

Package ‘dyndimred’

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Type Package

Title Dimensionality Reduction Methods in a Common Format

Version 1.0.4

Description Provides a common interface for applying dimensionality reduction methods, such as Principal Component Analysis ('PCA'), Independent Component Analysis ('ICA'), diffusion maps, Locally-Linear Embedding ('LLE'), t-distributed Stochastic Neighbor Embedding ('t-SNE'), and Uniform Manifold Approximation and Projection ('UMAP'). Has built-in support for sparse matrices.

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Encoding UTF-8

LazyData true

Imports dynutils (>= 1.0.5), irlba, lmds, stats, tibble

Suggests testthat, diffusionMap, fastICA, igraph, lle, MASS, Matrix, RANN, Rtsne, smacof, uwot

RoxygenNote 7.1.1

URL <https://github.com/dynverse/dyndimred>

BugReports <https://github.com/dynverse/dyndimred/issues>

NeedsCompilation no

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dimred	<i>Perform simple dimensionality reduction</i>
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Description

Perform simple dimensionality reduction

Usage

```
dimred(x, method, ndim, ...)

dimred_dm_destiny(
  x,
  ndim = 2,
  distance_method = c("euclidean", "spearman", "cosine")
)

dimred_dm_diffusionmap(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_ica(x, ndim = 3)

dimred_knn_fr(
  x,
  ndim = 2,
  lmds_components = 10,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski"),
  n_neighbors = 10
)

dimred_landmark_mds(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
```

```

    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_lle(x, ndim = 3)

dimred_mds(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_mds_isomds(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_mds_sammon(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_mds_smacof(
  x,
  ndim = 2,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)

dimred_pca(x, ndim = 2)

list_dimred_methods()

```

Arguments

<code>x</code>	Log transformed expression data, with rows as cells and columns as features
<code>method</code>	The name of the dimensionality reduction method to use
<code>ndim</code>	The number of dimensions
<code>...</code>	Any arguments to be passed to the dimensionality reduction method
<code>distance_method</code>	The name of the distance metric, see dynutils::calculate_distance
<code>lmds_components</code>	The number of lmds components to use. If NULL, LMDS will not be performed first. If this is a matrix, it is assumed it is a dimred for x.

`n_neighbors` The size of local neighborhood (in terms of number of neighboring sample points).

Examples

```
library(Matrix)
x <- abs(Matrix::rsparsematrix(100, 100, .5))
dimred(x, "pca", ndim = 3)
dimred(x, "ica", ndim = 3)

if (interactive()) {
  dimred_dm_destiny(x)
  dimred_dm_diffusionmap(x)
  dimred_ica(x)
  dimred_landmark_mds(x)
  dimred_lle(x)
  dimred_mds(x)
  dimred_mds_isomds(x)
  dimred_mds_sammon(x)
  dimred_mds_smacof(x)
  dimred_pca(x)
  dimred_tsne(x)
  dimred_umap(x)
}
```

dimred_tsne	<i>tSNE</i>
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Description

tSNE

Usage

```
dimred_tsne(
  x,
  ndim = 2,
  perplexity = 30,
  theta = 0.5,
  initial_dims = 50,
  distance_method = c("pearson", "spearman", "cosine", "euclidean", "chisquared",
    "hamming", "kullback", "manhattan", "maximum", "canberra", "minkowski")
)
```

Arguments

`x` Log transformed expression data, with rows as cells and columns as features

`ndim` The number of dimensions

perplexity	numeric; Perplexity parameter (should not be bigger than $3 * \text{perplexity} < \text{nrow}(X) - 1$, see details for interpretation)
theta	numeric; Speed/accuracy trade-off (increase for less accuracy), set to 0.0 for exact TSNE (default: 0.5)
initial_dims	integer; the number of dimensions that should be retained in the initial PCA step (default: 50)
distance_method	The name of the distance metric, see dynutils::calculate_distance

See Also

[Rtsne::Rtsne\(\)](#)

Examples

```
library(Matrix)
dataset <- abs(Matrix::rsparsematrix(100, 100, .5))
dimred_tsne(dataset, ndim = 3)
```

dimred_umap

UMAP

Description

UMAP

Usage

```
dimred_umap(
  x,
  ndim = 2,
  distance_method = c("euclidean", "cosine", "manhattan"),
  pca_components = 50,
  n_neighbors = 15L,
  init = "spectral",
  n_threads = 1
)
```

Arguments

x	Log transformed expression data, with rows as cells and columns as features
ndim	The number of dimensions
distance_method	The name of the distance metric, see dynutils::calculate_distance
pca_components	The number of pca components to use for UMAP. If NULL, PCA will not be performed first

<code>n_neighbors</code>	The size of local neighborhood (in terms of number of neighboring sample points).
<code>init</code>	Type of initialization for the coordinates. Options are: • "spectral" Spectral embedding using the normalized Laplacian of the fuzzy 1-skeleton, with Gaussian noise added. • "normlaplacian". Spectral embedding using the normalized Laplacian of the fuzzy 1-skeleton, without noise. • "random". Coordinates assigned using a uniform random distribution between -10 and 10. • "lvrandom". Coordinates assigned using a Gaussian distribution with standard deviation 1e-4, as used in LargeVis (Tang et al., 2016) and t-SNE. • "laplacian". Spectral embedding using the Laplacian Eigenmap (Belkin and Niyogi, 2002). • "pca". The first two principal components from PCA of X if X is a data frame, and from a 2-dimensional classical MDS if X is of class "dist". • "spca". Like "pca", but each dimension is then scaled so the standard deviation is 1e-4, to give a distribution similar to that used in t-SNE. This is an alias for <code>init = "pca"</code>, <code>init_sdev = 1e-4</code>. • "agspectral" An "approximate global" modification of "spectral" which all edges in the graph to a value of 1, and then sets a random number of edges (<code>negative_sample_rate</code> edges per vertex) to 0.1, to approximate the effect of non-local affinities. • A matrix of initial coordinates. For spectral initializations, ("spectral", "normlaplacian", "laplacian"), if more than one connected component is identified, each connected component is initialized separately and the results are merged. If <code>verbose = TRUE</code> the number of connected components are logged to the console. The existence of multiple connected components implies that a global view of the data cannot be attained with this initialization. Either a PCA-based initialization or increasing the value of <code>n_neighbors</code> may be more appropriate.
<code>n_threads</code>	Number of threads to use (except during stochastic gradient descent). Default is half the number of concurrent threads supported by the system. For nearest neighbor search, only applies if <code>nn_method = "annoy"</code> . If <code>n_threads > 1</code> , then the Annoy index will be temporarily written to disk in the location determined by tempfile .

See Also

[uwot::umap\(\)](#)

Examples

```
library(Matrix)
dataset <- abs(Matrix::rsparsematrix(100, 100, .5))
dimred_umap(dataset, ndim = 2, pca_components = NULL)
```

dyndimred*Common dimensionality reduction methods*

Description

Provides a common interface for applying common dimensionality reduction methods, Such as PCA, ICA, diffusion maps, LLE, t-SNE, and umap.

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