

Package ‘EmbedSOM’

February 12, 2020

Version 2.1.1

Title Fast Embedding Guided by Self-Organizing Map

Depends R (>= 3.2)

Suggests knitr, rmarkdown

Imports FNN, ggplot2, igraph, Matrix, Rtsne, umap, uwot

Description Provides a smooth mapping of multidimensional points into low-dimensional space defined by a self-organizing map. Designed to work with 'FlowSOM' and flow-cytometry use-cases. See Kratochvil et al. (2019) <doi:10.1101/496869>.

License GPL (>= 3)

LazyData true

URL <https://bioinfo.uochb.cas.cz/embedsom>

Encoding UTF-8

RoxygenNote 6.1.1

VignetteBuilder knitr

NeedsCompilation yes

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Repository CRAN

Date/Publication 2020-02-12 12:20:10 UTC

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ClusterPalette	<i>An acceptable cluster color palette</i>
----------------	--

Description

An acceptable cluster color palette

Usage

```
ClusterPalette(n, vcycle = c(1, 0.7), scycle = c(0.7, 1), alpha = 1)
```

Arguments

n	How many colors to generate
vcycle, scycle	Small vectors with cycles of saturation/value for hsv
alpha	Opacity of the colors

Examples

```
EmbedSOM::ClusterPalette(10)
```

EmbedSOM

*Process the cells with SOM into a nice embedding***Description**

Process the cells with SOM into a nice embedding

Usage

```
EmbedSOM(data = NULL, map = NULL, fsom = NULL, smooth = NULL,
          k = NULL, adjust = NULL, importance = NULL, coordsFn = NULL,
          coords = NULL, emcoords = NULL, emcoords.pow = 1, parallel = F,
          threads = if (parallel) 0 else 1)
```

Arguments

data	Data matrix with points that optionally overrides the one from fsom\$data
map	Map object in FlowSOM format, to optionally override fsom\$map
fsom	FlowSOM object with a built SOM (used if data or map are missing)
smooth	Produce smoother (positive values) or more rough approximation (negative values).
k	How many neighboring landmarks (e.g. SOM nodes) to take into the whole computation
adjust	How much non-local information to remove from the approximation
importance	Scaling of the landmarks, will be used to scale the incoming data (should be same as used for training the SOM or to select the landmarks)
coordsFn	A coordinates-generating function (e.g. <code>tSNECoords()</code>) that overrides the existing map\$grid.
coords	A matrix of embedding-space coordinates that correspond to map\$codes (i.e. the "embedded landmarks"). Overrides map\$grid if not NULL.
emcoords	Provided for backwards compatibility, will be removed. Use coords and coordsFn instead.
emcoords.pow	Provided for backwards compatibility, will be removed. Use a parametrized coordsFn instead.
parallel	Boolean flag whether the computation should be parallelized (this flag is just a nice name for threads and does not do anything directly – default FALSE sets threads=1, TRUE sets threads=0)
threads	Number of threads used for computation, 0 chooses hardware concurrency, 1 (default) turns off parallelization.

Value

matrix with 2D or 3D coordinates of the embedded data, depending on the map

Examples

```
d <- cbind(rnorm(10000), 3*runif(10000), rexp(10000))
colnames(d) <- paste0("col",1:3)
map <- EmbedSOM::SOM(d, xdim=10, ydim=10)
e <- EmbedSOM::EmbedSOM(data=d, map=map)
EmbedSOM::PlotEmbed(e, data=d, 'col1', pch=16)
```

ExprColors	<i>Generate colors for multi-color marker expression labeling in a single plot</i>
------------	--

Description

Generate colors for multi-color marker expression labeling in a single plot

Usage

```
ExprColors(exprs, base = exp(1), scale = 1, cutoff = 0, pow = NULL,
  col = ClusterPalette(dim(exprs)[2], alpha = alpha),
  nocolor = grDevices::rgb(0.75, 0.75, 0.75, alpha/2), alpha = 0.5)
```

Arguments

exprs	Matrix-like object with marker expressions (extract it manually from your data)
base, scale	Base(s) and scale(s) for softmax (convertible to numeric vectors of size 1+ncol(exprs))
cutoff	Gray level (expressed in sigmas of the sample distribution)
pow	Obsolete, now renamed to scale.
col	Colors to use, defaults to colors taken from 'ClusterPalette'
nocolor	The color to use for sub-gray-level expression, default gray.
alpha	Default alpha value.

Examples

```
d <- cbind(rnorm(1e5), rexp(1e5))
EmbedSOM::PlotEmbed(d, col=EmbedSOM::ExprColors(d, pow=2))
```

ExpressionGradient *The ggplot2 scale gradient from ExpressionPalette.*

Description

The ggplot2 scale gradient from ExpressionPalette.

Usage

```
ExpressionGradient(...)
```

Arguments

... Arguments passed to `ggplot2::scale_color_gradientn()`

Examples

```
library(EmbedSOM)
library(ggplot2)

# simulate a simple dataset
e <- cbind(rnorm(10000), rnorm(10000))

data <- data.frame(Val=log(1+e[,1]^2+e[,2]^2))
PlotGG(e, data=data) +
  geom_point(aes_string(color="Val"), alpha=.5) +
  ExpressionGradient(guide=FALSE)
```

ExpressionPalette *Marker expression palette generator based off ColorBrewer's RdYlBu, only better for plotting of half-transparent cells*

Description

Marker expression palette generator based off ColorBrewer's RdYlBu, only better for plotting of half-transparent cells

Usage

```
ExpressionPalette(n, alpha = 1)
```

Arguments

n How many colors to generate
alpha Opacity of the colors

Examples

```
EmbedSOM::ExpressionPalette(10)
```

GQTSOM

Train a Growing Quadtree Self-Organizing Map

Description

Train a Growing Quadtree Self-Organizing Map

Usage

```
GQTSOM(data, init.dim = c(3, 3), target_codes = 100, rlen = 10,
        radius = c(sqrt(sum(init.dim^2)), 0.5), epochRadii = seq(radius[1],
        radius[2], length.out = rlen), coords = NULL, codes = NULL,
        coordsFn = NULL, importance = NULL, distf = 2, nhbr.distf = 2,
        noMapping = F, parallel = F, threads = if (parallel) 0 else 1)
```

Arguments

<code>data</code>	Input data matrix
<code>init.dim</code>	Initial size of the SOM, default <code>c(3, 3)</code>
<code>target_codes</code>	Make the SOM grow linearly to at most this amount of nodes (default 100)
<code>rlen</code>	Number of training iterations
<code>radius</code>	Start and end training radius, as in <code>SOM()</code>
<code>epochRadii</code>	Precise radii for each epoch (must be of length <code>rlen</code>)
<code>coords</code>	Quadtree coordinates of the initial SOM nodes.
<code>codes</code>	Initial codebook
<code>coordsFn</code>	Function to generate/transform grid coordinates (e.g. <code>tSNECoords()</code>). If NULL (default), the grid is the 2D coordinates of GQTSOM map.
<code>importance</code>	Weights of input data dimensions
<code>distf</code>	Distance measure to use in input data space (1=manhattan, 2=euclidean, 3=chebyshev, 4=cosine)
<code>nhbr.distf</code>	Distance measure to use in output space (as in <code>distf</code>)
<code>noMapping</code>	If TRUE, do not compute the assignment of input data to SOM nodes
<code>parallel</code>	Parallelize the training by setting appropriate threads. Defaults to FALSE.
<code>threads</code>	Number of threads to use for training. Defaults to 0 (chooses maximum available hardware threads) if <code>parallel=TRUE</code> or 1 (single thread) if <code>parallel=FALSE</code> .

GraphCoords

Add Kamada-Kawai-generated embedding coordinates to the map

Description

This uses a complete graph on the map codebook, which brings overcrowding problems. It is therefore useful to transform the distances for avoiding that (e.g. by exponentiating them slightly).

Usage

```
GraphCoords(dim = NULL, dist.method = NULL, distFn = function(x) x,
            layoutFn = igraph::layout_with_kk)
```

Arguments

dim	Dimension of the result (passed to layoutFn)
dist.method	The method to compute distances, passed to <code>stats::dist()</code> as parameter method
distFn	Custom transformation function of the distance matrix
layoutFn	iGraph-compatible graph layouting function (default <code>igraph::layout_with_kk</code>)

Value

a function that transforms the map, usable as `coordsFn` parameter

Initialize_PCA

Create a grid from first 2 PCA components

Description

Create a grid from first 2 PCA components

Usage

```
Initialize_PCA(data, xdim, ydim, zdim = NULL)
```

Arguments

data	matrix in which each row represents a point
xdim, ydim, zdim	Dimensions of the SOM grid

Value

array containing the selected selected rows

kMeansMap	<i>Create a map from k-Means clusters</i>
-----------	---

Description

May give better results than 'RandomMap' on data where random sampling is complicated. This does not use actual kMeans clustering, but re-uses the batch version of [SOM\(\)](#) with tiny radius (which makes it work the same as kMeans). In consequence, the speedup of SOM function is applied here as well. Additionally, because we don't need that amount of clustering precision, parameters 'batch=F, rlen=1' may give a satisfactory result very quickly.

Usage

```
kMeansMap(data, k, coordsFn, batch = T, ...)
```

Arguments

data	Input data matrix, with individual data points in rows
k	How many points to sample
coordsFn	a function to generate embedding coordinates (default none)
batch	Use batch-SOM training (effectively kMeans, default TRUE)
...	Passed to SOM() , useful e.g. for 'parallel=T' or 'rlen=5'

Value

map object (without the grid, if coordsFn was not specified)

Examples

```
d <- iris[,1:4]
EmbedSOM::PlotEmbed(
  EmbedSOM::EmbedSOM(
    data = d,
    map = EmbedSOM::kMeansMap(d, 10, EmbedSOM::GraphCoords()),
    pch=19, clust=iris[,5]
  )
)
```

kNNCoords	<i>Add KNN-topology-based embedding coordinates to the map</i>
-----------	--

Description

Internally, this uses `FNN::get.knn()` to compute the k-neighborhoods. That function only supports Euclidean metric, therefore `kNNCoords` throws a warning whenever a different metric is used.

Usage

```
kNNCoords(k = 4, dim = NULL, distFn = function(x) x,
  layoutFn = igraph::layout_with_kk)
```

Arguments

<code>k</code>	Size of the neighborhoods (default 4)
<code>dim</code>	Dimension of the result (passed to <code>layoutFn</code>)
<code>distFn</code>	Custom transformation function of the distance matrix
<code>layoutFn</code>	igraph-compatible graph layouting function (default <code>igraph::layout_with_kk</code>)

Value

a function that transforms the map, usable as `coordsFn` parameter

MapDataToCodes	<i>Assign nearest node to each datapoint</i>
----------------	--

Description

Assign nearest node to each datapoint

Usage

```
MapDataToCodes(codes, data, distf = 2, parallel = F, threads = if
  (parallel) 0 else 1)
```

Arguments

<code>codes</code>	matrix with nodes of the SOM
<code>data</code>	datapoints to assign
<code>distf</code>	Distance function (1=manhattan, 2=euclidean, 3=chebyshev, 4=cosine)
<code>threads, parallel</code>	Use parallel computation (see <code>SOM()</code>)

Value

array with nearest node id for each datapoint

MSTCoords *Add MST-style embedding coordinates to the map*

Description

Add MST-style embedding coordinates to the map

Usage

```
MSTCoords(dim = NULL, dist.method = NULL, distFn = function(x) x,
  layoutFn = igraph::layout_with_kk)
```

Arguments

dim	Dimension of the result (passed to layoutFn)
dist.method	The method to compute distances, passed to <code>stats::dist()</code> as parameter method
distFn	Custom transformation function of the distance matrix
layoutFn	iGraph-compatible graph layouting function (default <code>igraph::layout_with_kk()</code>)

Value

a function that transforms the map, usable as `coordsFn` parameter

NormalizeColor *Helper for computing colors for embedding plots*

Description

Helper for computing colors for embedding plots

Usage

```
NormalizeColor(data, low = NULL, high = NULL, pow = 0, sds = 1)
```

Arguments

data	Vector of scalar values to normalize between 0 and 1
low, high	Originally quantiles for clamping the color. Only kept for backwards compatibility, now ignored.
pow	The scaled data are transformed to $\text{data}^{(2^{\text{pow}})}$. If set to 0, nothing happens. Positive values highlight differences in the data closer to 1, negative values highlight differences closer to 0.
sds	Inverse scale factor for measured standard deviation (greater value makes data look more extreme)

Examples

```
EmbedSOM::NormalizeColor(c(1,100,500))
```

PlotData	<i>Export a data frame for plotting with marker intensities and density.</i>
----------	--

Description

Export a data frame for plotting with marker intensities and density.

Usage

```
PlotData(embed, fsom, data = fsom$data, cols, names, normalize = cols,
  pow = 0, sds = 1, vf = PlotId, density = "Density",
  densBins = 256, densLimit = NULL, fdens = sqrt)
```

Arguments

embed, fsom, data, cols	The embedding data, columns to select
names	Column names for output
normalize	List of columns to normalize using NormalizeColor() , default all
pow, sds	Parameters for the normalization
vf	Custom value-transforming function
density	Name of the density column
densBins	Number of bins for density calculation
densLimit	Upper limit of density (prevents outliers)
fdens	Density-transforming function; default sqrt

PlotDefault	<i>Default plot</i>
-------------	---------------------

Description

Default plot

Usage

```
PlotDefault(pch = ".", cex = 1, ...)
```

Arguments

pch, cex, ...	correctly defaulted and passed to 'plot'
---------------	--

PlotEmbed

*Helper function for plotting the embedding***Description**

Convenience plotting function. Takes the embed matrix which is the output of [EmbedSOM\(\)](#), together with a multitude of arguments that set how the plotting is done.

Usage

```
PlotEmbed(embed, value = 0, red = 0, green = 0, blue = 0,
  fr = PlotId, fg = PlotId, fb = PlotId, fv = PlotId, powr = 0,
  powg = 0, powb = 0, powv = 0, sdsr = 1, sdsg = 1, sdsb = 1,
  sdsv = 1, clust = NULL, nbin = 256, maxDens = NULL,
  fdens = sqrt, limit = NULL, alpha = NULL, fsom, data, col,
  cluster.colors = ClusterPalette,
  expression.colors = ExpressionPalette,
  na.color = grDevices::rgb(0.75, 0.75, 0.75, if (is.null(alpha)) 0.5
  else alpha/2), plotf = PlotDefault, ...)
```

Arguments

embed	The embedding from EmbedSOM() , or generally any 2-column matrix of coordinates
value	The column of data to use for coloring the plotted points
red, green, blue	The same, for individual RGB components
fv, fr, fg, fb	Functions to transform the values before they are normalized
powv, powr, powg, powb	Passed to corresponding NormalizeColor() calls as pow
sdsv, sdsr, sdsg, sdsb	Passed to NormalizeColor() as sds
clust	Cluster labels (used as a factor)
nbin, maxDens, fdens	Parameters of density calculation, see PlotData()
limit	Low/high offset for NormalizeColor() (obsolete&ignored, will be removed)
alpha	Default alpha value of points
fsom	FlowSOM object
data	Data matrix, taken from fsom parameter by default
col	Overrides the computed point colors with exact supplied colors.
cluster.colors	Function to generate cluster colors, default ClusterPalette()
expression.colors	Function to generate expression color scale, default ExpressionPalette()

na.color	Color to assign to NA values
plotf	Plot function, defaults to <code>graphics::plot()</code> slightly decorated with <code>pch='.'</code> , <code>cex=1</code>
...	Extra params passed to the plot function

Examples

```
EmbedSOM::PlotEmbed(cbind(rnorm(1e5),rnorm(1e5)))
```

PlotGG	<i>Wrap PlotData result in ggplot object.</i>
--------	---

Description

This creates a ggplot2 object for plotting.

Usage

```
PlotGG(embed, ...)
```

Arguments

embed	Embedding data
...	Extra arguments passed to <code>PlotData()</code>

Examples

```
library(EmbedSOM)
library(ggplot2)

# simulate a simple dataset
e <- cbind(rnorm(10000),rnorm(10000))

PlotGG(e, data=data.frame(Expr=runif(10000))) +
  geom_point(aes_string(color="Expr"))
```

PlotId	<i>Identity on whatever</i>
--------	-----------------------------

Description

Identity on whatever

Usage

```
PlotId(x)
```

Arguments

x Just the x.

Value

The x.

RandomMap

Create a map by randomly selecting points

Description

Create a map by randomly selecting points

Usage

```
RandomMap(data, k, coordsFn)
```

Arguments

data Input data matrix, with individual data points in rows
k How many points to sample
coordsFn a function to generate embedding coordinates (default none)

Value

map object (without the grid, if coordsFn was not specified)

Examples

```
d <- iris[,1:4]
EmbedSOM::PlotEmbed(
  EmbedSOM::EmbedSOM(
    data = d,
    map = EmbedSOM::RandomMap(d, 30, EmbedSOM::GraphCoords()),
    pch=19, clust=iris[,5]
  )
)
```

SOM

*Build a self-organizing map***Description**

Build a self-organizing map

Usage

```
SOM(data, xdim = 10, ydim = 10, zdim = NULL, batch = F,
     rlen = 10, alphaA = c(0.05, 0.01),
     radiusA = stats::quantile(nhbrdist, 0.67) * c(1, 0), alphaB = alphaA
     * c(-negAlpha, -0.1 * negAlpha), radiusB = negRadius * radiusA,
     negRadius = 1.33, negAlpha = 0.1, epochRadii = seq(radiusA[1],
     radiusA[2], length.out = rlen), init = FALSE, initf = Initialize_PCA,
     distf = 2, codes = NULL, importance = NULL, coordsFn = NULL,
     nhbr.method = "maximum", noMapping = F, parallel = F,
     threads = if (parallel) 0 else 1)
```

Arguments

data	Matrix containing the training data
xdim	Width of the grid
ydim	Hight of the grid
zdim	Depth of the grid, causes the grid to be 3D if set
batch	Use batch training (default FALSE chooses online training, which is more like FlowSOM)
rlen	Number of training epochs; or number of times to loop over the training data in online training
alphaA	Start and end learning rate for online learning (only for online training)
radiusA	Start and end radius
alphaB	Start and end learning rate for the second radius (only for online training)
radiusB	Start and end radius (only for online training; make sure it is larger than radiusA)
negRadius	easy way to set radiusB as a multiple of default radius (use lower value for higher dimensions)
negAlpha	the same for alphaB
epochRadii	Vector of length rlen with precise epoch radii (only for batch training)
init	Initialize cluster centers in a non-random way
initf	Use the given initialization function if init==T (default: Initialize_PCA)
distf	Distance function (1=manhattan, 2=euclidean, 3=chebyshev, 4=cosine)
codes	Cluster centers to start with

importance	array with numeric values. Columns of data will be scaled according to importance.
coordsFn	Function to generate/transform grid coordinates (e.g. <code>tSNECoords()</code>). If NULL (default), the grid is the canonical SOM grid.
nhbr.method	Way of computing grid distances, passed as <code>method=</code> to <code>stats::dist()</code> function. Defaults to <code>maximum</code> (square neighborhoods); use <code>euclidean</code> for round neighborhoods.
noMapping	If TRUE, do not compute the mapping (default FALSE). Makes the process quicker by <code>1 rlen</code> .
parallel	Parallelize the batch training by setting appropriate threads. Defaults to FALSE. Always use <code>batch=TRUE</code> for fully parallelized version, online training is not parallelizable. Passed to <code>MapDataToCodes()</code> .
threads	Number of threads of the batch training (has no effect on online training). Defaults to 0 (chooses maximum available hardware threads) if <code>parallel==TRUE</code> or 1 (single thread) if <code>parallel==FALSE</code> . Passed to <code>MapDataToCodes()</code> .

Value

A map useful for embedding (`EmbedSOM()` function) or further analysis, e.g. clustering.

See Also

`FlowSOM::SOM`

tSNECoords	<i>Add tSNE-based coordinates to a map</i>
------------	--

Description

Add tSNE-based coordinates to a map

Usage

```
tSNECoords(dim = NULL, tSNEFn = Rtsne::Rtsne, ...)
```

Arguments

dim	Dimension of the result (passed to tSNEFn as <code>dims</code>)
tSNEFn	tSNE function to run (default <code>Rtsne::Rtsne</code>)
...	passed to tSNEFn

Value

a function that transforms the map, usable as `coordsFn` parameter

UMAPCoords *Add UMAP-based coordinates to a map*

Description

Add UMAP-based coordinates to a map

Usage

```
UMAPCoords(dim = NULL, UMAPFn = NULL)
```

Arguments

dim Dimension of the result (passed to UMAPFn as n_components)
 UMAPFn UMAP function to run (default `umap::umap` configured by `umap::umap.defaults`)

Value

a function that transforms the map, usable as `coordsFn` parameter

UMatrixCoords *Add U-Matrix-optimized embedding coordinates to the map*

Description

The map must already contain a SOM grid with corresponding `xdim,ydim` (possibly `zdim`)

Usage

```
UMatrixCoords(dim = NULL, dist.method = NULL, distFn = function(x) x,  

  layoutFn = igraph::layout_with_kk)
```

Arguments

dim Dimension of the result (passed to layoutFn)
 dist.method The method to compute distances, passed to `stats::dist()` as parameter method
 distFn Custom transformation function of the distance matrix
 layoutFn iGraph-compatible graph layouting function (default `igraph::layout_with_kk`)

Value

a function that transforms the map, usable as `'coordsFn'` parameter

`uwotCoords`*Add UMAP-based coordinates to a map, using the 'uwot' package*

Description

Add UMAP-based coordinates to a map, using the 'uwot' package

Usage

```
uwotCoords(dim = NULL, uwotFn = uwot::umap, ...)
```

Arguments

<code>dim</code>	Dimension of the result (passed to uwotFn as <code>dims</code>)
<code>uwotFn</code>	UMAP function to run (default uwot::umap)
<code>...</code>	passed to uwotFn

Value

a function that transforms the map, usable as `coordsFn` parameter

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