

# Package ‘CICA’

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**Type** Package

**Title** Clusterwise Independent Component Analysis

**Version** 0.1.0

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**Depends** ica, NMFN, R (>= 2.10)

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## Description

Clustering multi-subject resting state functional Magnetic Resonance Imaging data. This method enables the clustering of subjects based on multi-subject resting state functional Magnetic Resonance Imaging data. Objects are clustered based on similarities and differences in cluster-specific estimated components obtained by Independent Component Analysis.

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**URL** <https://hdl.handle.net/1887/35077>,  
<https://github.com/jeffreydurieux/CICA>

**NeedsCompilation** no

**Repository** CRAN

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**Description**

Main function to perform Clusterwise Independent Component Analysis

**Usage**

```
CICA(
  DataList,
  nStarts,
  nComp,
  nClus,
  scale = TRUE,
  center = TRUE,
  rational = NULL,
  maxiter = 100,
  verbose = TRUE
)
```

**Arguments**

DataList	a list of matrices
nStarts	number of multiple starts
nComp	number of ICA components per cluster
nClus	number of clusters
scale	scale each matrix to have an equal sum of squares
center	mean center matrices
rational	a rational starting seed, if NULL no rational starting seed is used
maxiter	maximum number of iterations for each start
verbose	print loss information to console

**Value**

CICA returns an object of `class` "CICA". It contains the estimated clustering, cluster specific component matrices and subject specific time course matrices

P	partitioning vector of size <code>length(DataList)</code>
Sr	list of size <code>nClus</code> , containing cluster specific independent components
Ais	list of size <code>length(DataList)</code> , containing subject specific time courses
Loss	loss function value of the best start
LossStarts	loss function values of all starts

**Author(s)**

Jeffrey Durieux

**Examples**

```
data('ExampleData', package = 'CICA')
output <- CICA(DataList = ExampleData, nStarts = 3, nComp = 5, nClus = 3, verbose = FALSE)
summary(output)
```

---

`ExampleData`*Example data for CICA*

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**Description**

Example data for CICA

**Usage**`data(ExampleData)`**Format**An object of class `list` of length 9.**Examples**`data(ExampleData)`

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`summary.CICA`*Summary method for class CICA*

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**Description**

Summarize a CICA analysis

**Usage**

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

**Arguments**

<code>object</code>	Object of the type produced by <a href="#">CICA</a>
<code>...</code>	Additional arguments

**Value**

`summary.CICA` returns an overview of the estimated clustering of a [CICA](#) analysis

PM	Partitioning matrix
tab	tabulation of the clustering
Loss	Loss function value of the solution

**Examples**

```
data('ExampleData', package = 'CICA')
output <- CICA(DataList = ExampleData, nStarts = 3, nComp = 5, nClus = 3, verbose = FALSE)
summary(output)
```

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