

Package ‘tourr’

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Title Tour Methods for Multivariate Data Visualisation

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Description Implements geodesic interpolation and basis generation functions that allow you to create new tour methods from R.

Depends R (>= 3.6)

Imports MASS, tibble, dplyr, stats, utils, grDevices

Suggests TeachingDemos, ash, energy, testthat, colorspace, ggplot2, mgcv, gifski, geozoo, knitr, rmarkdown, tidy, covr

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LazyData true

URL <https://github.com/ggobi/tourr>

BugReports <https://github.com/ggobi/tourr/issues>

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anchored_orthogonal_distance
Calculate orthogonal distances

Description

For each datapoint this function calculates the orthogonal distance from the anchored projection plane.

Usage

anchored_orthogonal_distance(plane, data, anchor = NULL)

Arguments

- plane matrix specifying the projection plane
- data data frame or matrix
- anchor A vector specifying the reference point to anchor the plane If NULL (default) the slice will be anchored at the origin.

Value

distance vector

andrews	<i>Compute Andrews' curves</i>
---------	--------------------------------

Description

This function takes a numeric vector of input, and returns a function which allows you to compute the value of the Andrew's curve at every point along its path from $-\pi$ to π .

Usage

```
andrews(x)
```

Arguments

x input a new parameter

Value

a function with single argument, theta

Examples

```
a <- andrews(1:2)
a(0)
a(-pi)
grid <- seq(-pi, pi, length = 50)
a(grid)

plot(grid, andrews(1:2)(grid), type = "l")
plot(grid, andrews(runif(5))(grid), type = "l")
```

angular_breaks	<i>Returns n equidistant bins between -pi and pi</i>
----------------	------------------------------------------------------

Description

Returns n equidistant bins between $-\pi$ and π

Usage

```
angular_breaks(n)
```

Arguments

n number of bins

animate	<i>Animate a tour path.</i>
---------	-----------------------------

Description

This is the function that powers all of the tour animations. If you want to write your own tour animation method, the best place to start is by looking at the code for animation methods that have already implemented in the package.

Usage

```
animate(  
  data,  
  tour_path = grand_tour(),  
  display = display_xy(),  
  start = NULL,  
  aps = 1,  
  fps = 10,  
  max_frames = Inf,  
  rescale = TRUE,  
  sphere = FALSE,  
  ...  
)
```

Arguments

data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour
display	takes the display that is suppose to be used, defaults to the xy display
start	projection to start at, if not specified, uses default associated with tour path
aps	target angular velocity (in radians per second)
fps	target frames per second (defaults to 15, to accommodate RStudio graphics device)
max_frames	the maximum number of bases to generate. Defaults to Inf for interactive use (must use Ctrl + C to terminate), and 1 for non-interactive use.
rescale	if true, rescale all variables to range [0,1]?
sphere	if true, sphere all variables
...	ignored

Details

See [render](#) to render animations to disk.

Value

an (invisible) list of bases visited during this tour

Examples

```
f <- flea[, 1:6]
animate(f, grand_tour(), display_xy())
# or in short
animate(f)
animate(f, max_frames = 30)

animate(f, max_frames = 10, fps = 1, aps = 0.1)
```

areColors	<i>Test if all entries are colors</i>
-----------	---------------------------------------

Description

Test if all entries are colors

Usage

```
areColors(x)
```

Arguments

x vector

center	<i>Center a numeric vector by subtracting off its mean.</i>
--------	-------------------------------------------------------------

Description

Center a numeric vector by subtracting off its mean.

Usage

```
center(x)
```

Arguments

x numeric vector

cmass	<i>Central mass index.</i>
-------	----------------------------

Description

Calculates the central mass index. See Cook and Swayne (2007) Interactive and Dynamic Graphics for Data Analysis for equations.

Usage

```
cmass()
```

dcor2d	<i>Distance correlation index.</i>
--------	------------------------------------

Description

Computes the distance correlation based index on 2D projections of the data.

Usage

```
dcor2d()
```

dependence_tour	<i>A dependence tour path.</i>
-----------------	--------------------------------

Description

The dependence tour combines a set of independent 1d tours to produce a nd tour. For the special case of 2d, this is known as a correlation tour. This tour corresponds to the multivariate method known as generalised canonical correlation, and is used to investigate dependence between groups of variables.

Usage

```
dependence_tour(pos)
```

Arguments

pos	a numeric vector describing which variables are mapped to which dimensions: 1 corresponds to first, 2 to second etc.
-----	----------------------------------------------------------------------------------------------------------------------

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
animate_xy(flea[, 1:3], dependence_tour(c(1, 2, 2)))
animate_xy(flea[, 1:4], dependence_tour(c(1, 2, 1, 2)))
animate_pcp(flea[, 1:6], dependence_tour(c(1, 2, 3, 2, 1, 3)))
```

display_andrews	<i>Andrews' curves tour path animation.</i>
-----------------	---------------------------------------------

Description

Animate a nD tour path with Andrews' curves. For more details about Andrew's curves, see [andrews](#)

Usage

```
display_andrews(col = "black", ...)

animate_andrews(data, tour_path = grand_tour(3), col = "black", ...)
```

Arguments

col	color to be plotted. Defaults to "black"
...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

See Also

[animate](#) for options that apply to all animations

Examples

```
animate_andrews(flea[, 1:6])
animate_andrews(flea[, 1:6], grand_tour(d = 3))
animate_andrews(flea[, 1:6], grand_tour(d = 6))

# It's easy to experiment with different tour paths:
animate_andrews(flea[, 1:6], guided_tour(cmass()))
```

display_density2d *Display tour path with a density and scatterplot*

Description

Animate a 2D tour path with density contour(s) and a scatterplot.

Usage

```
display_density2d(
  center = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  cex = 1,
  contour_quartile = c(0.25, 0.5, 0.75),
  edges = NULL,
  ...
)

animate_density2d(data, tour_path = grand_tour(), ...)
```

Arguments

center	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch	shape of the point to be plotted. Defaults to 20.
cex	size of the point to be plotted. Defaults to 1.
contour_quartile	Vector of quartiles to plot the contours at. Defaults to 5.
edges	A two column integer matrix giving indices of ends of lines.
...	other arguments passed on to animate and display_density2d
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Examples

```

animate_density2d(flea[, 1:6])
animate(flea[, 1:6], tour_path = grand_tour(), display = display_density2d())
animate(flea[, 1:6],
  tour_path = grand_tour(),
  display = display_density2d(axes = "bottomleft")
)
animate(flea[, 1:6],
  tour_path = grand_tour(),
  display = display_density2d(half_range = 0.5)
)
animate_density2d(flea[, 1:6], tour_path = little_tour())

animate_density2d(flea[, 1:3], tour_path = guided_tour(holes()), sphere = TRUE)
animate_density2d(flea[, 1:6], center = FALSE)

# The default axes are centered, like a biplot, but there are other options
animate_density2d(flea[, 1:6], axes = "bottomleft")
animate_density2d(flea[, 1:6], axes = "off")
animate_density2d(flea[, 1:6], dependence_tour(c(1, 2, 1, 2, 1, 2)),
  axes = "bottomleft"
)

animate_density2d(flea[, -7], col = flea$species)

# You can also draw lines
edges <- matrix(c(1:5, 2:6), ncol = 2)
animate(
  flea[, 1:6], grand_tour(),
  display_density2d(axes = "bottomleft", edges = edges)
)

```

display_depth

Display 3d projection with depth cues

Description

Suggestion to use gray background and colour saturation (instead of gray shading) by Graham Wills.

Usage

```

display_depth(center = TRUE, half_range = NULL, ...)

animate_depth(data, tour_path = grand_tour(3), ...)

```

Arguments

center	should projected data be centered to have mean zero (default: TRUE). This pins the centre of the data to the same place, and makes it easier to focus on the shape.
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

See Also

[animate](#) for options that apply to all animations

Examples

```
animate_depth(flea[, 1:6])
```

display_dist

1d distribution tour path animation.

Description

Animate a 1d tour path with a density plot or histogram.

Usage

```
display_dist(
  method = "density",
  center = TRUE,
  half_range = NULL,
  rug = FALSE,
  ...
)
```

```
animate_dist(data, tour_path = grand_tour(1), ...)
```

Arguments

method	display method, histogram or density plot
center	should 1d projection be centered to have mean zero (default: TRUE). This pins the centre of distribution to the same place, and makes it easier to focus on the shape of the distribution.
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.

rug	draw rug plot showing position of actual data points?
...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

See Also

[animate](#) for options that apply to all animations

Examples

```
animate_dist(flea[, 1:6])

# When the distribution is not centred, it tends to wander around in a
# distracting manner
animate_dist(flea[, 1:6], center = FALSE)

# Alternatively, you can display the distribution with a histogram
animate_dist(flea[, 1:6], method = "hist")
```

display_faces

Chernoff faces tour path animation.

Description

Animate a nD tour path with Chernoff's faces. Can display up to 18 dimensions.

Usage

```
display_faces(...)

animate_faces(data, tour_path = grand_tour(3), ...)
```

Arguments

...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Details

This function requires the TeachingDemos package to draw the Chernoff faces. See [faces2](#) for more details.

See Also

[animate](#) for options that apply to all animations

Examples

```
# The drawing code is fairly slow, so this animation works best with a
# limited number of cases
animate_faces(flea[1:2, 1:6])
animate_faces(flea[1:4, 1:6])

animate_faces(flea[1:2, 1:6], grand_tour(5))
```

display_groupxy

Display 2D tour projections displayed separately by groups

Description

This function is designed to allow comparisons across multiple groups, especially for examining things like two (or more) different models on the same data. The primary display is a scatterplot, with lines or contours overlaid.

Usage

```
display_groupxy(
  centr = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  cex = 1,
  edges = NULL,
  group_by = NULL,
  plot_xgp = TRUE,
  ...
)

animate_groupxy(data, tour_path = grand_tour(), ...)
```

Arguments

centr	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch	shape of the point to be plotted. Defaults to 20.
cex	size of the point to be plotted. Defaults to 1.
edges	A two column integer matrix giving indices of ends of lines.

group_by	variable to group by. Must have less than 25 unique values.
plot_xgp	if TRUE, plots points from other groups in light grey
...	other arguments passed on to animate and display_groupxy
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Examples

```
f <- flea[, 1:6]
col <- rainbow(length(unique(flea$species)))[as.numeric(as.factor(flea$species))]
pch <- as.numeric(flea$species) + 14

animate_groupxy(f, col = col, pch = pch, group_by = flea$species)
animate_groupxy(f, col = col, pch = pch, group_by = flea$species, plot_xgp = FALSE)
```

display_image	<i>Image tour path animation.</i>
---------------	-----------------------------------

Description

Animate a 1d tour path with an image plot. This animation requires a different input data structure, a 3d array. The first two dimensions are locations on a grid, and the 3rd dimension gives the observations to be mixed with the tour.

Usage

```
display_image(xs, ys, ...)

animate_image(data, tour_path = grand_tour(1), ...)
```

Arguments

xs	x limit that is used in making the size of the plot
ys	y limit that is used in making the size of the plot
...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

See Also

[animate](#) for options that apply to all animations

Examples

```
str(ozone)
animate_image(ozone)
```

display_pca	<i>Display tour path with principal component scores with original axes</i>
-------------	-----------------------------------------------------------------------------

Description

Animate a 2D tour path on data that has been transformed into principal components, and also show the original variable axes.

Usage

```
display_pca(
  center = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  cex = 1,
  pc_coefs = NULL,
  edges = NULL,
  edges.col = "black",
  ...
)
```

```
animate_pca(data, tour_path = grand_tour(), rescale = TRUE, ...)
```

Arguments

center	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black".
pch	shape of the point to be plotted. Defaults to 20.
cex	size of the point to be plotted. Defaults to 1.
pc_coefs	coefficients relating the original variables to principal components. This is required.
edges	A two column integer matrix giving indices of ends of lines.
edges.col	colour of edges to be plotted, Defaults to "black".
...	other arguments passed on to animate and display_slice
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour
rescale	if true, rescale all variables to range [0,1].

Examples

```
flea_std <- scale(flea[, 1:6])
flea_pca <- prcomp(flea_std, center = FALSE, )
flea_coefs <- flea_pca$rotation[, 1:3]
flea_scores <- flea_pca$x[, 1:3]
animate_pca(flea_scores, pc_coefs = flea_coefs)
```

display_pcp

Parallel coordinates tour path animation.

Description

Animate a nD tour path with a parallel coordinates plot.

Usage

```
display_pcp(...)
```

```
animate_pcp(data, tour_path = grand_tour(3), ...)
```

Arguments

...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Details

The lines show the observations, and the points, the values of the projection matrix.

See Also

[animate](#) for options that apply to all animations

Examples

```
animate_pcp(flea[, 1:6], grand_tour(3))
animate_pcp(flea[, 1:6], grand_tour(5))
```


display_sage

*Display tour path with a sage scatterplot***Description**

Animate a 2D tour path with a sage scatterplot that uses a radial transformation on the projected points to re-allocate the volume projected across the 2D plane.

Usage

```
display_sage(
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  gam = 1,
  R = NULL,
  ...
)

animate_sage(data, tour_path = grand_tour(), ...)
```

Arguments

axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch	marker for points. Defaults to 20.
gam	scaling of the effective dimensionality for rescaling. Defaults to 1.
R	scale for the radial transformation. If not set, defaults to maximum distance from origin to each row of data.
...	other arguments passed on to animate and display_sage
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Examples

```
# Generate uniform samples in a 10d sphere using the geozoo package
sphere10 <- geozoo::sphere.solid.random(10)$points
# Columns need to be named before launching the tour
colnames(sphere10) <- paste0("x", 1:10)
# Standard grand tour display, points cluster near center
animate_xy(sphere10)
# Sage display, points are uniformly distributed across the disk
animate_sage(sphere10)
```

display_scattermat	<i>Scatterplot matrix tour path animation.</i>
--------------------	------------------------------------------------

Description

Animate a nD tour path with a scatterplot matrix.

Usage

```
display_scattermat(...)
```

```
animate_scattermat(data, tour_path = grand_tour(3), ...)
```

Arguments

...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Details

The lines show the observations, and the points, the values of the projection matrix.

See Also

[animate](#) for options that apply to all animations

Examples

```
animate_scattermat(flea[, 1:6], grand_tour(2))  
animate_scattermat(flea[, 1:6], grand_tour(6))
```

display_slice	<i>Display tour path with a sliced scatterplot</i>
---------------	----------------------------------------------------

Description

Animate a 2D tour path with a sliced scatterplot.

Usage

```

display_slice(
  center = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch_slice = 20,
  pch_other = 46,
  cex_slice = 2,
  cex_other = 1,
  v_rel = NULL,
  anchor = NULL,
  anchor_nav = "off",
  edges = NULL,
  edges.col = "black",
  ...
)

animate_slice(data, tour_path = grand_tour(), rescale = TRUE, ...)

```

Arguments

center	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch_slice	marker for plotting points inside the slice. Defaults to 20.
pch_other	marker for plotting points outside the slice. Defaults to 46.
cex_slice	size of the points inside the slice. Defaults to 2.
cex_other	size if the points outside the slice. Defaults to 1.
v_rel	relative volume of the slice. If not set, suggested value is calculated and printed to the screen.
anchor	A vector specifying the reference point to anchor the slice. If NULL (default) the slice will be anchored at the data center.
anchor_nav	position of the anchor: center, topright or off
edges	A two column integer matrix giving indices of ends of lines.
edges.col	colour of edges to be plotted, Defaults to "black.
...	other arguments passed on to animate and display_slice
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour
rescale	if true, rescale all variables to range [0,1].

Examples

```
# Generate samples on a 3d and 5d hollow sphere using the geozoo package
sphere3 <- geozoo::sphere.hollow(3)$points
sphere5 <- geozoo::sphere.hollow(5)$points

# Columns need to be named before launching the tour
colnames(sphere3) <- c("x1", "x2", "x3")
colnames(sphere5) <- c("x1", "x2", "x3", "x4", "x5")

# Animate with the slice display using the default parameters
animate_slice(sphere3)
animate_slice(sphere5)

# Animate with off-center anchoring
anchor3 <- matrix(rep(0.7, 3), ncol=3)
anchor5 <- matrix(rep(0.3, 5), ncol=5)
animate_slice(sphere3, anchor = anchor3)
# Animate with thicker slice to capture more points in each view
animate_slice(sphere5, anchor = anchor5, v_rel = 0.02)
```

display_stars

Star glyph tour path animation.

Description

Animate a nD tour path with star glyphs.

Usage

```
display_stars(...)
```

```
animate_stars(data, tour_path = grand_tour(3), ...)
```

Arguments

...	other arguments passed on to stars
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Details

Currently, scaling doesn't seem to be computed absolutely correctly, as centres move around as well as outside points.

See Also

[animate](#) for options that apply to all animations

Examples

```

animate_stars(flea[1:10, 1:6])
animate_stars(flea[1:10, 1:6], grand_tour(5))
animate_stars(flea[, 1:6], grand_tour(5))
animate_stars(flea[1:10, 1:6], grand_tour(5),
  col.stars = rep("grey50", 10), radius = FALSE
)

```

display_stereo *Anaglyph tour path animation.*

Description

Uses red-blue anaglyphs to display a 3d tour path. You'll need some red- blue glasses to get much out of this displays!

Usage

```
display_stereo(blue, red, cex = 1, ...)
```

```

animate_stereo(
  data,
  tour_path = grand_tour(3),
  blue = rgb(0, 0.91, 0.89),
  red = rgb(0.98, 0.052, 0),
  ...
)

```

Arguments

blue	blue colour (for right eye)
red	red colour (for left eye)
cex	size of the point to be plotted. Defaults to 1.
...	other arguments passed on to animate
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Examples

```
animate_stereo(flea[, 1:6])
```

display_trails *Display tour path with trails*

Description

Animate a 2D tour path with a point trails

Usage

```
display_trails(
  center = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  cex = 1,
  past = 3,
  ...
)
```

```
animate_trails(data, tour_path = grand_tour(), ...)
```

Arguments

center	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch	shape of the point to be plotted. Defaults to 20.
cex	magnification of plotting text relative to default. Defaults to 1.
past	draw line between current projection and projection past steps ago
...	other arguments passed on to animate and display_xy
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

display_xy	<i>Display tour path with a scatterplot</i>
------------	---------------------------------------------

Description

Animate a 2D tour path with a scatterplot.

Usage

```
display_xy(
  center = TRUE,
  axes = "center",
  half_range = NULL,
  col = "black",
  pch = 20,
  cex = 1,
  edges = NULL,
  edges.col = "black",
  ...
)
```

```
animate_xy(data, tour_path = grand_tour(), ...)
```

Arguments

center	if TRUE, centers projected data to (0,0). This pins the center of data cloud and make it easier to focus on the changing shape rather than position.
axes	position of the axes: center, bottomleft or off
half_range	half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.
col	color to be plotted. Defaults to "black"
pch	shape of the point to be plotted. Defaults to 20.
cex	size of the point to be plotted. Defaults to 1.
edges	A two column integer matrix giving indices of ends of lines.
edges.col	colour of edges to be plotted, Defaults to "black"
...	other arguments passed on to animate and display_xy
data	matrix, or data frame containing numeric columns
tour_path	tour path generator, defaults to 2d grand tour

Examples

```

animate_xy(flea[, 1:6])
animate(flea[, 1:6], tour_path = grand_tour(), display = display_xy())
animate(flea[, 1:6],
  tour_path = grand_tour(),
  display = display_xy(axes = "bottomleft")
)
animate(flea[, 1:6],
  tour_path = grand_tour(),
  display = display_xy(half_range = 0.5)
)
animate_xy(flea[, 1:6], tour_path = little_tour())
animate_xy(flea[, 1:3], tour_path = guided_tour(holes()), sphere = TRUE)
animate_xy(flea[, 1:6], center = FALSE)

# The default axes are centered, like a biplot, but there are other options
animate_xy(flea[, 1:6], axes = "bottomleft")
animate_xy(flea[, 1:6], axes = "off")
animate_xy(flea[, 1:6], dependence_tour(c(1, 2, 1, 2, 1, 2)),
  axes = "bottomleft"
)

animate_xy(flea[, -7], col = flea$species)

# You can also draw lines
edges <- matrix(c(1:5, 2:6), ncol = 2)
animate(
  flea[, 1:6], grand_tour(),
  display_xy(axes = "bottomleft", edges = edges)
)

```

estimate_eps

Estimate cutoff eps for section pursuit.

Description

Estimate cutoff eps for section pursuit.

Usage

```
estimate_eps(N, p, res, K, K_theta, r_breaks)
```

Arguments

N	total number of points in the input data.
p	number of dimensions of the input data.
res	resolution, (slice radius)/(data radius)
K	total number of bins

K_theta number of angular bins
r_breaks boundaries of the radial bins

Flea measurements *Flea beetle measurements*

Description

This data is from a paper by A. A. Lubischew, "On the Use of Discriminant Functions in Taxonomy", Biometrics, Dec 1962, pp.455-477.

Format

A 74 x 7 numeric array

Details

- tars1, width of the first joint of the first tarsus in microns (the sum of measurements for both tarsi)
- tars2, the same for the second joint
- head, the maximal width of the head between the external edges of the eyes in 0.01 mm
- ade1, the maximal width of the aedeagus in the fore-part in microns
- ade2, the front angle of the aedeagus (1 unit = 7.5 degrees)
- ade3, the aedeagus width from the side in microns
- species, which species is being examined - concinna, heptapotamica, heikertingeri

Examples

```
head(flea)
animate_xy(flea[, -7])
animate_xy(flea[, -7], col = flea[, 7])
```

frozen_guided_tour *The frozen guided tour*

Description

The frozen guided tour

Usage

```
frozen_guided_tour(frozen, index_f, d = 2, max.tries = 25)
```

Arguments

frozen	matrix of frozen variables, as described in freeze
index_f	the index function to optimise.
d	target dimensionality
max.tries	the maximum number of unsuccessful attempts to find a better projection before giving up

See Also

[cmass](#), [holes](#) and [lda_pp](#) for examples of index functions. The function should take a numeric matrix and return a single number, preferably between 0 and 1.

Examples

```
frozen <- matrix(NA, nrow = 4, ncol = 2)
frozen[3, ] <- .5
animate_xy(flea[, 1:4], frozen_guided_tour(frozen, holes()))
```

frozen_tour	<i>A frozen tour path.</i>
-------------	----------------------------

Description

A frozen tour fixes some of the values of the orthonormal projection matrix and allows the others to vary freely according to any of the other tour methods. This frozen tour is a frozen grand tour. See [frozen_guided_tour](#) for a frozen guided tour.

Usage

```
frozen_tour(d = 2, frozen)
```

Arguments

d	target dimensionality
frozen	matrix of frozen variables, as described in freeze

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
frozen <- matrix(NA, nrow = 4, ncol = 2)
frozen[3, ] <- .5
animate_xy(flea[, 1:4], frozen_tour(2, frozen))

frozen <- matrix(NA, nrow = 4, ncol = 2)
frozen[1, 1] <- 0.5
animate_xy(flea[, 1:4], frozen_tour(2, frozen))

# Doesn't work - a bug?
frozen <- matrix(NA, nrow = 4, ncol = 2)
frozen[1:2, 1] <- 1 / 4
animate_xy(flea[, 1:4], frozen_tour(2, frozen))

## Not run:
# This freezes one entire direction which causes a problem,
# and is caught by error handling.
# If you want to do this it would be best with a dependence
# tour, with one variable set one axis, eg 3rd variable to
# x axis would be indicated from the code below
frozen <- matrix(NA, nrow = 4, ncol = 2)
frozen[3, ] <- c(0, 1)
animate_xy(flea[, 1:4], frozen_tour(2, frozen))

## End(Not run)

# Two frozen variables in five.
frozen <- matrix(NA, nrow = 5, ncol = 2)
frozen[3, ] <- .5
frozen[4, ] <- c(-.2, .2)
animate_xy(flea[, 1:5], frozen_tour(2, frozen))
```

grand_tour

A grand tour path.

Description

This method generates target bases by randomly sampling on the space of all d-dimensional planes in p-space.

Usage

```
grand_tour(d = 2, ...)
```

Arguments

d	target dimensionality
...	arguments sent to the generator

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
# All animation methods use the grand tour path by default
animate_dist(flea[, 1:6])
animate_xy(flea[, 1:6])
animate_pcp(flea[, 1:6])
animate_pcp(flea[, 1:6], grand_tour(4))

# The grand tour is a function:
tour2d <- grand_tour(2)
is.function(tour2d)

# with two parameters, the previous projection and the data set
args(tour2d)
# if the previous projection is null, it will generate a starting
# basis, otherwise the argument is ignored
tour2d(NULL, mtcars)
# the data argument is just used to determine the correct dimensionality
# of the output matrix
tour2d(NULL, mtcars[, 1:2])
```

guided_section_tour *A guided section tour path.*

Description

The guided section tour is a variation of the guided tour that is using a section pursuit index for the selection of target planes.

Usage

```
guided_section_tour(
  index_f,
  d = 2,
  alpha = 0.5,
  cooling = 0.99,
  max.tries = 25,
  max.i = Inf,
  v_rel = NULL,
  anchor = NULL,
  search_f = search_geodesic,
  ...
)
```

Arguments

<code>index_f</code>	the section pursuit index function to optimise. The function needs to take three arguments, the projected data, the vector of distances from the current projection plane, and the slice thickness <code>h</code> .
<code>d</code>	target dimensionality
<code>alpha</code>	the initial size of the search window, in radians
<code>cooling</code>	the amount the size of the search window should be adjusted by after each step
<code>max.tries</code>	the maximum number of unsuccessful attempts to find a better projection before giving up
<code>max.i</code>	the maximum index value, stop search if a larger value is found
<code>v_rel</code>	relative volume of the slice. If not set, suggested value is calculated and printed to the screen.
<code>anchor</code>	A vector specifying the reference point to anchor the slice. If NULL (default) the slice will be anchored at the data center.
<code>search_f</code>	the search strategy to use
<code>...</code>	arguments sent to the <code>search_f</code>

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate_slice](#), [save_history](#) or [render](#).

See Also

[slice_index](#) for an example of an index functions. [search_geodesic](#), [search_better](#), [search_better_random](#) for different search strategies

Examples

```
# Generate samples on a 3d hollow sphere using the geozoo package
set.seed(12345)
sphere3 <- geozoo::sphere.hollow(3)$points
# Columns need to be named before launching the tour
colnames(sphere3) <- c("x1", "x2", "x3")
# Off-center anchoring
anchor3 <- matrix(rep(0.75, 3), ncol=3)
# Index setup
r_breaks <- linear_breaks(5, 0, 1)
a_breaks <- angular_breaks(10)
eps <- estimate_eps(nrow(sphere3), ncol(sphere3), 0.1 / 1, 5 * 10, 10, r_breaks)
idx <- slice_index(r_breaks, a_breaks, eps, bintype = "polar", power = 1, reweight = TRUE, p = 3)
# Running the guided section tour select sections showing a big hole in the center
animate_slice(sphere3, guided_section_tour(idx, v_rel = 0.1, anchor = anchor3, max.tries = 5),
  v_rel = 0.1, anchor = anchor3
)
```

 guided_tour

A guided tour path.

Description

Instead of choosing new projections at random like the grand tour, the guided tour always tries to find a projection that is more interesting than the current projection.

Usage

```
guided_tour(
  index_f,
  d = 2,
  alpha = 0.5,
  cooling = 0.99,
  max.tries = 25,
  max.i = Inf,
  search_f = search_geodesic,
  n_sample = 100,
  ...
)
```

Arguments

<code>index_f</code>	the index function to optimise.
<code>d</code>	target dimensionality
<code>alpha</code>	the initial size of the search window, in radians
<code>cooling</code>	the amount the size of the search window should be adjusted by after each step
<code>max.tries</code>	the maximum number of unsuccessful attempts to find a better projection before giving up
<code>max.i</code>	the maximum index value, stop search if a larger value is found
<code>search_f</code>	the search strategy to use: search_geodesic , search_better , search_better_random , search_polish . Default is search_geodesic .
<code>n_sample</code>	number of samples to generate if <code>search_f</code> is search_polish
<code>...</code>	arguments sent to the <code>search_f</code>

Details

Currently the index functions only work in 2d.

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

See Also

[cmass](#), [holes](#) and [lda_pp](#) for examples of index functions. The function should take a numeric matrix and return a single number, preferably between 0 and 1. [search_geodesic](#), [search_better](#), [search_better_random](#) for different search strategies

Examples

```
animate_xy(flea[, 1:6], guided_tour(holes()), sphere = TRUE)

animate_xy(flea[, 1:6], guided_tour(holes(), search_f = search_better_random), sphere = TRUE)
animate_dist(flea[, 1:6], guided_tour(holes(), 1), sphere = TRUE)
animate_xy(flea[, 1:6], guided_tour(lda_pp(flea$species)), sphere = TRUE, col = flea$species)

# save_history is particularly useful in conjunction with the
# guided tour as it allows us to look at the tour path in many different
# ways
f <- flea[, 1:3]
tries <- replicate(5, save_history(f, guided_tour(holes()))), simplify = FALSE)
```

holes

Holes index.

Description

Calculates the holes index. See Cook and Swayne (2007) Interactive and Dynamic Graphics for Data Analysis for equations.

Usage

```
holes()
```

interpolate

Interpolate geodesically between bases.

Description

This function takes a set of bases and produces a tour by geodesically interpolating between each basis

Usage

```
interpolate(basis_set, angle = 0.05, cycle = FALSE)
```

Arguments

basis_set	input basis set
angle	target distance (in radians) between bases
cycle	For <code>planned_tour</code> cycle through continuously (TRUE) or stop after first pass (FALSE)

Examples

```
t1 <- save_history(flea[, 1:6], grand_tour(1), max = 10)
dim(t1)
dim(interpolate(t1, 0.01))
dim(interpolate(t1, 0.05))
dim(interpolate(t1, 0.1))
```

Laser measurements *Turnable laser measurements from Bellcore*

Description

This data came from an investigation of an experimental laser at Bellcore. It was a tunable laser, in the sense that both its wavelength and power output were controllable.

Format

A 64 x 4 numeric array

Details

Rotation helped the experimental physicists to characterize the laser, which turned out not to be a very good one, due to its unstable operating region.

This data initially came to the statistics research group when Janette Cooper asked Paul Tukey to help her analyze the data she had collected to describe the laser.

- ifront, current applied to the front of the laser
- iback, current applied to the back of the laser
- power, output power
- lambda, output wavelength

Examples

```
head(laser)
animate_xy(laser[, -4])
```

lda_pp	<i>LDA projection pursuit index.</i>
--------	--------------------------------------

Description

Calculate the LDA projection pursuit index. See Cook and Swayne (2007) Interactive and Dynamic Graphics for Data Analysis for equations.

Usage

```
lda_pp(c1)
```

Arguments

c1	class to be used. Such as "color"
----	-----------------------------------

linear_breaks	<i>Returns n equidistant bins between a and b</i>
---------------	---------------------------------------------------

Description

Returns n equidistant bins between a and b

Usage

```
linear_breaks(n, a, b)
```

Arguments

n	number of bins
a	lower bound
b	upper bound

little_tour	<i>A little tour path.</i>
-------------	----------------------------

Description

The little tour is a planned tour that travels between all axis parallel projections. (John McDonald named this type of tour.)

Usage

```
little_tour(d = 2)
```

Arguments

d	target dimensionality
---	-----------------------

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
animate_xy(flea[, 1:6], little_tour())
animate_pcp(flea[, 1:6], little_tour(3))
animate_scatmat(flea[, 1:6], little_tour(3))
animate_pcp(flea[, 1:6], little_tour(4))
```

local_tour	<i>A local tour path.</i>
------------	---------------------------

Description

The local tour alternates between the starting position and a nearby random projection.

Usage

```
local_tour(start, angle = pi/4)
```

Arguments

start	initial projection matrix
angle	distance in radians to stay within

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
animate_xy(flea[, 1:3], local_tour(basis_init(3, 2)))
animate_xy(flea[, 1:3], local_tour(basis_init(3, 2), 0.2))
animate_xy(flea[, 1:3], local_tour(basis_random(3, 2), 0.2))
```

manual_slice	<i>Manually slice along a variable axis.</i>
--------------	----------------------------------------------

Description

The manual slice tour takes the current projection, with `display_slice`, and changes the slice center.

Usage

```
manual_slice(
  data,
  proj,
  var = 1,
  nsteps = 20,
  v_rel = 0.01,
  rescale = FALSE,
  sphere = FALSE,
  col = "black",
  half_range = NULL,
  anchor_nav = "topright",
  ...
)
```

Arguments

<code>data</code>	numeric matrix, with <code>n</code> rows and <code>p</code> columns
<code>proj</code>	projection from which slices are constructed
<code>var</code>	variable axis to run the center along: 1, ..., <code>p</code>
<code>nsteps</code>	number of changes in center to make
<code>v_rel</code>	relative volume of the slice. If not set, suggested value is calculated and printed to the screen.
<code>rescale</code>	if true, rescale all variables to range [0,1]?
<code>sphere</code>	if true, sphere all variables
<code>col</code>	color to be plotted. Defaults to "black"

half_range half range to use when calculating limits of projected. If not set, defaults to maximum distance from origin to each row of data.

anchor_nav position of the anchor: center, topright or off

... other options passed to output device

Examples

```
# Note that you might need to use the quartz()
# on OSX to see the animation
sphere5 <- data.frame(geozoo::sphere.hollow(5)$points)
proj <- basis_random(5, 2)
manual_slice(sphere5, proj, var=3, nsteps=10, rescale=TRUE, half_range=1.5)
```

mapColors *Map vector of factors to color*

Description

Map vector of factors to color

Usage

```
mapColors(x)
```

Arguments

x vector

norm_bin *Normality index.*

Description

Compares the similarity between the projected distribution and a normal distribution.

- norm_bin compares the count in 100 histogram bins
- norm_kol compares the cdf based on the Kolmogorov–Smirnov test (KS test)

Usage

```
norm_bin(nr)
```

```
norm_kol(nr)
```

Arguments

nr The number of rows in the target matrix

Examples

```
# manually compute the norm_kol index
# create the index function
set.seed(123)
index <- norm_kol(nrow(flea[, 1:3]))
# create the projection
proj <- matrix(c(1, 0, 0), nrow = 3)
# pre-process the example data
flea_s <- sphere_data(flea[, 1:3])
# produce the index value
index(flea_s %*% proj)
```

Olive oil measurements

Olive oil samples from Italy

Description

This data is from a paper by Forina, Armanino, Lanteri, Tiscornia (1983) Classification of Olive Oils from their Fatty Acid Composition, in Martens and Russwurm (ed) Food Research and Data Analysis. We thank Prof. Michele Forina, University of Genova, Italy for making this dataset available.

Format

A 572 x 10 numeric array

Details

- region Three super-classes of Italy: North, South and the island of Sardinia
- area Nine collection areas: three from North, four from South and 2 from Sardinia
- palmitic, palmitoleic, stearic, oleic, linoleic, linolenic, arachidic, eicosenoic fatty acids percent x 100

Examples

```
head(olive)
animate_xy(olive[, c(7, 9, 10)])
animate_xy(olive[, c(7, 9, 10)], col = olive[, 1])
```

Ozone measurements *Monthly ozone measurements over Central America*

Description

This data set is a subset of the data from the 2006 ASA Data expo challenge. The data are monthly ozone averages on a very coarse 24 by 24 grid covering Central America, from Jan 1995 to Dec 2000. The data is stored in a 3d area with the first two dimensions representing latitude and longitude, and the third representing time.

Format

A 24 x 24 x 72 numeric array

Examples

```
example(display_image)
```

path_curves *Draw the path that the geodesics took.*

Description

This computes the projected values of each observation at each step, and allows you to recreate static views of the animated plots.

Usage

```
path_curves(history, data = attr(history, "data"))
```

Arguments

history list of bases produced by `save_history` (or otherwise)
 data dataset to be projected on to bases

Examples

```
path1d <- save_history(flea[, 1:6], grand_tour(1), 3)
path2d <- save_history(flea[, 1:6], grand_tour(2), 3)

if (require("ggplot2")) {
  plot(path_curves(path1d))
  plot(path_curves(interpolate(path1d)))

  plot(path_curves(path2d))
}
```

```

plot(path_curves(interpolate(path2d)))

# Instead of relying on the built in plot method, you might want to
# generate your own. Here are few examples of alternative displays:

df <- path_curves(path2d)
ggplot(data = df, aes(x = step, y = value, group = obs:var, colour = var)) +
  geom_line() +
  facet_wrap(~obs)

library(tidyr)
ggplot(
  data = pivot_wider(df,
    id_cols = c(obs, step),
    names_from = var, names_prefix = "Var",
    values_from = value
  ),
  aes(x = Var1, y = Var2)
) +
  geom_point() +
  facet_wrap(~step) +
  coord_equal()
}

```

path_dist

Compute distance matrix from bases.

Description

Compute distance matrix from bases.

Usage

```
path_dist(history)
```

Arguments

history history of the plots

Examples

```

# This code is to be used as an example but you should increase
# the max from 2 to 50, say, to check tour coverage.
grand <- interpolate(save_history(flea[, 1:6], max = 2), 0.2)
# The grand tour -----
# Look at the tour path in a tour, how well does it cover a sphere
# Using MDS to summarise the high-d space of projections
d <- path_dist(grand)
ord <- as.data.frame(MASS::isoMDS(d)$points)
require(ggplot2)

```

```

ggplot(data = ord, aes(x=V1, y=V2)) +
  geom_path() +
  coord_equal() +
  labs(x = NULL, y = NULL)

# Compare five guided tours -----
holes1d <- guided_tour(holes(), 1)
tour_reps <- replicate(5, save_history(flea[, 1:6], holes1d, max = 2),
  simplify = FALSE
)
tour_reps2 <- lapply(tour_reps, interpolate, 0.2)

bases <- unlist(lapply(tour_reps2, as.list), recursive = FALSE)
class(bases) <- "history_list"
index_values <- paths_index(tour_reps2, holes())
d <- path_dist(bases)
ord <- as.data.frame(cmdscale(d, 2))

info <- cbind(ord, index_values)
if (require("ggplot2")) {
  ggplot(data = info, aes(x = step, y = value, group = try)) +
    geom_line()
  ##ggplot(data = info, aes(x = V1, y = V2, group = try)) +
  ## geom_path() +
  ## geom_point(aes(size = value)) +
  ## coord_equal()
  ##last_plot() + facet_wrap(~try)
}

```

path_index

Compute index values for a tour history.

Description

Compute index values for a tour history.

Usage

```
path_index(history, index_f, data = attr(history, "data"))
```

Arguments

history	list of bases produced by save_history (or otherwise)
index_f	index function to apply to each basis
data	dataset to be projected on to bases

See Also

[save_history](#) for options to save history

Examples

```

fl_holes <- save_history(flea[, 1:6], guided_tour(holes()), sphere = TRUE)
path_index(fl_holes, holes())
## path_index(fl_holes, cmass())

plot(path_index(fl_holes, holes()), type = "l")
## plot(path_index(fl_holes, cmass()), type = "l")

# Use interpolate to show all intermediate bases as well
hi <- path_index(interpolate(fl_holes), holes())
hi
plot(hi)

```

pda_pp

PDA projection pursuit index.

Description

Calculate the PDA projection pursuit index. See Lee and Cook (2009) A Projection Pursuit Index for Large p, Small n Data

Usage

```
pda_pp(c1, lambda = 0.2)
```

Arguments

c1	class to be used. Such as "color"
lambda	shrinkage parameter (0 = no shrinkage, 1 = full shrinkage)

Places Ratings

Ratings of different locations across North America

Description

The "places data" were distributed to interested ASA members a few years ago so that they could apply contemporary data analytic methods to describe these data and then present results in a poster session at the ASA annual conference. Latitude and longitude have been added by Paul Tukey.

Format

A 329 x 14 numeric array

Details

The first dataset is taken from the Places Rated Almanac, by Richard Boyer and David Savageau, copyrighted and published by Rand McNally. This book order (SBN) number is 0-528-88008-X, and it retails for \$14.95 . The data are reproduced on disk by kind permission of the publisher, and with the request that the copyright notice of Rand McNally, and the names of the authors appear in any paper or presentation using these data.

The nine rating criteria used by Places Rated Almanac are: Climate and Terrain Housing Health Care and Environment Crime Transportation Education The Arts Recreation Economics

For all but two of the above criteria, the higher the score, the better. For Housing and Crime, the lower the score the better.

The scores are computed using the following component statistics for each criterion (see the Places Rated Almanac for details):

Climate and Terrain: very hot and very cold months, seasonal temperature variation, heating- and cooling-degree days, freezing days, zero-degree days, ninety-degree days.

Housing: utility bills, property taxes, mortgage payments.

Health Care and Environment: per capita physicians, teaching hospitals, medical schools, cardiac rehabilitation centers, comprehensive cancer treatment centers, hospices, insurance/hospitalization costs index, flouridation of drinking water, air pollution.

Crime: violent crime rate, property crime rate.

Transportation: daily commute, public transportation, Interstate highways, air service, passenger rail service.

Education: pupil/teacher ratio in the public K-12 system, effort index in K-12, accademic options in higher education.

The Arts: museums, fine arts and public radio stations, public television stations, universities offering a degree or degrees in the arts, symphony orchestras, theatres, opera companies, dance companies, public libraries.

Recreation: good restaurants, public golf courses, certified lanes for tenpin bowling, movie theatres, zoos, aquariums, family theme parks, sanctioned automobile race tracks, pari-mutuel betting attractions, major- and minor- league professional sports teams, NCAA Division I football and basketball teams, miles of ocean or Great Lakes coastline, inland water, national forests, national parks, or national wildlife refuges, Consolidated Metropolitan Statistical Area access.

Economics: average household income adjusted for taxes and living costs, income growth, job growth.

Examples

```
head(places)
animate_xy(places[, 1:9])
```

planned_tour	<i>A planned tour path.</i>
--------------	-----------------------------

Description

The planned tour takes you from one basis to the next in a set order. Once you have visited all the planned bases, you either stop or start from the beginning once more (if `cycle = TRUE`).

Usage

```
planned_tour(basis_set, cycle = FALSE)
```

Arguments

<code>basis_set</code>	the set of bases as a list of projection matrices or a 3d array
<code>cycle</code>	cycle through continuously (<code>TRUE</code>) or stop after first pass (<code>FALSE</code>)

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

See Also

The [little_tour](#), a special type of planned tour which cycles between all axis parallel projections.

Examples

```
twod <- save_history(flea[, 1:3], max = 5)
str(twod)
animate_xy(flea[, 1:3], planned_tour(twod))
animate_xy(flea[, 1:3], planned_tour(twod, TRUE))

oned <- save_history(flea[, 1:6], grand_tour(1), max = 3)
animate_dist(flea[, 1:6], planned_tour(oned))
```

proj_dist	<i>Calculate the distance between two bases.</i>
-----------	--------------------------------------------------

Description

Computes the Frobenius norm between two bases, in radians. This is equals to the Euclidean norm of the vector of principal angles between the two subspaces.

Usage

```
proj_dist(x, y)
```

Arguments

x projection matrix a
y projection matrix b

radial_tour *A radial tour path.*

Description

The radial tour rotates a chosen variable axis out of the current projection.

Usage

```
radial_tour(start, mvar = 1, ...)
```

Arguments

start initial projection matrix
mvar variable chosen to rotate out
... additional arguments for drawing

Details

Usually, you will not call this function directly, but will pass it to a method that works with tour paths like [animate](#), [save_history](#) or [render](#).

Examples

```
animate_xy(flea[, 1:6], radial_tour(basis_random(6, 2), mvar = 4))
animate_dist(flea[, 1:6], radial_tour(basis_random(6, 1), mvar = 4))
animate_scatmat(flea[, 1:6], radial_tour(basis_random(6, 3), mvar = 4))
```

Rat CNS *Rat CNS Gene Expression*

Description

Columns:

Format

A 112 x 11 numeric array

Details

e11 e13 e15 e18 e21 p0 p7 p14 a class1 class2

- e11, an embryonic timepoint from the original data with the number corresponding to the day
- e13, an embryonic timepoint from the original data with the number corresponding to the day
- e15, an embryonic timepoint from the original data with the number corresponding to the day
- e18, an embryonic timepoint from the original data with the number corresponding to the day
- e21, an embryonic timepoint from the original data with the number corresponding to the day
- p0, a postnatal timepoint from the original data with the number corresponding to the day
- p7, a postnatal timepoint from the original data with the number corresponding to the day
- p14, a postnatal timepoint from the original data with the number corresponding to the day
- a, a postnatal timepoint from the original data. It is equivalent to p90.
- class1, is the high-level class: its range is 1:4
- class2, breaks down the high-level classes, so its range is 1:14

Rows: Each case is a gene (or gene family?) And each cell is the gene expression level for that gene at time t, averaging a few measured values and normalizing using the maximum expression value for that gene.

Reference (available on the web at [pnas.org](https://www.pnas.org)): Large-scale temporal gene expression mapping of central nervous system development by X. Wen, S. Fuhrman, G. S. Michaels, D. B. Carr, S. Smith, J. L. Barker, R. Somogyi in the Proceedings of the National Academy of Science, Vol 95, pp. 334-339, January 1998

References

<https://www.pnas.org>

Examples

```
head(ratcns)
animate_xy(ratcns[, 1:8], col = ratcns[, 10])
```

render

Render frames of animation to disk

Description

Render frames of animation to disk

Usage

```
render(
  data,
  tour_path,
  display,
  dev,
  ...,
  apf = 1/10,
  frames = 50,
  rescale = TRUE,
  sphere = FALSE,
  start = NULL
)
```

Arguments

<code>data</code>	matrix, or data frame containing numeric columns
<code>tour_path</code>	tour path generator
<code>display</code>	the method used to render the projected data, e.g. display_xy , display_pcp
<code>dev</code>	name of output device to use (e.g. png , pdf)
<code>...</code>	other options passed to output device
<code>apf</code>	angle (in radians) per frame
<code>frames</code>	number of frames in output
<code>rescale</code>	if true, rescale all variables to range [0,1]
<code>sphere</code>	if true, sphere all variables
<code>start</code>	starting projection. If NULL, uses path default.

Examples

```
tmp_path <- tempdir()
render(flea[, 1:6], grand_tour(), display_xy(), "pdf",
  frames = 3,
  file.path(tmp_path, "test.pdf")
)
render(flea[, 1:6], grand_tour(), display_xy(), "png",
  frames = 3,
  file.path(tmp_path, "test-%03d.png")
)
```

render_gif	<i>Render frames of animation to a gif file</i>
------------	-------------------------------------------------

Description

Render frames of animation to a gif file

Usage

```
render_gif(
  data,
  tour_path,
  display,
  gif_file = "animation.gif",
  ...,
  apf = 1/10,
  frames = 50,
  rescale = TRUE,
  sphere = FALSE,
  start = NULL
)
```

Arguments

data	matrix, or data frame containing numeric columns
tour_path	tour path generator
display	the method used to render the projected data, e.g. display_xy , display_pcp
gif_file	Name of gif file (default = "animation.gif")
...	other options passed to png
apf	angle (in radians) per frame
frames	number of frames in output
rescale	if true, rescale all variables to range [0,1]
sphere	if true, sphere all variables
start	starting projection. If NULL, uses path default.

Examples

```
## Not run:
# gifski needs to be installed to render a gif
if (requireNamespace("gifski", quietly = TRUE)) {
  gif_file <- file.path(tempdir(), "test.gif")
  render_gif(flea[, 1:6], grand_tour(), display_xy(), gif_file)
  utils::browseURL(gif_file)
  unlink(gif_file)
}

## End(Not run)
```

rescale	<i>Rescale a matrix or data frame</i>
---------	---------------------------------------

Description

Standardise each column to have range [0, 1]

Usage

```
rescale(df)
```

Arguments

df	data frame or matrix
----	----------------------

save_history	<i>Save tour history.</i>
--------------	---------------------------

Description

Save a tour path so it can later be displayed in many different ways.

Usage

```
save_history(
  data,
  tour_path = grand_tour(),
  max_bases = 100,
  start = NULL,
  rescale = TRUE,
  sphere = FALSE,
  step_size = Inf,
  ...
)
```

Arguments

data	matrix, or data frame containing numeric columns
tour_path	tour path generator
max_bases	maximum number of new bases to generate. Some tour paths (like the guided tour) may generate less than the maximum.
start	starting projection, if you want to specify one
rescale	if true, rescale all variables to range [0,1]?
sphere	if true, sphere all variables


```

step_size      distance between each step - defaults to Inf which forces new basis generation
                at each step.
...           additional arguments passed to tour path

```

Examples

```

# You can use a saved history to replay tours with different visualisations

t1 <- save_history(flea[, 1:6], max = 3)
animate_xy(flea[, 1:6], planned_tour(t1))
## andrews_history(t1)
## andrews_history(interpolate(t1))

## t1 <- save_history(flea[, 1:6], grand_tour(4), max = 3)
## animate_pcp(flea[, 1:6], planned_tour(t1))
## animate_scatter(flea[, 1:6], planned_tour(t1))

## t1 <- save_history(flea[, 1:6], grand_tour(1), max = 3)
## animate_dist(flea[, 1:6], planned_tour(t1))

testdata <- matrix(rnorm(100 * 3), ncol = 3)
testdata[1:50, 1] <- testdata[1:50, 1] + 10
testdata <- sphere_data(testdata)
t2 <- save_history(testdata, guided_tour(holes(), max.tries = 10),
  max = 5, rescale = FALSE
)
animate_xy(testdata, planned_tour(t2))

# Or you can use saved histories to visualise the path that the tour took.
plot(path_index(interpolate(t2), holes()))

```

search_better

Search for a better projection near the current projection.

Description

Search for a better projection near the current projection.

Usage

```

search_better(
  current,
  alpha = 0.5,
  index,
  tries,
  max.tries = Inf,
  ...,
  method = "linear",
  cur_index = NA
)

```

Arguments

current	starting projection
alpha	the angle used to search the target basis from the current basis
index	index function
tries	the counter of the outer loop of the optimiser
max.tries	maximum number of iteration before giving up
...	other arguments being passed into the search_better()
method	whether the nearby bases are found by a linear/ geodesic formulation
cur_index	the index value of the current basis

Examples

```
animate_xy(flea[, 1:6], guided_tour(holes(), search_f = search_better))
```

search_better_random *Search for a better projection using simulated annealing*

Description

Given an initial t_0 , the cooling scheme updates temperature at

$$T = t_0 / \log(i + 1)$$

The candidate basis is sampled via

$$B_j = (1 - \alpha) * B_i + \alpha * B$$

where alpha defines the neighbourhood, B_i is the current basis, B is a randomly generated basis
The acceptance probability is calculated as

$$prob = \exp(-abs(I(B_i) - I(B_j)) / T)$$

For more information, see https://projecteuclid.org/download/pdf_1/euclid.ss/1177011077

Usage

```
search_better_random(
  current,
  alpha = 0.5,
  index,
  tries,
  max.tries = Inf,
  method = "linear",
  cur_index = NA,
  t0 = 0.01,
  ...
)
```

Arguments

current	starting projection
alpha	the angle used to search the target basis from the current basis
index	index function
tries	the counter of the outer loop of the optimiser
max.tries	maximum number of iteration before giving up
method	whether the nearby bases are found by a linear/ geodesic formulation
cur_index	the index value of the current basis
t0	initial decrease in temperature
...	other arguments being passed into the search_better_random()

Examples

```
animate_xy(flea[, 1:6], guided_tour(holes(), search_f = search_better_random))
```

search_geodesic *A pseudo-derivative, line search algorithm.*

Description

This is a novel method for finding more interesting projections for the guided tour. It works by first taking a small step in n random directions, and then picking the direction that looks most promising (based on the height of the index function), which is effectively a gradient search. Then it performs a linear search along the geodesic in that direction, traveling up to half way around the sphere.

Usage

```
search_geodesic(
  current,
  alpha = 1,
  index,
  tries,
  max.tries = 5,
  ...,
  n = 5,
  delta = 0.01,
  cur_index = NA
)
```

Arguments

current	starting projection
alpha	maximum distance to travel (currently ignored)
index	interestingness index function
tries	the counter of the outer loop of the optimiser
max.tries	maximum number of failed attempts before giving up
...	other arguments being passed into the search_geodesic()
n	number of random steps to take to find best direction
delta	step size for evaluation of best direction
cur_index	index value for starting projection, set NA if it needs to be calculated

Details

You should not to have call this function directly, but should supply it to the [guided_tour](#) as a search strategy.

Examples

```
animate_xy(flea[, 1:6], guided_tour(holes(), search_f = search_geodesic))
```

search_polish	<i>Search very locally to find slightly better projections to polish a broader search.</i>
---------------	--------------------------------------------------------------------------------------------

Description

Search very locally to find slightly better projections to polish a broader search.

Usage

```
search_polish(
  current,
  alpha = 0.5,
  index,
  tries,
  polish_max_tries = 30,
  cur_index = NA,
  n_sample = 100,
  polish_cooling = 1,
  ...
)
```

Arguments

current	the current projection basis
alpha	the angle used to search the target basis from the current basis
index	index function
tries	the counter of the outer loop of the optimiser
polish_max_tries	maximum number of iteration before giving up
cur_index	the index value of the current basis
n_sample	number of samples to generate
polish_cooling	percentage of reduction in polish_alpha when no better basis is found
...	other arguments being passed into the search_polish()

Examples

```

data(t1)
best_proj <- t1[, , dim(t1)[3]]
attr(best_proj, "data") <- NULL
best_proj <- unclass(drop(best_proj))
animate_xy(
  flea[, 1:6],
  guided_tour(holes()),
  search_f = search_polish(
    polish_max_tries = 5),
  start = best_proj
)

```

search_posse

Search for a better projection based on Poss, 1995

Description

Search for a better projection based on Poss, 1995

Usage

```

search_posse(
  current,
  alpha = 0.5,
  index,
  tries,
  max.tries = 300,
  cur_index = NA,
  ...
)

```

Arguments

current	starting projection
alpha	the angle used to search the target basis from the current basis
index	index function
tries	the counter of the outer loop of the optimiser
max.tries	maximum number of iteration before giving up
cur_index	the index value of the current basis
...	other arguments being passed into the search_better()

 slice_index

Section pursuit index.

Description

Calculates a section pursuit index that compares the distribution inside and outside a slice.

Usage

```

slice_index(
  breaks_x,
  breaks_y,
  eps,
  bintype = "polar",
  power = 1,
  flip = 1,
  reweight = FALSE,
  p = 4
)

```

Arguments

breaks_x	binning on the first variable (x or radius).
breaks_y	binning on the second variable (y or angle).
eps	cutoff values to suppress summing up small differences. Vector with one entry for each bin, can be estimated using estimate_eps .
bintype	select polar (default) or square binning.
power	exponent q used in the index computation.
flip	sign of the index computation, select +1 when searching for low densities and -1 when searching for high densities.
reweight	if TRUE will reweight according to the expected distribution in a uniform hypersphere (default is FALSE).
p	number of variables in the data (needed for accurate reweighting, default is 4).

sphere_data	<i>Sphere a matrix (or data frame) by transforming variables to principal components.</i>
-------------	-------------------------------------------------------------------------------------------

Description

Sphering is often useful in conjunction with the guided tour, as it removes simpler patterns that may conceal more interesting findings.

Usage

```
sphere_data(df)
```

Arguments

df	data frame or matrix
----	----------------------

splines2d	<i>Spline based index.</i>
-----------	----------------------------

Description

Compares the variance in residuals of a fitted spline model to the overall variance to find functional dependence in 2D projections of the data.

Usage

```
splines2d()
```

t1	<i>Saved history of guided tour with holes</i>
----	------------------------------------------------

Description

This data was generated from the following code: `set.seed(2020) t1 <- save_history(flea[, 1:6], guided_tour(holes()), max = 100) attr(t1, "class") <- NULL` And used as an example for `search_polish()` to start optimising from the best projection from `search_geodesic`. `t1` is a 3D array or 2D projections.

Tropical Atmosphere Ocean

Tropical Atmosphere Ocean data

Description

This is a subset of data taken from the NOAA web site <https://www.pmel.noaa.gov/tao/>. The data is generated from recording instruments on a grid of buoys laid out over the Pacific Ocean. The grid was setup to monitor El Nino and La Nina events. This subset contains measurements from 5 locations (0deg/110W, 2S/110W, 0deg/95W, 2S/95W, 5S/95W) and two time points Nov-Jan 1993 (normal), 1997 (El Nino). There are missing values in this data set, which need to be removed, or imputed before running a tour.

Format

A 736 x 8 numeric array

References

<https://www.pmel.noaa.gov/tao/>

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