

# Package: mp (via r-universe)

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**Type** Package

**Title** Multidimensional Projection Techniques

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**Author** Francisco M. Fatore, Samuel G. Fadel

**Maintainer** Francisco M. Fatore <fmfatore@gmail.com>

**Description** Multidimensional projection techniques are used to create two dimensional representations of multidimensional data sets.

**License** GPL

**Depends** R (>= 1.8.0)

**Imports** Rcpp (>= 0.11.0)

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 5.0.1

**NeedsCompilation** yes

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`forceScheme`*Force Scheme Projection*

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### Description

Creates a 2D representation of the data based on a dissimilarity matrix. A few modifications have been made in relation to the method described in the literature: shuffled indices are used to minimize the order dependency factor, only a fraction of delta is used for better stability and a tolerance factor was introduced as a second stop criterion.

### Usage

```
forceScheme(D, Y = NULL, max.iter = 50, tol = 0, fraction = 8,  
            eps = 1e-05)
```

### Arguments

<code>D</code>	A dissimilarity structure such as that returned by <code>dist</code> or a full symmetric matrix containing the dissimilarities.
<code>Y</code>	Initial 2D configuration. A random configuration will be used when omitted.
<code>max.iter</code>	Maximum number of iterations that the algorithm will run.
<code>tol</code>	The tolerance for the accumulated error between iterations. If set to 0, the algorithm will run <code>max.iter</code> times.
<code>fraction</code>	Controls the point movement. Larger values means less freedom to move.
<code>eps</code>	Minimum distance between two points.

### Value

The 2D representation of the data.

### References

Eduardo Tejada, Rosane Minghim, Luis Gustavo Nonato: On improved projection techniques to support visual exploration of multi-dimensional data sets. *Information Visualization* 2(4): 218-231 (2003)

### See Also

[dist](#) (stats) and [dist](#) (proxy) for d computation

**Examples**

```
# Eurodist example
emb <- forceScheme(eurodist)
plot(emb, type = "n", xlab = "", ylab = "", asp=1, axes=FALSE, main="")
text(emb, labels(eurodist), cex = 0.6)

# Iris example
emb <- forceScheme(dist(iris[,1:4]))
plot(emb, col=iris$Species)
```

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is.symmetric	<i>Tests whether the given matrix is symmetric.</i>
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**Description**

Tests whether the given matrix is symmetric.

**Usage**

```
is.symmetric(mat)
```

**Arguments**

mat                    Matrix to be tested for symmetry.

**Value**

Whether the matrix is symmetric.

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lamp	<i>Local Affine Multidimensional Projection</i>
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**Description**

Creates a 2D representation of the data. Requires a subsample (sample.indices) and its 2D representation (Ys).

**Usage**

```
lamp(X, sample.indices = NULL, Ys = NULL, cp = 1)
```

**Arguments**

X	A data frame or matrix.
sample.indices	The indices of data points in X used as subsamples. If not given, some points from X will be randomly selected and Ys will be generated by calling forceScheme on them.
Ys	Initial 2D configuration of the data subsamples (will be ignored if sample.indices is NULL). Scaling the columns to [-0.5, 0.5] is recommended to avoid scaling problems.
cp	Proportion of nearest control points to be used.

**Value**

The 2D representation of the data.

**References**

Joia, P.; Paulovich, F.V.; Coimbra, D.; Cuminato, J.A.; Nonato, L.G., "Local Affine Multidimensional Projection," Visualization and Computer Graphics, IEEE Transactions on , vol.17, no.12, pp.2563,2571, Dec. 2011

**Examples**

```
# Iris example
emb <- lamp(iris[, 1:4])
plot(emb, col=iris$Species)
```

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lsp

*Least-Square Projection*


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**Description**

Creates a q-dimensional representation of multidimensional data. Requires a subsample (sample.indices) and its qD representation (Ys).

**Usage**

```
lsp(X, sample.indices = NULL, Ys = NULL, k = 15, q = 2)
```

**Arguments**

X	A data frame or matrix.
sample.indices	The indices of data points in X used as subsamples. If not given, some rows from X will be randomly selected and Ys will be generated by calling forceScheme on them.

Ys	Initial kD configuration of the data subsamples (will be ignored if sample.indices is NULL).
k	Number of neighbors used to build the neighborhood graph.
q	The target dimensionality.

**Value**

The qD representation of the data.

**References**

F. V. Paulovich, L. Nonato, R. Minghim, and H. Levkowitz, Least-Square Projection: A fast high-precision multidimensional projection technique and its application to document mapping, vol. 14, no. 3, pp. 564-575.

**Examples**

```
# Iris example
emb <- lsp(iris[, 1:4])
plot(emb, col=iris$Species)
```

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mp *Multidimensional Projection Techniques*

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**Description**

Implementation of multidimensional projection techniques

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pekalska *Pekalska's approach to speeding up Sammon's mapping.*

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**Description**

Creates a k-dimensional representation of the data. As input, a subsample and its k-dimensional mapping are required. The method approximates the subsample mapping to a linear mapping based on the distances matrix of the subsample and then applies the same mapping to all instances.

**Usage**

```
pekalska(D, sample.indices = NULL, Ys = NULL)
```

**Arguments**

D	dist object or distances matrix.
sample.indices	The indices of subsamples.
Ys	The subsample mapping (k-dimensional).

**Value**

The low-dimensional representation of the data.

**References**

Pekalska, E., de Ridder, D., Duin, R. P., & Kraaijveld, M. A. (1999). A new method of generalizing Sammon mapping with application to algorithm speed-up (pp. 221-228).

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plmp

*Part-Linear Multidimensional Projection*

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**Description**

Creates a k-dimensional representation of the data. As input, a subsample and its k-dimensional mapping (control points) are required. The method approximates the subsample mapping to a linear mapping and then applies the same mapping to all instances.

**Usage**

```
plmp(X, sample.indices = NULL, Ys = NULL, k = 2)
```

**Arguments**

X	A dataframe or matrix representing the data.
sample.indices	The indices of subsamples used as control points.
Ys	The control points.
k	The target dimensionality.

**Value**

The low-dimensional representation of the data.

**References**

Paulovich, F.V.; Silva, C.T.; Nonato, L.G., "Two-Phase Mapping for Projecting Massive Data Sets," Visualization and Computer Graphics, IEEE Transactions on , vol.16, no.6, pp.1281,1290, Nov.-Dec. 2010.

**Examples**

```
# Iris example
emb <- plmp(iris[,1:4])
plot(emb, col=iris$Species)
```

tSNE

*t-Distributed Stochastic Neighbor Embedding***Description**

Creates a k-dimensional representation of the data by modeling the probability of picking neighbors using a Gaussian for the high-dimensional data and t-Student for the low-dimensional map and then minimizing the KL divergence between them. This implementation uses the same default parameters as defined by the authors.

**Usage**

```
tSNE(X, Y = NULL, k = 2, perplexity = 30, n.iter = 1000, eta = 500,
      initial.momentum = 0.5, final.momentum = 0.8, early.exaggeration = 4,
      gain.fraction = 0.2, momentum.threshold.iter = 20,
      exaggeration.threshold.iter = 100, max.binsearch.tries = 50)
```

**Arguments**

X	A data frame, data matrix, dissimilarity (distance) matrix or dist object.
Y	Initial k-dimensional configuration. If NULL, the method uses a random initial configuration.
k	Target dimensionality. Avoid anything other than 2 or 3.
perplexity	A rough upper bound on the neighborhood size.
n.iter	Number of iterations to perform.
eta	The "learning rate" for the cost function minimization
initial.momentum	The initial momentum used before changing
final.momentum	The momentum to use on remaining iterations
early.exaggeration	The early exaggeration applied to initial iterations
gain.fraction	Undocumented
momentum.threshold.iter	Number of iterations before using the final momentum
exaggeration.threshold.iter	Number of iterations before using the real probabilities
max.binsearch.tries	Maximum number of tries in binary search for parameters to achieve the target perplexity

**Value**

The k-dimensional representation of the data.

**References**

L.J.P. van der Maaten and G.E. Hinton. *Visualizing High-Dimensional Data Using t-SNE.* Journal of Machine Learning Research 9(Nov): 2579-2605, 2008.

**Examples**

```
# Iris example
emb <- tSNE(iris[, 1:4])
plot(emb, col=iris$Species)
```



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