

# Package ‘RMixtCompUtilities’

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**Type** Package

**Title** Utility Functions for 'MixtComp' Outputs

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**Description** Mixture Composer <<https://github.com/modal-inria/MixtComp>> is a project to build mixture models with heterogeneous data sets and partially missing data management. This package contains graphical, getter and some utility functions to facilitate the analysis of 'MixtComp' output.

**URL** <https://github.com/modal-inria/MixtComp>,  
<https://massiccc.lille.inria.fr/>

**BugReports** <https://github.com/modal-inria/MixtComp/issues>

**Imports** plotly, ggplot2, scales

**Suggests** testthat, xml2, RMixtCompIO (>= 4.0.4), Rmixmod, blockcluster

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**RMixtCompUtilities-package**  
*RMixtCompUtilities*

---

### Description

MixtComp (Mixture Composer, <https://github.com/modal-inria/MixtComp>) is a model-based clustering package for mixed data originating from the Modal team (Inria Lille).

It has been engineered around the idea of easy and quick integration of all new univariate models, under the conditional independence assumption. Five basic models (Gaussian, Multinomial, Poisson, Weibull, NegativeBinomial) are implemented, as well as two advanced models (Func\_CS and Rank\_ISR). MixtComp has the ability to natively manage missing data (completely or by interval). MixtComp is used as an R package, but its internals are coded in C++ using state of the art libraries for faster computation.

Online SaaS version (not up-to-date): <https://massiccc.lille.inria.fr/>

This package contains plots, getters and format functions to simplify the use of RMixtComp and RMixtCompIO packages. It is recommended to use RMixtComp (instead of RMixtCompIO) which is more user-friendly.

## Details

`createAlgo` gives you default values for required parameters.

`convertFunctionalToVector`, `createFunctional` and `refactorCategorical` functions help to transform data to the required format.

Getters are available to easily access some results: `getBIC`, `getICL`, `getCompletedData`, `getParam`, `getTik`, `getEmpiricTik`, `getPartition`, `getType`, `getModel`, `getVarNames`.

You can compute discriminative powers and similarities with functions: `computeDiscrimPowerClass`, `computeDiscrimPowerVar`, `computeSimilarityClass`, `computeSimilarityVar`.

Graphics functions are `plot.MixtComp`, `heatmapClass`, `heatmapTikSorted`, `heatmapVar`, `histMisclassif`, `plotConvergence`, `plotDataBoxplot`, `plotDataCI`, `plotDiscrimClass`, `plotDiscrimVar`, `plotProportion`.

## See Also

`RMixtComp` `RMixtCompIO` `Rmixmod`, `blockcluster` packages

---

availableModels

*Available models*

---

## Description

Get information about models implemented in MixtComp

## Usage

```
availableModels()
```

## Value

a data.frame containing models implemented in MixtComp

**model** model name

**data.type** data type

**format** Special format required for individuals

**missing.formats** accepted formats (separated by a ;) for missing values

**hyperparameter** Required hyperparameters in the paramStr elements of model object

**comments** comments about the model

**reference** link to article

**Author(s)**

Quentin Grimonprez

**See Also**

`mixtCompLearn`

**Examples**

```
availableModels()
```

`computeDiscrimPowerVar`

*Discriminative power*

**Description**

Compute the discriminative power of each variable or classe

**Usage**

```
computeDiscrimPowerVar(outMixtComp, class = NULL)
computeDiscrimPowerClass(outMixtComp)
```

**Arguments**

- |                          |  |
|--------------------------|--|
| <code>outMixtComp</code> | object of class <i>MixtCompLearn</i> or <i>MixtComp</i> obtained using <code>mixtCompLearn</code> or <code>mixtCompPredict</code> functions from <code>RMixtComp</code> package or <code>rmcMultiRun</code> from <code>RMixtCompIO</code> package. |
| <code>class</code>       | NULL or a number of classes. If NULL, return the discriminative power of variables globally otherwise return the discriminative power of variables in the given class  |

**Details**

The discriminative power of variable  $j$  is defined by  $1 - C(j)$

$$C(j) = - \sum_{k=1}^K \sum_{i=1}^n P(Z_i = k | x_{ij}) \log(P(Z_i = k | x_{ij})) / (n * \log(K))$$

A high value (close to one) means that the variable is highly discriminating. A low value (close to zero) means that the variable is poorly discriminating.

The discriminative power of variable  $j$  in class  $k$  is defined by  $1 - C(j)$

$$C(j) = - \sum_{i=1}^n P(Z_i! = k | x_{ij}) \log(P(Z_i! = k | x_{ij})) / (n * \log(2))$$

The discriminative power of class k is defined by  $1 - D(k)$

$$D(k) = - \sum_{i=1}^n P(Z_i = k|x_i) \log(P(Z_i = k|x_i))/(n * \exp(-1))$$

### Value

the discriminative power

### Author(s)

Matthieu Marbac

### See Also

[plotDiscrimClass](#) [plotDiscrimVar](#)

### Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))
```

```
model <- list(var1 = list(type = "Gaussian", paramStr = ""),
               var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

discVar <- computeDiscrimPowerVar(resLearn)
discVarInClass1 <- computeDiscrimPowerVar(resLearn, class = 1)
discClass <- computeDiscrimPowerClass(resLearn)

# graphic representation of discriminant variables
plotDiscrimVar(resLearn)
# graphic representation of discriminant classes
plotDiscrimClass(resLearn)
```

**computeSimilarityVar** *Similarity*

## Description

Compute the similarity between variables (or classes)

## Usage

```
computeSimilarityVar(outMixtComp)
computeSimilarityClass(outMixtComp)
```

## Arguments

**outMixtComp** object of class *MixtCompLearn* or *MixtComp* obtained using *mixtCompLearn* or *mixtCompPredict* functions from *RMixtComp* package or *rmcMultiRun* from *RMixtCompIO* package.

## Details

The similarities between variables j and h is defined by  $\Delta(j,h)$

$$\Delta(j,h)^2 = 1 - \sqrt{(1/n) * \sum_{i=1}^n \sum_{k=1}^K (P(Z_i = k|x_{ij}) - P(Z_i = k|x_{ih}))^2}$$

The similarities between classes k and g is defined by  $1 - \Sigma(k,g)$

$$\Sigma(k,g)^2 = (1/n) * \sum_{i=1}^n (P(Z_i = k|x_i) - P(Z_i = g|x_i))^2$$

## Value

a similarity matrix

## Author(s)

Quentin Grimonprez

## See Also

[heatmapVar](#) [heatmapClass](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

simVar <- computeSimilarityVar(resLearn)
simClass <- computeSimilarityClass(resLearn)
```

## convertFunctionalToVector

*Convert a mixtcomp string into a list of 2 vectors*

## Description

Convert a mixtcomp string into a list of 2 vectors

## Usage

```
convertFunctionalToVector(x)
```

## Arguments

x	a string containing a functional observation (cf example)
---	---

## Value

a list of 2 vectors: time and value

**Author(s)**

Quentin Grimonprez

**Examples**

```
convertFunctionalToVector("1:5,1.5:12,1.999:2.9")
```

**createAlgo**

*Create algo object*

**Description**

create an algo object required by `mixtCompLearn` and `mixtCompPredict` from `RMixtComp`.

**Usage**

```
createAlgo(
  nbBurnInIter = 50,
  nbIter = 50,
  nbGibbsBurnInIter = 50,
  nbGibbsIter = 50,
  nInitPerClass = 10,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.99,
  nStableCriterion = 20
)
```

**Arguments**

<code>nbBurnInIter</code>	Number of iterations of the burn-in part of the SEM algorithm.
<code>nbIter</code>	Number of iterations of the SEM algorithm.
<code>nbGibbsBurnInIter</code>	Number of iterations of the burn-in part of the Gibbs algorithm.
<code>nbGibbsIter</code>	Number of iterations of the Gibbs algorithm.
<code>nInitPerClass</code>	Number of individuals used to initialize each cluster (default = 10).
<code>nSemTry</code>	Number of try of the algorithm for avoiding an error.
<code>confidenceLevel</code>	confidence level for confidence bounds for parameter estimation
<code>ratioStableCriterion</code>	stability partition required to stop earlier the SEM
<code>nStableCriterion</code>	number of iterations of partition stability to stop earlier the SEM

**Value**

a list with the parameters values

**Author(s)**

Quentin Grimonprez

**Examples**

```
# default values  
algo <- createAlgo()  
  
# change some values  
algo <- createAlgo(nbIter = 200)
```

---

createFunctional

*Create a functional in MixtComp format*

---

**Description**

Create a functional in MixtComp format

**Usage**

```
createFunctional(time, value)
```

**Arguments**

time	vector containing the time of the functional
value	vector containing the value of the functional

**Value**

The functional data formatted to the mixtcomp standard

**Author(s)**

Quentin Grimonprez

**Examples**

```
mat <- matrix(c(1, 2, 3, 9, 1, 1.5, 15, 1000), ncol = 2)  
createFunctional(mat[,1], mat[,2])
```

**getBIC***Get criterion value***Description**

Get criterion value

**Usage**

```
getBIC(outMixtComp)
getICL(outMixtComp)
```

**Arguments**

`outMixtComp` object of class *MixtCompLearn* or *MixtComp* obtained using `mixtCompLearn` or `mixtCompPredict` functions from `RMixtComp` package or `rmcMultiRun` from `RMixtCompIO` package.

**Value**

value of the criterion

**Author(s)**

Quentin Grimonprez

**See Also**

Other getter: `getCompletedData()`, `getEmpiricTik()`, `getParam()`, `getPartition()`, `getType()`

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))
model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))
algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
```

```

confidenceLevel = 0.95,
ratioStableCriterion = 0.95,
nStableCriterion = 10,
mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# get criterion
bic <- getBIC(resLearn)
icl <- getICL(resLearn)

```

**getCompletedData***Get the completed data from MixtComp object***Description**

Get the completed data from MixtComp object (does not manage functional models)

**Usage**

```
getCompletedData(outMixtComp, var = NULL, with.z_class = FALSE)
```

**Arguments**

- |              |  |
|--------------|--|
| outMixtComp  | object of class <i>MixtCompLearn</i> or <i>MixtComp</i> obtained using <code>mixtCompLearn</code> or <code>mixtCompPredict</code> functions from <code>RMixtComp</code> package or <code>rmcMultiRun</code> from <code>RMixtCompIO</code> package. |
| var          | Name of the variables for which to extract the completed data. Default is NULL (all variables are extracted)   |
| with.z_class | if TRUE, z_class is returned with the data.  |

**Value**

a matrix with the data completed by MixtComp (z\_class is in the first column and then variables are sorted in alphabetic order, it may differ from the original order of the data).

**Author(s)**

Quentin Grimonprez

**See Also**

Other getter: [getBIC\(\)](#), [getEmpiricTik\(\)](#), [getParam\(\)](#), [getPartition\(\)](#), [getType\(\)](#)

## Examples

```

require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

# add missing values
dataLearn$var1[12] = "?"
dataLearn$var2[72] = "?"

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# get completedData
completedData <- getCompletedData(resLearn)
completedData2 <- getCompletedData(resLearn, var = "var1")

```

getEmpiricTik

*Get the tik*

## Description

Get the a posteriori probability to belong to each class for each individual

## Usage

```

getEmpiricTik(outMixtComp)

getTik(outMixtComp, log = TRUE)

```

**Arguments**

- `outMixtComp` object of class *MixtCompLearn* or *MixtComp* obtained using `mixtCompLearn` or `mixtCompPredict` functions from `RMixtComp` package or `rmcMultiRun` from `RMixtCompIO` package.
- `log` if TRUE, `log(tik)` are returned

**Details**

`getTik` returns a posteriori probabilities computed with the returned parameters. `getEmpiricTik` returns an estimation based on the sampled  $z_i$  during the algorithm.

**Value**

a matrix containing the tik for each individuals (in row) and each class (in column).

**Author(s)**

Quentin Grimonprez

**See Also**

[heatmapTikSorted](#)

Other getter: `getBIC()`, `getCompletedData()`, `getParam()`, `getPartition()`, `getType()`

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)
```

```
# get tik
tikEmp <- getEmpiricTik(resLearn)
tik <- getTik(resLearn, log = FALSE)
```

**getParam***Get the estimated parameter***Description**

Get the estimated parameter

**Usage**

```
getParam(outMixtComp, var)

getProportion(outMixtComp)
```

**Arguments**

<code>outMixtComp</code>	object of class <i>MixtCompLearn</i> or <i>MixtComp</i> obtained using <code>mixtCompLearn</code> or <code>mixtCompPredict</code> functions from <code>RMixtComp</code> package or <code>rmcMultiRun</code> from <code>RMixtCompIO</code> package.
<code>var</code>	name of the variable to get parameter

**Value**

the parameter of the variable

**Author(s)**

Quentin Grimonprez

**See Also**

[plotDataBoxplot](#) [plotDataCI](#)  
 Other getter: [getBIC\(\)](#), [getCompletedData\(\)](#), [getEmpiricTik\(\)](#), [getPartition\(\)](#), [getType\(\)](#)

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
```

```

nClass = 2,
nInd = 100,
nbBurnInIter = 100,
nbIter = 100,
nbGibbsBurnInIter = 100,
nbGibbsIter = 100,
nInitPerClass = 3,
nSemTry = 20,
confidenceLevel = 0.95,
ratioStableCriterion = 0.95,
nStableCriterion = 10,
mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# get estimated parameters for variable var1
param <- getParam(resLearn, "var1")
prop <- getProportion(resLearn)

```

**getPartition***Get the estimated class from MixtComp object***Description**

Get the estimated class from MixtComp object

**Usage**

```
getPartition(outMixtComp, empiric = TRUE)
```

**Arguments**

- |             |  |
|-------------|--|
| outMixtComp | object of class <i>MixtCompLearn</i> or <i>MixtComp</i> obtained using <code>mixtCompLearn</code> or <code>mixtCompPredict</code> functions from <code>RMixtComp</code> package or <code>rmcMultiRun</code> from <code>RMixtCompIO</code> package. |
| empiric     | if TRUE, use the partition obtained at the end of the gibbs algorithm. If FALSE, use the partition obtained with the observed probabilities.   |

**Value**

a vector containing the estimated class for each individual.

**Author(s)**

Quentin Grimonprez

**See Also**

Other getter: `getBIC()`, `getCompletedData()`, `getEmpiricTik()`, `getParam()`, `getType()`

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))
model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))
algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# get class
estimatedClass <- getPartition(resLearn)
```

`getType`

*Names and Types Getters*

**Description**

`getType` returns the type output of a MixtComp object, `getModel` returns the model object, `getVarNames` returns the name for each variable

**Usage**

```
getType(outMixtComp, with.z_class = FALSE)

getModel(outMixtComp, with.z_class = FALSE)

getVarNames(outMixtComp, with.z_class = FALSE)
```

**Arguments**

- `outMixtComp` object of class *MixtCompLearn* or *MixtComp* obtained using `mixtCompLearn` or `mixtCompPredict` functions from `RMixtComp` package or `rmcMultiRun` from `RMixtCompIO` package.
- `with.z_class` if TRUE, the type of `z_class` is returned.

**Value**

a vector containing the type of models, names associated with each individual.

**Author(s)**

Quentin Grimonprez

**See Also**

Other getter: `getBIC()`, `getCompletedData()`, `getEmpiricTik()`, `getParam()`, `getPartition()`

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# get type
type <- getType(resLearn)

# get model object
model <- getModel(resLearn)

# get variable names
```

```
varNames <- getVarNames(resLearn)
```

**heatmapClass**

*Heatmap of the similarities between classes about clustering*

## Description

Heatmap of the similarities between classes about clustering

## Usage

```
heatmapClass(output, pkg = c("ggplot2", "plotly"), ...)
```

## Arguments

<b>output</b>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<b>pkg</b>	"ggplot2" or "plotly". Package used to plot
<b>...</b>	arguments to be passed to <i>plot_ly</i> . For <i>pkg</i> = "ggplot2", <i>addValues</i> = TRUE prints similarity values on the heatmap

## Details

The similarities between classes k and g is defined by  $1 - \text{Sigma}(k, g)$

$$\text{Sigma}(k, g)^2 = (1/n) * \sum_{i=1}^n (P(Z_i = k|x_i) - P(Z_i = g|x_i))^2$$

## Author(s)

Matthieu MARBAC

## See Also

[computeSimilarityClass](#)

Other plot: [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```

require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
heatmapClass(resLearn)

```

heatmapTikSorted

*Heatmap of the tik =  $P(Z_{-i}=k|x_{-i})$* 

## Description

Heatmap of the tik =  $P(Z_{-i}=k|x_{-i})$

## Usage

```
heatmapTikSorted(output, pkg = c("ggplot2", "plotly"), ...)
```

## Arguments

output	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
pkg	"ggplot2" or "plotly". Package used to plot
...	arguments to be passed to <i>plot_ly</i>

## Details

Observation are sorted according to the hard partition then for each component they are sorted by decreasing order of their tik's

## Author(s)

Matthieu MARBAC

## See Also

[getTik](#)

Other plot: [heatmapClass\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8)))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
heatmapTikSorted(resLearn)
```

---

heatmapVar	<i>Heatmap of the similarities between variables about clustering</i>
------------	---

---

## Description

Heatmap of the similarities between variables about clustering

## Usage

```
heatmapVar(output, pkg = c("ggplot2", "plotly"), ...)
```

## Arguments

- |        |  |
|--------|--|
| output | object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>          |
| pkg    | "ggplot2" or "plotly". Package used to plot  |
| ...    | arguments to be passed to <i>plot_ly</i> . For <i>pkg</i> = "ggplot2", <i>addValues</i> = TRUE prints similarity values on the heatmap |

## Details

The similarities between variables *j* and *h* is defined by *Delta(j,h)*

$$\Delta(j, h) = 1 - \sqrt{(1/n) * \sum_{i=1}^n \sum_{k=1}^K (P(Z_i = k|x_{ij}) - P(Z_i = k|x_{ih}))^2}$$

## Author(s)

Matthieu MARBAC

## See Also

[computeSimilarityVar](#)

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
```

```

nClass = 2,
nInd = 100,
nbBurnInIter = 100,
nbIter = 100,
nbGibbsBurnInIter = 100,
nbGibbsIter = 100,
nInitPerClass = 3,
nSemTry = 20,
confidenceLevel = 0.95,
ratioStableCriterion = 0.95,
nStableCriterion = 10,
mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
heatmapVar(resLearn)

```

**histMisclassif**      *Histogram of the misclassification probabilities*

## Description

Histogram of the misclassification probabilities

## Usage

```
histMisclassif(output, pkg = c("ggplot2", "plotly"), ...)
```

## Arguments

<b>output</b>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<b>pkg</b>	"ggplot2" or "plotly". Package used to plot
<b>...</b>	arguments to be passed to <code>plot_ly</code>

## Details

Missclassification probability of observation  $i$  is denoted  $\text{err}_i$   $\text{err}_i = 1 - \max_{k=1,\dots,K} P(Z_i=k|x_i)$   
 Histograms of  $\text{err}_i$ 's can be plotted for a specific class, all classes or every class

## Author(s)

Matthieu MARBAC

**See Also**

Other plot: `heatmapClass()`, `heatmapTikSorted()`, `heatmapVar()`, `plot.MixtComp()`, `plotConvergence()`, `plotDataBoxplot()`, `plotDataCI()`, `plotDiscrimClass()`, `plotDiscrimVar()`, `plotParamConvergence()`, `plotProportion()`

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8)))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
               var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
histMisclassif(resLearn)
```

plot.MixtComp

*Plot of a MixtComp object***Description**

Plot of a *MixtComp* object

**Usage**

```
## S3 method for class 'MixtComp'
plot(
  x,
  nVarMaxToPlot = 3,
  pkg = c("ggplot2", "plotly"),
```

```
plotData = c("CI", "Boxplot"),
...
)
```

## Arguments

x	<i>MixtComp</i> object
nVarMaxToPlot	number of variables to display
pkg	"ggplot2" or "plotly". Package used to plot
plotData	"CI" or "Boxplot". If "CI", uses <a href="#">plotDataCI</a> function. If "Boxplot", uses <a href="#">plotDataBoxplot</a>
...	extra parameter for <a href="#">plotDataCI</a>

## Author(s)

Quentin Grimonprez

## See Also

[mixtCompLearn](#) [mixtCompPredict](#)

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
               var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)
```

```
plot(resLearn)
```

plotConvergence	<i>Convergence of algorithm</i>
-----------------	---------------------------------

## Description

Plot the evolution of the completed loglikelihood during the SEM algorithm. The vertical line denotes the end of the burn-in phase.

## Usage

```
plotConvergence(output, ...)
```

## Arguments

- |        |   |
|--------|---|
| output | object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i> |
| ...    | graphical parameters  |

## Details

This function can be used to check the convergence and choose the parameters `nbBurnInIter` and `nbIter` from `mcStrategy`.

## Author(s)

Quentin Grimonprez

## See Also

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8)))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
```

```

nbIter = 100,
nbGibbsBurnInIter = 100,
nbGibbsIter = 100,
nInitPerClass = 3,
nSemTry = 20,
confidenceLevel = 0.95,
ratioStableCriterion = 0.95,
nStableCriterion = 10,
mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotConvergence(resLearn)

```

**plotDataBoxplot**      *Boxplot per class*

## Description

Display a boxplot (5

## Usage

```

plotDataBoxplot(
  output,
  var,
  class = 1:output$algo$nClass,
  grl = TRUE,
  pkg = c("ggplot2", "plotly"),
  ...
)

```

## Arguments

<b>output</b>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<b>var</b>	name of the variable
<b>class</b>	classes to plot
<b>grl</b>	if TRUE plot the general distribution of the data
<b>pkg</b>	"ggplot2" or "plotly". Package used to plot
<b>...</b>	other parameters (see <i>Details</i> )

## Details

For functional data, three other parameters are available:

- add.obs** if TRUE, observations are added to the plot. Default = FALSE.
- ylim** ylim of the plot.
- xlim** xlim of the plot.

## Author(s)

Matthieu MARBAC

## See Also

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))
model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))
algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotDataBoxplot(resLearn, "var1")
```

**plotDataCI***Mean and 95%-level confidence intervals per class*

## Description

Mean and 95%-level confidence intervals per class

## Usage

```
plotDataCI(
  output,
  var,
  class = 1:output$algo$nClass,
  grl = FALSE,
  pkg = c("ggplot2", "plotly"),
  ...
)
```

## Arguments

<b>output</b>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<b>var</b>	name of the variable
<b>class</b>	class to plot
<b>grl</b>	if TRUE plot the CI for the dataset and not only classes
<b>pkg</b>	"ggplot2" or "plotly". Package used to plot
<b>...</b>	other parameters (see <i>Details</i> )

## Details

For functional data, three other parameters are available:

**add.obs** if TRUE, observations are added to the plot. Default = FALSE.

**add.CI** if FALSE, confidence intervals are removed from the plot. Default = TRUE.

**xlim** xlim of the plot.

**ylim** ylim of the plot.

## Author(s)

Matthieu MARBAC

## See Also

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```

require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotDataCI(resLearn, "var1")

```

plotDiscrimClass

*Barplot of the discriminative power of the classes*

## Description

Barplot of the discriminative power of the classes

## Usage

```
plotDiscrimClass(output, ylim = c(0, 1), pkg = c("ggplot2", "plotly"), ...)
```

## Arguments

output	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
ylim	vector of length 2 defining the range of y-axis
pkg	"ggplot2" or "plotly". Package used to plot
...	arguments to be passed to <i>plot_ly</i>

## Details

The discriminative power of class k is defined by  $1 - D(k)$

$$D(k) = - \sum_{i=1}^n P(Z_i = k|x_i) \log(P(Z_i = k|x_i))/(n * \exp(-1))$$

## Author(s)

Matthieu MARBAC

## See Also

[computeDiscrimPowerClass](#)

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

plotDiscrimClass(resLearn)
```

---

plotDiscrimVar	<i>Barplot of the discriminative power of the variables</i>
----------------	---

---

**Description**

Barplot of the discriminative power of the variables

**Usage**

```
plotDiscrimVar(
  output,
  class = NULL,
  ylim = c(0, 1),
  pkg = c("ggplot2", "plotly"),
  ...
)
```

**Arguments**

output	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
class	NULL or a number of classes. If NULL, return the discriminative power of variables globally otherwise return the discriminative power of variables in the given class
ylim	vector of length 2 defining the range of y-axis
pkg	"ggplot2" or "plotly". Package used to plot
...	arguments to be passed to plot_ly

**Details**

The discriminative power of variable  $j$  is defined by  $1 - C(j)$

$$C(j) = - \sum_{k=1}^K \sum_{i=1}^n P(Z_i = k|x_{ij}) \ln(P(Z_i = k|x_{ij})) / (n * \log(K))$$

**Author(s)**

Matthieu MARBAC

**See Also**

[computeDiscrimPowerVar](#)

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotParamConvergence\(\)](#), [plotProportion\(\)](#)

## Examples

```

require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotDiscrimVar(resLearn)

plotDiscrimVar(resLearn, class = 1)

```

*plotParamConvergence Evolution of parameters*

## Description

Plot the evolution of estimated parameters after the burn-in phase.

## Usage

```
plotParamConvergence(output, var, ...)
```

## Arguments

<b>output</b>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<b>var</b>	name of the variable
<b>...</b>	graphical parameters

**Author(s)**

Quentin Grimonprez

**See Also**

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotProportion\(\)](#)

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8)))))

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotParamConvergence(resLearn, "var1")
plotParamConvergence(resLearn, "var2")
```

---

plotProportion      *Plot the mixture's proportions*

---

**Description**

Plot the mixture's proportions

**Usage**

```
plotProportion(output, pkg = c("ggplot2", "plotly"), ...)
```

**Arguments**

<code>output</code>	object returned by <i>mixtCompLearn</i> function from <i>RMixtComp</i> or <i>rmcMultiRun</i> function from <i>RMixtCompIO</i>
<code>pkg</code>	"ggplot2" or "plotly". Package used to plot
...	arguments to be passed to <code>plot_ly</code>

**Author(s)**

Quentin Grimonprez

**See Also**

Other plot: [heatmapClass\(\)](#), [heatmapTikSorted\(\)](#), [heatmapVar\(\)](#), [histMisclassif\(\)](#), [plot.MixtComp\(\)](#), [plotConvergence\(\)](#), [plotDataBoxplot\(\)](#), [plotDataCI\(\)](#), [plotDiscrimClass\(\)](#), [plotDiscrimVar\(\)](#), [plotParamConvergence\(\)](#)

**Examples**

```
require(RMixtCompIO) # for learning a mixture model
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))),

model <- list(var1 = list(type = "Gaussian", paramStr = ""),
              var2 = list(type = "Poisson", paramStr = ""))

algo <- list(
  nClass = 2,
  nInd = 100,
  nbBurnInIter = 100,
  nbIter = 100,
  nbGibbsBurnInIter = 100,
  nbGibbsIter = 100,
  nInitPerClass = 3,
  nSemTry = 20,
  confidenceLevel = 0.95,
  ratioStableCriterion = 0.95,
  nStableCriterion = 10,
  mode = "learn"
)

resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)

# plot
plotProportion(resLearn)
```

---

print.MixtComp	<i>Print Values</i>
----------------	---------------------

---

## Description

Print a *MixtComp* object

## Usage

```
## S3 method for class 'MixtComp'  
print(x, nVarMaxToPrint = 5, ...)
```

## Arguments

x	<i>MixtComp</i> object
nVarMaxToPrint	number of variables to display (including z_class)
...	parameter of head function

## Author(s)

Quentin Grimonprez

## See Also

`mixtCompLearn` `mixtCompPredict`

## Examples

```
require(RMixtCompIO) # for learning a mixture model  
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),  
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))  
  
model <- list(var1 = list(type = "Gaussian", paramStr = ""),  
               var2 = list(type = "Poisson", paramStr = ""))  
  
algo <- list(  
  nClass = 2,  
  nInd = 100,  
  nbBurnInIter = 100,  
  nbIter = 100,  
  nbGibbsBurnInIter = 100,  
  nbGibbsIter = 100,  
  nInitPerClass = 3,  
  nSemTry = 20,  
  confidenceLevel = 0.95,  
  ratioStableCriterion = 0.95,  
  nStableCriterion = 10,  
  mode = "learn"  
)
```

```
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)
print(resLearn)
```

**refactorCategorical**    *Rename a categorical value*

### Description

Rename a categorical value

### Usage

```
refactorCategorical(
  data,
  oldCateg = unique(data),
  newCateg = 1:length(oldCateg)
)
```

### Arguments

<code>data</code>	matrix/data.frame/vector containing the data
<code>oldCateg</code>	vector containing categories to change
<code>newCateg</code>	vector containing new categorical values

### Value

Data with new categorical values

### Author(s)

Quentin Grimonprez

### Examples

```
dat <- c("single", "married", "married", "divorced", "single")
refactorCategorical(dat, c("single", "married", "divorced"), 1:3)
```

---

summary.MixtComp      *MixtComp Object Summaries*

---

## Description

Summary of a *MixtComp* object

## Usage

```
## S3 method for class 'MixtComp'  
summary(object, ...)
```

## Arguments

object	<i>MixtComp</i> object
...	Not used.

## Author(s)

Quentin Grimonprez

## See Also

`mixtCompLearn` [print.MixtComp](#)

## Examples

```
require(RMixtCompIO) # for learning a mixture model  
dataLearn <- list(var1 = as.character(c(rnorm(50, -2, 0.8), rnorm(50, 2, 0.8))),  
                  var2 = as.character(c(rnorm(50, 2), rpois(50, 8))))  
  
model <- list(var1 = list(type = "Gaussian", paramStr = ""),  
               var2 = list(type = "Poisson", paramStr = ""))  
  
algo <- list(  
  nClass = 2,  
  nInd = 100,  
  nbBurnInIter = 100,  
  nbIter = 100,  
  nbGibbsBurnInIter = 100,  
  nbGibbsIter = 100,  
  nInitPerClass = 3,  
  nSemTry = 20,  
  confidenceLevel = 0.95,  
  ratioStableCriterion = 0.95,  
  nStableCriterion = 10,  
  mode = "learn"  
)
```

```
resLearn <- rmcMultiRun(algo, dataLearn, model, nRun = 3)  
summary(resLearn)
```

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