
spiral_track(height = 0.8, background_gp = gpar(fill = "red"), gradient = 5)
```

spiral_yaxis

Draw y-axis

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 in <code>spiral_initialize()</code> , then "start" corresponds to the outside of the spiral.
at	Break points.
labels	Corresponding labels for the break points.
ticks_length	Length of the tick. Value should be a <code>grid::unit()</code> object.

ticks_gp	Graphical parameters for ticks.
labels_gp	Graphical 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

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

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

## S3 method for class 'TRACK_META'
x[[i, exact = TRUE]]

## S3 method for class 'TRACK_META'
x[i]

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

Arguments

x	The TRACK_META object.
name	Name of the meta name. For all supported names, type names(TRACK_META).
i	Name of the meta name. For all supported names, type names(TRACK_META).
exact	Please ignore.
...	Additional parameters.

Format

An object of class TRACK_META of length 1.

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 change 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 the 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

```
spiral_initialize(xlim = c(0, 1))
spiral_track(ylim = c(0, 1))
for(nm in names(TRACK_META)) {
  cat(nm, ":\n", sep = "")
  print(TRACK_META[[nm]])
  cat("\n")
}
names(TRACK_META)
```

<code>xy_to_cartesian</code>	<i>Transform between coordinate systems</i>
------------------------------	---

Description

Transform between coordinate systems

Usage

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

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

polar_to_cartesian(theta, r)

cartesian_to_polar(x, y)

cartesian_to_xy(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.
<code>flip</code>	If it is FALSE, it returns theta for the original spiral (before flipping).
<code>theta</code>	Angles, in radians.
<code>r</code>	Radius.

Details

There are three coordinate systems: the data coordinate system (xy), the polar coordinate system (polar) and the canvas coordinate system (cartesian). The canvas coordinates correspond to the "native" coordinates of the viewport where the graphics are drawn.

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

`xy_to_cartesian()` converts from the data coordinate system to the canvas coordinate system.

`xy_to_polar()` converts from the data coordinate system to the polar coordinate system.

`polar_to_cartesian()` converts from the polar coordinate system to the canvas coordinate system.

`cartesian_to_polar()` converts from the canvas coordinate system to the polar coordinate system.

`cartesian_to_xy()` converts from the canvas coordinate system to the data coordinate system. The data points are assigned to the nearest inner spiral loops (if the point is located inside a certain spiral loop, the distance is zero).

Value

`xy_to_cartesian()` returns A data frame with two columns: `x` and `y`.
`xy_to_polar()` returns a data frame with two columns: `theta` (in radians) and `r` (the radius).
`polar_to_cartesian()` returns a data frame with two columns: `x` and `y`.
`cartesian_to_polar()` returns a data frame with two columns: `theta` (in radians) and `r` (the radius).
`cartesian_to_xy()` returns a data frame with two columns: `x` and `y`.

Examples

```
x = runif(2)
y = runif(2)
spiral_initialize(xlim = c(0, 1))
spiral_track(ylim = c(0, 1))
spiral_points(x, y)
xy_to_cartesian(x, y)
xy_to_polar(x, y)

x = runif(100, -4, 4)
y = runif(100, -4, 4)
spiral_initialize(xlim = c(0, 1))
spiral_track(ylim = c(0, 1))
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, gp = gpar(col = 2))
```

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