

# Package ‘ltmix’

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**Type** Package

**Title** Left-Truncated Mixtures of Gamma, Weibull, and Lognormal Distributions

**Version** 0.2.1

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**Description** Mixture modelling of one-dimensional data using combinations of left-truncated Gamma, Weibull, and Lognormal Distributions. Blostein, Martin & Miljkovic, Tatjana. (2019) <doi:10.1016/j.insmatheco.2018.12.001>.

**License** GPL-3

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**NeedsCompilation** no

**Repository** CRAN

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## R topics documented:

|                        |   |
|------------------------|---|
| createLtmObj . . . . . | 2 |
| ltmix . . . . .        | 2 |
| ltmm . . . . .         | 3 |
| ltmmCombo . . . . .    | 4 |
| secura . . . . .       | 6 |

|              |          |
|--------------|----------|
| <b>Index</b> | <b>8</b> |
|--------------|----------|

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|--------------|---|
| createLtmObj | <i>Create an ltm model object given data and parameters</i> |
|--------------|---|

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### Description

This function is useful for comparing models produced using the ltmix package to models fit using other, or for computing fit criteria and risk measures for a known set of parameters.

### Usage

```
createLtmObj(x, distributions, trunc, Pars, Pi, npars = NULL)
```

### Arguments

|               |   |
|---------------|---|
| x             | data vector   |
| distributions | densities to combine  |
| trunc         | left truncation point (optional)  |
| Pars          | list of length G of parameter values  |
| Pi            | vector of length G of component proportions   |
| npars         | Can optionally be used to overwrite the number of free parameters (used in the calculation of AIC & BIC), if the model has additional constraints |

### Value

An ltm model object

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|       |  |
|-------|--|
| ltmix | <i>ltmix: Left-Truncated Mixtures of Gamma, Weibull, and Lognormal Distributions</i> |
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### Description

Mixture modelling of one-dimensional data using combinations of left-truncated Gamma, Weibull, and Lognormal Distributions.

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ltmm

*Fit a Left-truncated mixture model (LTMM)*


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### Description

This function generates a mixture model combining left-truncated lognormal, gamma, and weibull distributions

### Usage

```
ltmm(
  x,
  G,
  distributions,
  trunc = NULL,
  EM_init_method = "emEM",
  EM_starts = 5,
  init_pars = NULL,
  init_pi = NULL,
  init_classes = NULL,
  one_group_reps = 50,
  eps = 1e-06,
  max.it = 1000,
  verbose = FALSE
)
```

### Arguments

|                |  |
|----------------|--|
| x              | data vector  |
| G              | number of components   |
| distributions  | densities to combine   |
| trunc          | left truncation point (optional)   |
| EM_init_method | initialization method for EM algorithm                                       |
| EM_starts      | number of random starts for initialization of EM algorithm. (only for G > 1) |
| init_pars      | initial parameter values (list of length G)                                  |
| init_pi        | manually specified initial component proportions (for init_method=specified) |
| init_classes   | manually specified initial classes. will overwrite init_pars and init_pi     |
| one_group_reps | number of random starts for each numerical optimization in 1-component model |
| eps            | stopping tolerance for EM algorithm  |
| max.it         | maximum number of iterations of EM algorithm                                 |
| verbose        | print information as fitting progresses?                                     |

**Value**

An ltmm model object, with the following properties:

**x** Copy of the input data

**distributions** The selected distributions

**trunc** The left truncation value, if specified

**fitted\_pdf** The probability density function of the fitted model

**fitted\_cdf** The cumulative density function of the fitted model

**VaR** The value-at-risk of the fitted model (function with p taken as onl yargument)

**ES** The expected shortfall of the fitted model (function with p taken as onl yargument)

**G** The number of components in the model

**Pi** The estimated probabillites of component membership

**Pars** The estimated model parameters

**ll** The log-likelihood of the fitted model

**bic** The BIC of the fitted model

**aic** The AIC of the fitted model

**id** The MAP component membership for each observation

**iter** The number of iterations until convergence for the EM algorithm

**npars** The total number of model parameters for the fitted model

**ll.history** The value of log-likelihood at each iteration of the EM algorithm

**Examples**

```
x <- securaloss
fit <- ltmm(x, G = 2, distributions = c('gamma', 'gamma', 'weibull'), trunc = 1.2e6)
summary(fit)
plot(fit)
```

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ltmmCombo

*Fit a Left-truncated mixture model (LTMM)*


---

**Description**

This function fits a family of finite mixture models using every combination of the left-truncated lognormal, gamma, and weibull distributions.

**Usage**

```

ItmmCombo(
  x,
  G,
  distributions = c("lognormal", "gamma", "weibull"),
  trunc = NULL,
  EM_init_method = "emEM",
  EM_starts = 5,
  init_pars = NULL,
  init_pi = NULL,
  init_classes = NULL,
  one_group_reps = 50,
  eps = 1e-06,
  max.it = 1000,
  verbose = FALSE,
  parallel = FALSE,
  cores = NULL,
  save_each_fit = FALSE
)

```

**Arguments**

|                |  |
|----------------|--|
| x              | data vector  |
| G              | number of components   |
| distributions  | densities to combine   |
| trunc          | left truncation point (optional)   |
| EM_init_method | initialization method for EM algorithm   |
| EM_starts      | number of random starts for initialization of EM algorithm. (only for G > 1)   |
| init_pars      | initial parameter values (list of length G)                                    |
| init_pi        | manually specified initial component proportions (for init_method=specified)   |
| init_classes   | manually specified initial classes. will overwrite init_pars and init_pi       |
| one_group_reps | number of random starts for each numerical optimization in 1-component model   |
| eps            | stopping tolerance for EM algorithm  |
| max.it         | maximum number of iterations of EM algorithm                                   |
| verbose        | print information as fitting progresses?                                       |
| parallel       | fit models in parallel?  |
| cores          | number of processes used for parallel computation. if NULL detect.cores() used |
| save_each_fit  | save each model as it is produced, in a time-stamped directory (safer)         |

**Value**

An ItmmCombo model object, with the following properties:

**x** Copy of the input data

**distributions** The selected distributions  
**combos** List of all combinations of distributions considered  
**all.fits** List of all ltmm fit objects  
**all.bic** Vector of BIC values for each model  
**best.bic.fit** The best ltmm fit by BIC  
**best.bic** The best BIC value of all fits  
**best.bic.combo** The combination of distributions used for the best fit by BIC  
**all.aic** Vector of AIC value for each model  
**best.aic.fit** The best ltmm fit by AIC  
**best.aic** The best AIC value of all fits  
**best.aic.combo** The combination of distributions used for the best fit by AIC  
**all.ll** Vector of log-likelihood value for each model  
**summary\_table** Table summarizing the AIC, BIC, LL, and risk measures for each fitted model

## References

Blostein, Martin & Miljkovic, Tatjana. (2019). On modeling left-truncated loss data using mixtures of distributions. *Insurance Mathematics and Economics*. 85. 35-46. 10.1016/j.insmatheco.2018.12.001.

## Examples

```
x <- securaData$Loss

fits_GL <- ltmmCombo(x, G = 2, distributions = c('gamma', 'lognormal'), trunc = 1.2e6)
summary(fits_GL)
```

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secura

*The Secura Belgian Re Data*

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## Description

"The Secura Belgian Re data set contains automobile claims from 1988 until 2001, which are at least as large as 1,200,000 Euros." (Beirlant, Goegebeur, Segers & Teugels, 2004).

## Usage

secura

## Format

An object of class `data.frame` with 371 rows and 2 columns.

## References

Beirlant, J., Goegebeur Y., Segers, J., & Teugels, J. Statistics of extremes : theory and applications. Hoboken, NJ: Wiley, 2004. Print.

<https://lstat.kuleuven.be/Wiley/>

# Index

## \* datasets

secura, 6

createLtmObj, 2

ltmix, 2

ltmm, 3

ltmmCombo, 4

secura, 6