

# Package ‘superMDS’

February 20, 2015

**Type** Package

**Title** Implements the supervised multidimensional scaling (superMDS) proposal of Witten and Tibshirani (2011)

**Version** 1.0.2

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**Author** Daniela M. Witten

**Maintainer** Daniela Witten <dwitten@u.washington.edu>

**Description** Witten and Tibshirani (2011) Supervised multidimensional scaling for visualization, classification, and bipartite ranking. Computational Statistics and Data Analysis 55(1): 789-801.

**License** GPL-2

**LazyLoad** yes

**NeedsCompilation** no

**Repository** CRAN

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superMDS-package	<i>Supervised multidimensional scaling for visualization, classification, and bipartite ranking</i>
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## Description

A method for implementing the supervised multidimensional scaling proposal of Witten and Tibshirani (2011)

**Details**

```

Package:    superMDS
Type:      Package
Version:   1.0.2
Date:      2013-01-02
License:   GPL-2
LazyLoad:  yes

```

Supervised multidimensional scaling (MDS) is a supervised version of least squares MDS. Suppose that we have a  $n \times n$  dissimilarity matrix  $D$  and we want to find a set of  $n$  configuration points  $z_1, \dots, z_n$ , each a vector of length  $s$ , so that  $D$  is well-approximated by the Euclidean distances between the configuration points. Then least squares MDS can be used. However, suppose that we also have a vector of binary class labels associated with the dissimilarity matrix,  $y_i = 1$  or  $2$  for  $i=1, \dots, n$ . Then we might want configuration points whose Euclidean distances approximate  $D$ , and also that have the property that  $z_i \cdot z_j > z_i \cdot z_k$  when  $y_i > y_j$ . This is the objective of supervised MDS. It leads to a method for visualizing observations, as well as a classification method. Details can be found in the paper below.

#### Author(s)

Daniela M. Witten

Maintainer: Daniela Witten <dwitten@u.washington.edu>

#### References

Witten and Tibshirani (2011) Supervised multidimensional scaling for visualization, classification, and bipartite ranking. *Computational Statistics and Data Analysis* 55(1): 789-801.

#### Examples

```

##### Generate some data #####
n <- 30
p <- 10
x <- matrix(rnorm(n*p),ncol=p)
y <- c(rep(1,n/2),rep(2,n/2))
xte <- matrix(rnorm(n*p),ncol=p)
yte <- c(rep(1,n/2),rep(2,n/2))
x[y==1,1:(p)] <- x[y==1,1:(p)] + .4
x[y==2,1:(p)] <- x[y==2,1:(p)] - .4
xte[yte==1,1:(p)] <- xte[yte==1,1:(p)] + .4
xte[yte==2,1:(p)] <- xte[yte==2,1:(p)] - .4
# Done generating data #

##### Perform SuperMDS #####
out <- TrainSuperMDS(x=x,y=y,alpha=.4,S=2, silent=TRUE)
# A plot of the training configuration points #
par(mfrow=c(1,2))
plot(out$x, col=yte, main="Training Data", xlab="Dimension 1", ylab="Dimension 2")
testout <- TestSuperMDS(trout=out,xte=xte)
ytehat <- testout$ytehat

```

```
# A table showing the true vs predicted class labels #
print(table(ytehat,yte))
# A plot of the test configuration points #
plot(testout$ztc, col=yte, main="Test Data", xlab="Dimension 1", ylab="Dimension 2")
```

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TestSuperMDS	<i>Given the configuration points of a set of training observations, and the dissimilarities between the training and test observations, compute the configuration points for a set of test observations.</i>
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### Description

Suppose that we have training data observations  $x$  (of dimension  $n \times p$ ) with an associated binary outcome vector  $y$  of length  $n$ , and that TrainSuperMDS has already been run on the training observations. Furthermore, we have test observations  $x_{te}$  (of dimension  $m \times p$ ) for which we do not have an outcome vector. This function will predict the class of the test observations, and also to compute configuration points for the test observations.

### Usage

```
TestSuperMDS(trout, xte = NULL, dtetr = NULL)
```

### Arguments

trout	The output of a call to TrainSuperMDS on the training data. We assume that there were $n$ training observations which were either passed into TrainSuperMDS as a $n \times n$ dissimilarity matrix, or as a $n \times p$ data matrix.
xte	The test observations, a matrix of dimension $m \times p$ . If this is NULL then must pass in dtetr. Can pass in xte only if previously passed in $x$ when TrainSuperMDS was called. Otherwise, pass in dtetr instead.
dtetr	A $m \times n$ data matrix with the dissimilarity between each test observation and each training observation; if NULL then must pass in xte.

### Value

ytehat	Predicted class labels for test data.
ztc	Predicted configuration points for test data; should be a matrix of dimension $m \times S$ where $S$ is the dimension of training configuration points.

### Author(s)

Daniela M Witten

### References

Witten and Tibshirani (2011) Supervised multidimensional scaling for visualization, classification, and bipartite ranking. CSDA.

**See Also**[TrainSuperMDS](#)**Examples**

```
# Try ?superMDS for examples.
```

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TrainSuperMDS	<i>Find a set of configuration points that agree with a dissimilarity matrix D and a vector of class labels y</i>
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**Description**

Given a  $n \times n$  dissimilarity matrix  $D$  and a  $n$ -vector of binary (1,2) class labels  $y$ , this function outputs a set of configuration points  $z_1, \dots, z_n$ , each a  $S$ -vector, such that the distances between the configuration points approximate the dissimilarity matrix  $D$ , AND such that  $z_i \geq z_j$  tends to occur when  $y_i \geq y_j$ .

**Usage**

```
TrainSuperMDS(d = NULL, y, alpha = 0.5, S = 2, x = NULL, nstarts = 5, silent = FALSE)
```

**Arguments**

<code>d</code>	A $n \times n$ dissimilarity matrix. If NULL, then <code>x</code> , a $n \times p$ data matrix, must be input instead.
<code>y</code>	A $n$ -vector of binary labels, in the form of 1's and 2's. For instance, <code>c(1,1,1,2,2)</code> could be input if $D$ is a $5 \times 5$ matrix.
<code>alpha</code>	A scalar between 0 and 1. If <code>alpha=0</code> then this is just least squares MDS, and if <code>alpha=1</code> then it's completely supervised.
<code>S</code>	The number of dimensions of the configuration points $z_1, \dots, z_n$ . Must be at least equal to 1.
<code>x</code>	A $n \times p$ data matrix, to be input only if $D$ is NULL.
<code>nstarts</code>	The supervised MDS algorithm finds a local minimum for the objective. Here, specify the number of initial values to try. If <code>nstarts&gt;1</code> then the set of configuration points corresponding to the optimal (smallest) value of the objective will be reported.
<code>silent</code>	Set to TRUE in order to turn off printing output to screen.

**Value**

<code>z</code>	A $n \times S$ matrix of the configuration points obtained.
<code>crits</code>	The values of the criterion obtained at the iterations of the algorithm.
<code>stress</code>	The portion of the final criterion value that are due to the STRESS component of the objective function.
<code>super</code>	The portion of the final criterion value that are due to the SUPERVISED component of the objective function.

**Author(s)**

Daniela M Witten

**References**

Witten and Tibshirani (2011) Supervised multidimensional scaling for visualization, classification, and bipartite ranking. Computational Statistics and Data Analysis.

**See Also**

[TestSuperMDS](#)

**Examples**

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# Try ?superMDS for examples
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