

# Package ‘ltmix’

February 23, 2024

**Type** Package

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

**Version** 0.2.2

**Author** Martin Blostein <[martin.blostein@gmail.com](mailto:martin.blostein@gmail.com)> and Tatjana Miljkovic <[miljkot@miamioh.edu](mailto:miljkot@miamioh.edu)>

**Maintainer** Martin Blostein <[martin.blostein@gmail.com](mailto:martin.blostein@gmail.com)>

**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](https://doi.org/10.1016/j.insmatheco.2018.12.001)>.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.1

**Imports** gtools, pracma

**Depends** R (>= 3.5.0)

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2024-02-23 13:20:02 UTC

## R topics documented:

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

**Index**

**8**

`createLtmObj`*Create an ltm model object given data and parameters***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**

<code>x</code>	data vector
<code>distributions</code>	densities to combine
<code>trunc</code>	left truncation point (optional)
<code>Pars</code>	list of length G of parameter values
<code>Pi</code>	vector of length G of component proportions
<code>npars</code>	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

`ltmix`*ltmix: Left-Truncated Mixtures of Gamma, Weibull, and Lognormal Distributions***Description**

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

---

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

---

## 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 algoithm
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 probabilites 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 <- secura$Loss

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

summary(fit)
plot(fit)
```

## Description

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

**Usage**

```
ltmmCombo(
  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 algoithm
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 ltmmCombo 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 <- secura$Loss

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

secura

*The Secura Belgian Re Data*

## 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.

# Index

\* **datasets**

secura, [6](#)

\_PACKAGE (ltmix), [2](#)

createLtmObj, [2](#)

ltmix, [2](#)

ltmm, [3](#)

ltmmCombo, [4](#)

secura, [6](#)