

# Package ‘regda’

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**Type** Package

**Title** Regularised Discriminant Analysis

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## Description

Regularised discriminant analysis functions. The classical regularised discriminant analysis proposed by Friedman in 1989, including cross-validation, of which the linear and quadratic discriminant analyses are special cases. Further, the regularised maximum likelihood linear discriminant analysis, including cross-validation. References: Friedman J.H. (1989): ``Regularized Discriminant Analysis''. Journal of the American Statistical Association 84(405): 165--175. <doi:10.2307/2289860>. Friedman J., Hastie T. and Tibshirani R. (2009). ``The elements of statistical learning'', 2nd edition. Springer, Berlin. <doi:10.1007/978-0-387-84858-7>. Tsagris M., Preston S. and Wood A.T.A. (2016). ``Improved classification for compositional data using the alpha-transformation''. Journal of Classification, 33(2): 243--261. <doi:10.1007/s00357-016-9207-5>.

**License** GPL (>= 2)

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regda-package

*Regularised Discriminant Analysis*

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## Description

Description: Regularised discriminant analysis functions. The classical regularised discriminant analysis proposed by Friedman in 1989, including cross-validation, of which the linear and quadratic discriminant analyses are special cases. Further, the regularised maximum likelihood linear discriminant analysis, including cross-validation.

## Details

Package: regda  
Type: Package  
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License: GPL-2

## Maintainers

Michail Tsagris <mtsagris@uoc.gr>.

## Author(s)

Michail Tsagris <mtsagris@uoc.gr>

## References

- Friedman J.H. (1989): Regularized Discriminant Analysis. *Journal of the American Statistical Association* 84(405): 165–175.
- Friedman Jerome, Trevor Hastie and Robert Tibshirani (2009). *The elements of statistical learning*, 2nd edition. Springer, Berlin.
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Cross-validation for the regularised maximum likelihood linear discriminant analysis

*Cross-validation for the regularised maximum likelihood linear discriminant analysis*

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## Description

Cross-validation for the regularised maximum likelihood linear discriminant analysis.

## Usage

```
regmlelda.cv(x, ina, lambda = seq(0, 1, by = 0.1), folds = NULL, nfolds = 10,  
             stratified = TRUE, seed = FALSE, pred.ret = FALSE)
```

## Arguments

x	A matrix with numerical data.
ina	A numerical vector or factor with consecutive numbers indicating the group to which each observation belongs to.
lambda	A vector of regularization values $\lambda$ such as (0, 0.1, 0.2,...).
folds	A list with the indices of the folds.
nfolds	The number of folds to be used. This is taken into consideration only if "folds" is NULL.
stratified	Do you want the folds to be selected using stratified random sampling? This preserves the analogy of the samples of each group. Make this TRUE if you wish.
seed	If you set this to TRUE, the same folds will be created every time.
pred.ret	If you want the predicted values returned set this to TRUE.

## Details

Cross-validation for the regularised maximum likelihood linear discriminant analysis is performed. The function is not extremely fast, yet is pretty fast.

## Value

A list including:

preds	If pred.ret is TRUE the predicted values for each fold are returned as elements in a list.
crit	A vector whose length is equal to the number of k and is the accuracy metric for each k. For the classification case it is the percentage of correct classification. For the regression case the mean square of prediction error.

**Author(s)**

Michail Tsagris.

R implementation and documentation: Michail Tsagris <mtsagris@uoc.gr>.

**References**

Friedman J., Hastie T. and Tibshirani R. (2017). The elements of statistical learning. New York: Springer.

**See Also**

[reg.mle.lda](#)

**Examples**

```
x <- as.matrix(iris[, 1:4])
mod <- regmlelda.cv(x, iris[, 5])
```

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Regularised discriminant analysis for Euclidean data

*Regularised discriminant analysis for Euclidean data*

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**Description**

Regularised discriminant analysis for Euclidean data.

**Usage**

```
rda(xnew, x, ina, gam = 1, del = 0)
```

**Arguments**

xnew	A matrix with the new data whose group is to be predicted. They have to be continuous.
x	A matrix with the available data. They have to be continuous.
ina	A group indicator variable for the available data.
gam	This is a number between 0 and 1. It is the weight of the pooled covariance and the diagonal matrix.
del	This is a number between 0 and 1. It is the weight of the LDA and QDA.

**Details**

The covariance matrix of each group is calculated and then the pooled covariance matrix. The spherical covariance matrix consists of the average of the pooled variances in its diagonal and zeros in the off-diagonal elements.  $\text{gam}$  is the weight of the pooled covariance matrix and  $1-\text{gam}$  is the weight of the spherical covariance matrix,  $S_a = \text{gam} * S_p + (1-\text{gam}) * s_p$ . Then it is a compromise between LDA and QDA.  $\text{del}$  is the weight of  $S_a$  and  $1-\text{del}$  the weight of each group covariance group.

**Value**

A list including:

prob	The estimated probabilities of the new data of belonging to each group.
scores	The estimated scores of the new data of each group.
est	The estimated group membership of the new data.

**Author(s)**

Michail Tsagris.

R implementation and documentation: Michail Tsagris <mtsagris@uoc.gr>.

**References**

Friedman J.H. (1989): Regularized Discriminant Analysis. *Journal of the American Statistical Association* 84(405): 165–175.

Friedman Jerome, Trevor Hastie and Robert Tibshirani (2009). *The elements of statistical learning*, 2nd edition. Springer, Berlin.

Tsagris M., Preston S. and Wood A.T.A. (2016). Improved classification for compositional data using the  $\alpha$ -transformation. *Journal of Classification*, 33(2): 243–261.

**See Also**

[rda.tune](#)

**Examples**

```
x <- as.matrix(iris[, 1:4])
ina <- iris[, 5]
mod <- rda(x, x, ina)
table(ina, mod$est)
```

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Regularised maximum likelihood linear discriminant analysis

*Regularised maximum likelihood linear discriminant analysis*

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**Description**

Regularised maximum likelihood linear discriminant analysis.

**Usage**

```
reg.mle.lda(xnew, x, ina, lambda)
```

**Arguments**

xnew	A numerical vector or a matrix with the new observations, continuous data.
x	A matrix with numerical data.
ina	A numerical vector or factor with consecutive numbers indicating the group to which each observation belongs to.
lambda	A vector of regularization values $\lambda$ such as (0, 0.1, 0.2,...).

**Details**

Regularised maximum likelihood linear discriminant analysis is performed. The function is not extremely fast, yet is pretty fast.

**Value**

A matrix with the predicted group of each observation in "xnew". Every column corresponds to a  $\lambda$  value. If you have just one value of  $\lambda$ , then you will have one column only.

**Author(s)**

Michail Tsagris.

R implementation and documentation: Michail Tsagris <mtsagris@uoc.gr>.

**See Also**

[regmlelda.cv](#)

**Examples**

```
x <- as.matrix(iris[, 1:4])
ina <- iris[, 5]
a <- reg.mle.lda(x, x, ina, lambda = seq(0, 1, by = 0.1) )
```

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Tuning the parameters of the regularised discriminant analysis

*Tuning the parameters of the regularised discriminant analysis*

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**Description**

Tuning the parameters of the regularised discriminant analysis for Euclidean data.

**Usage**

```
rda.tune(x, ina, nfolds = 10, gam = seq(0, 1, by = 0.1), del = seq(0, 1, by = 0.1),
ncores = 1, folds = NULL, stratified = TRUE, seed = NULL)
```

**Arguments**

<code>x</code>	A matrix with the data.
<code>ina</code>	A group indicator variable for the available data.
<code>nfolds</code>	The number of folds in the cross validation.
<code>gam</code>	A grid of values for the $\gamma$ parameter as defined in Tsagris et al. (2016).
<code>del</code>	A grid of values for the $\delta$ parameter as defined in Tsagris et al. (2016).
<code>ncores</code>	The number of cores to use. If more than 1, parallel computing will take place. It is advisable to use it if you have many observations and or many variables, otherwise it will slow down the process.
<code>folds</code>	If you have the list with the folds supply it here. You can also leave it NULL and it will create folds.
<code>stratified</code>	Do you want the folds to be created in a stratified way? TRUE or FALSE.
<code>seed</code>	You can specify your own seed number here or leave it NULL.

**Details**

Cross validation is performed to select the optimal parameters for the regularised discriminant analysis and also estimate the rate of accuracy.

The covariance matrix of each group is calculated and then the pooled covariance matrix. The spherical covariance matrix consists of the average of the pooled variances in its diagonal and zeros in the off-diagonal elements. `gam` is the weight of the pooled covariance matrix and `1-gam` is the weight of the spherical covariance matrix,  $S_a = \text{gam} * S_p + (1-\text{gam}) * s_p$ . Then it is a compromise between LDA and QDA. `del` is the weight of  $S_a$  and `1-del` the weight of each group covariance group.

**Value**

A list including: If `graph` is TRUE a plot of a heatmap of the performance `s` will appear.

<code>per</code>	An array with the estimate rate of correct classification for every fold. For each of the <code>M</code> matrices, the row values correspond to <code>gam</code> and the columns to the <code>del</code> parameter.
<code>percent</code>	A matrix with the mean estimated rates of correct classification. The row values correspond to <code>gam</code> and the columns to the <code>del</code> parameter.
<code>se</code>	A matrix with the standard error of the mean estimated rates of correct classification. The row values correspond to <code>gam</code> and the columns to the <code>del</code> parameter.
<code>result</code>	The estimated rate of correct classification along with the best <code>gam</code> and <code>del</code> parameters.
<code>runtime</code>	The time required by the cross-validation procedure.

**Author(s)**

Michail Tsagris.

R implementation and documentation: Michail Tsagris <mtsagris@uoc.gr>.

**References**

Friedman J.H. (1989): Regularized Discriminant Analysis. *Journal of the American Statistical Association* 84(405): 165–175.

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**See Also**

[rda](#)

**Examples**

```
mod <- rda.tune(as.matrix(iris[, 1:4]), iris[, 5], gam = seq(0, 1, by = 0.2),  
del = seq(0, 1, by = 0.2) )  
mod
```

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