Package ‘FMmssnReg’

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Type Package

Title Regression Models with Finite Mixtures of Skew Heavy-Tailed Errors

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Imports mvtnorm

Description Fit linear regression models where the random errors follow a finite mixture of of Skew Heavy-Tailed Errors.

License GPL (>= 2)

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Description

This package contains a principal function that performs to estimate the parameters of a regression model considering an error that follows a finite mixture of Skew Heavy-Tailed Errors, using an analytically simple and efficient EM-type algorithm for iteratively computing maximum likelihood estimates of the parameters.

Details

Package: FMmsnReg
Type: Package
Version: 1.0
Date: 2016-03-30
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Author(s)

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References


See Also

FMmsnReg

Examples

#See examples for the FMmsnReg function linked above.
Description

Data on 102 male and 100 female athletes collected at the Australian Institute of Sport.

Format

This data frame contains the following columns:

- **Sex** (0 = male or 1 = female)
- **Ht** height (cm)
- **Wt** weight (kg)
- **LBM** lean body mass
- **RCC** red cell count
- **WCC** white cell count
- **He** Hematocrit
- **Hg** Hemoglobin
- **Ferr** plasma ferritin concentration
- **BMI** body mass index, weight/height^2
- **SSF** sum of skin folds
- **Bfat** Percent body fat
- **Label** Case Labels
- **Sport** Sport

References


Examples

```r
#Load the data
library(FMsmsnReg)
data(ais)
attach(ais)
#Set the response y and covariate x
x1 <- cbind(1,SSF,Ht)
y <- Bfat

#Fits a linear Regression Model with Finite Mixtures of Skew t
parST <- FMsmmsnReg(y, x1, g=2, get.init = TRUE, criteria = TRUE,
group = FALSE,family = "Skew.t", error = 10^-4,
```

## Fits a linear Regression Models with Finite Mixtures of Skew normal
parSN <- FMsmnReg(y, x1, g=2, get.init = TRUE, criteria = TRUE, group = FALSE, family = "Skew.normal", error = 10^-4, iter.max = 5000, obs.prob= FALSE, kmeans.param = NULL, show.converge=FALSE, cp=0.5)

## Fits a linear Regression Models with Finite Mixtures of Skew Contaminated Normal
parCN <- FMsmnReg(y, x1, g=2, get.init = TRUE, criteria = TRUE, group = FALSE, family = "Skew.cn", error = 10^-4, iter.max = 5000, obs.prob= FALSE, kmeans.param = NULL, show.converge=FALSE, cp=0.5)

### Description
Fits Linear Regression Models with Finite Mixtures of Skew Heavy-Tailed Errors

### Arguments
- **y**: the response matrix (dimension nx1)
- **x1**: Matrix or vector of covariates.
- **Abetas**: Parameters of vector regression dimension \((p + 1)\) include intercept
- **medj**: a list of \(g\) arguments of vectors of initial values (dimension \(p\)) for the location parameters
- **sigma2**: a list of \(g\) arguments of matrices of initial values (dimension \(pxp\)) for the scale parameters
- **shape**: a list of \(g\) arguments of vectors of initial values (dimension \(p\)) for the skewness parameters
- **pii**: Initial value for the EM algorithm. Each of them must be a vector of length \(g\). (the algorithm considers the number of components to be adjusted based on the size of these vectors)
- **g**: the number of cluster to be considered in fitting
- **get.init**: if TRUE, the initial values are generated via k-means
- **criteria**: It indicates if are calculated the criterion selection methods (AIC, BIC, EDC and ICL)
**group**  
if TRUE, the vector with the classification of the response is returned

**family**  
distribution family to be used in fitting (Skew.t", "Skew.cn", "Skew.slash", "Skew.normal")

**error**  
define the stopping criterion of the algorithm

**iter.max**  
the maximum number of iterations of the EM algorithm

**obs.prob**  
if TRUE, the posterior probability of each observation belonging to one of the g groups is reported

**kmeans.param**  
a list with alternative parameters for the kmeans function when generating initial values, list(iter.max = 10, n.start = 1, algorithm = "Hartigan-Wong")

**show.convergence**  
graphics of convergence for the parameters

**cp**  
Cut Point

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**Value**

The function returns a list with 16 elements detailed as

- **iter**: Number of iterations.
- **criteria**: Attained criteria value.
- **convergence**: Convergence reached or not.
- **nu**: Location parameter estimate.
- **sigma2**: Scale parameter estimate.
- **lambda**: Shape parameter estimate.
- **pii**: Weight parameter estimate.
- **nu**: Estimated degrees of freedom parameter.
- **SE**: Standard Error estimates, if the output shows NA the function does not provide the standard error for this parameter.
- **table**: Table containing the inference for the estimated parameters.
- **loglik**: Log-likelihood value.
- **AIC**: Akaike information criterion.
- **BIC**: Bayesian information criterion.
- **EDC**: Efficient Determination Criterion.
- **ICL**: Information Completed Likelihood.
- **time**: Processing time.

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and Victor Hugo Lachos <hlachos@ime.unicamp.br>
References


See Also

FMsmnReg, ais, horses

Examples

#See examples for the FMsmnReg function linked above.

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horses                  Horse Racing at Eagle Farm data

Description

Results of horse races at Eagle Farm, Brisbane, on 31 August 1998. The data, collected by Donald Forbes for his MS305 Data Analysis Project, give results for each horse in a sequence of 8 races.

Format

This data frame contains the following columns:

- **Position** (Finishing position)
- **Starters** Number of horses in race
- **Last** Finishing position in last race
- **Since** Days since last race
- **Number** Identifying number of horse in race
- **Carried** Weight carried
- **Weight** Handicap weight
- **Barrier** Barrier position at start of race
- **Distance** Length of race
- **Lengths** Number of lengths that horse finished from winner
- **Odds** Starting odds
- **Starts** Number of races previously started in
- **Age** Age of horse in years
- **Ratio** Proportion of wins in previous starts
Examples

```r
# Load the data
library(FMsmsnReg)
data(horses)
attach(horses)

# Set the response y and covariate x
x1 <- cbind(1, Last, Carried)
y <- Position

## Fits a linear Regression Model with Finite Mixtures of Skew Contaminated Normal
parCN <- FMsmReg(y, x1, g=2, get.init = TRUE, criteria = TRUE, group = FALSE, family = "Skew.cn", error = 10^-4, iter.max = 5000, obs.prob = FALSE, kmeans.param = NULL, show.converge = FALSE, cp=0.5)

## Fits a linear Regression Model with Finite Mixtures of Skew normal
parSN <- FMsmReg(y, x1, g=2, get.init = TRUE, criteria = TRUE, group = FALSE, family = "Skew.normal", error = 10^-4, iter.max = 5000, obs.prob = FALSE, kmeans.param = NULL, show.converge = FALSE, cp=0.5)
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