

# Package ‘ltmix’

October 13, 2022

**Type** Package

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

**Version** 0.2.1

**Author** Martin Blostein <martin.blostein@gmail.com> and Tatjana Miljkovic <miljkot@miamioh.edu>

**Maintainer** Martin Blostein <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>.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Imports** gtools, pracma

**Depends** R (>= 3.5.0)

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2021-07-05 06:10:05 UTC

## R topics documented:

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

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

---

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

---

### 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

---

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

---

### 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 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\_cfd** 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)
```

---

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**

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

---

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.

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

# Index

## \* datasets

secura, 6

createLtmObj, 2

ltmix, 2

ltmm, 3

ltmmCombo, 4

secura, 6