

Package ‘GSM’

April 17, 2009

Title Gamma Shape Mixture

Description This package implements a Bayesian approach for estimation of a mixture of gamma distributions in which the mixing occurs over the shape parameter. This family provides a flexible and novel approach for modeling heavy-tailed distributions, it is computationally efficient, and it only requires to specify a prior distribution for a single parameter.

Version 0.1-2

Date 2007-08-10

Author Sergio Venturini

Maintainer Sergio Venturini <sergio.venturini@unibocconi.it>

Depends gtools

License GPL (>= 2)

URL <http://www.bepress.com/jhubiostat/paper124>

Repository CRAN

Date/Publication 2007-08-09 16:48:51

R topics documented:

GSM-package	2
allcurves.q	2
gsm	3
gsm.plot	4
gsm.theta	6
GSMDist	7
prob.predict	8
Index	10

`GSM-package`*Estimation of a Gamma Shape Mixture Model*

Description

This package implements a Bayesian approach for estimation of a mixture of gamma distributions in which the mixing occurs over the shape parameter. This family provides a flexible and novel approach for modeling heavy-tailed distributions, it is computationally efficient, and it only requires to specify a prior distribution for a single parameter. See Venturini et al. (2006).

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

[gsm](#), [gsm.theta](#).

`allcurves.q`*Utility function*

Description

Utility function for plotting a Gamma Shape Mixture Model density.

Usage

```
allcurves.q(postdata, perc)
```

Arguments

<code>postdata</code>	matrix containing of a mixture's density posterior draws.
<code>perc</code>	percentile.

Details

This is a utility function used to generate the credibility bands for a Gamma Shape Mixture density within [gsm.plot](#).

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

See Also

[gsm.plot.](#)

gsm

Estimation of a Gamma Shape Mixture Model (GSM)

Description

This function provides the inferential algorithm to estimate a mixture of gamma distributions in which the mixing occurs over the shape parameter. It implements the collapsing approach for the GSM model, as discussed in Venturini et al. (2006).

Usage

```
gsm(y, J, G, M, a, b, alpha)
```

Arguments

y	vector of data.
J	number of mixture components.
G	number of points where to evaluate the GSM density.
M	number of MCMC runs.
a	hyperparameter of the rate parameter prior distribution.
b	hyperparameter of the rate parameter prior distribution.
alpha	hyperparameter of the mixture's weights prior distribution.

Details

Suggestions on how to choose J , a and b are provided in Venturini et al. (2006). In that work the α vector is always set at $(1/J, \dots, 1/J)$, but here one is free to choose the value of the generic element of α .

Value

List with the following components:

J	number of mixture components used in the GSM model.
a	hyperparameter of the rate parameter prior distribution used in the GSM model.
b	hyperparameter of the rate parameter prior distribution used in the GSM model.
alpha	hyperparameter of the mixture's weights prior distribution used in the GSM model.

<code>ff</code>	matrix containing the posterior draws for the mixture's density.
<code>y.grid</code>	vector of values used to evaluate the GSM density.
<code>theta</code>	vector containing the posterior draws for the mixture's rate parameter.
<code>label</code>	matrix containing the posterior draws for the mixture's hidden label.
<code>weight</code>	matrix containing the posterior draws for the mixture's weights.

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

[gsm.theta](#), [gsm.plot](#).

Examples

```
set.seed(2040)
y <- rgsm(500, c(.1, .3, .4, .2), 1)
burnin <- 100
J <- 250
gsm.out <- gsm(y, J, 300, burnin+500, 6500, 340, 1/J)
gsm.plot(gsm.out, y, ndens=5, nbins=20, histogram=TRUE)
```

`gsm.plot`

Plot of a Gamma Shape Mixture Model

Description

This function plots the output of a Gamma Shape Mixture estimation procedure.

Usage

```
gsm.plot(v, y, ndens=5, xlim=c(min(y), max(y)), ylim=c(0, max(v$ff)), xlab="x", ylab="de
```

Arguments

<code>v</code>	list returned by the <code>gsm</code> or <code>gsm.theta</code> functions.
<code>y</code>	vector of data to use in the analysis.
<code>ndens</code>	number of simulated density curves to plot.
<code>xlim</code>	the x limits (x1, x2) of the plot.
<code>ylim</code>	the y limits of the plot.
<code>xlab</code>	a title for the x axis.
<code>ylab</code>	a title for the y axis.
<code>nbin</code>	number of bins for the histogram.
<code>histogram</code>	logical; if TRUE the histogram is plotted on the figure.
<code>bands</code>	logical; if TRUE the 95% credibility bands are overlaid on the density graph.

Details

To produce a standard histogram with the estimated density curve superimposed on it, simply set `ndens` to 0 and `histogram` to TRUE.

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

`gsm`, `gsm.theta`, `rgsm`.

Examples

```
set.seed(2040)
y <- rgsm(500, c(.1, .3, .4, .2), 1)
burnin <- 100
J <- 250
gsm.out <- gsm(y, J, 300, burnin+500, 6500, 340, 1/J)
par(mfrow=c(2, 2))
gsm.plot(gsm.out, y, ndens=5, nbin=20)
gsm.plot(gsm.out, y, ndens=0, nbin=20, histogram=TRUE)
gsm.plot(gsm.out, y, ndens=0, nbin=20, histogram=TRUE, bands=TRUE)
gsm.plot(gsm.out, y, ndens=5, nbin=20, histogram=TRUE, bands=TRUE)
```

gsm.theta

Estimation of a Gamma Shape Mixture Model (GSM)

Description

This function provides the inferential algorithm to estimate a mixture of gamma distributions in which the mixing occurs over the shape parameter. It implements the standard approach for the GSM model, as discussed in Venturini et al. (2006).

Usage

```
gsm.theta(y, J, G, M, a, b, alpha)
```

Arguments

y	vector of data.
J	number of mixture components.
G	number of points where to evaluate the GSM density.
M	number of MCMC runs.
a	hyperparameter of the rate parameter prior distribution.
b	hyperparameter of the rate parameter prior distribution.
alpha	hyperparameter of the mixture's weights prior distribution.

Details

Suggestions on how to choose J , a and b are provided in Venturini et al. (2006). In that work the α vector is always set at $(1/J, \dots, 1/J)$, but here one is free to choose the value of the generic element of α .

Value

List with the following components:

J	number of mixture components used in the GSM model.
a	hyperparameter of the rate parameter prior distribution used in the GSM model.
b	hyperparameter of the rate parameter prior distribution used in the GSM model.
alpha	hyperparameter of the mixture's weights prior distribution used in the GSM model.
ff	matrix containing the posterior draws for the mixture's density.
y.grid	vector of values used to evaluate the GSM density.
theta	vector containing the posterior draws for the mixture's rate parameter.
label	matrix containing the posterior draws for the mixture's hidden label.
weight	matrix containing the posterior draws for the mixture's weights.

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

[gsm](#), [gsm.plot](#).

Examples

```
set.seed(2040)
y <- rgsm(500, c(.1, .3, .4, .2), 1)
burnin <- 100
J <- 250
gsm.out <- gsm.theta(y, J, 300, burnin+500, 6500, 340, 1/J)
gsm.plot(gsm.out, y, ndens=0, nbin=20, histogram=TRUE)
```

GSMDist

Utility function

Description

Density function evaluation for a Gamma Shape Mixture Model.

Usage

```
dgsm(x, weight, rateparam)
pgsm(q, weight, rateparam)
rgsm(n, weight, rateparam)
```

Arguments

<code>x</code> , <code>q</code>	vector of quantiles.
<code>n</code>	number of observations.
<code>weight</code>	vector of mixture weights.
<code>rateparam</code>	reciprocal of the shape parameter, as in GammaDist .

Details

The parametrisation implemented in this function is described in Venturini et al. (2006).

Value

`dgsm` gives the density, `pgsm` gives the distribution function, and `rgsm` generates random deviates.

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

`dgamma`, `pgamma`, `rgamma`.

prob.predict

Tail probability estimation for a Gamma Shape Mixture Model

Description

This function allows to estimate the tail probability of a Gamma Shape Mixture Model using the output of the `gsm` or `gsm.theta` procedures.

Usage

```
prob.predict(mcmc.w, mcmc.theta, thresh)
```

Arguments

<code>mcmc.w</code>	matrix of the mixture's weights posterior draws; it is part of the output of the <code>gsm</code> or <code>gsm.theta</code> functions.
<code>mcmc.theta</code>	vector of the mixture's rate parameter posterior draws; it is part of the output of the <code>gsm</code> or <code>gsm.theta</code> functions.
<code>thresh</code>	threshold value.

Details

The tail probability is estimated by applying the standard Rao-Blackwellized estimator on the Gibbs sampling realizations obtained through the `gsm` or `gsm.theta` procedures.

Value

A numerical vector containing the posterior draws for the tail probability exceeding the value of `thresh`.

Author(s)

Sergio Venturini (sergio.venturini@unibocconi.it)

References

Venturini, S., Dominici, F., and Parmigiani, G., "Gamma Shape Mixtures for Heavy-Tailed Distributions" (December 2006). Johns Hopkins University, Dept. of Biostatistics Working Papers. Working Paper 124. <http://www.bepress.com/jhubiostat/paper124>

See Also

[gsm](#), [gsm.plot](#).

Examples

```
set.seed(2040)
y <- rgsm(500,c(.1, .3, .4, .2),1)
burnin <- 100
J <- 250
gsm.out <- gsm(y, J, 300, burnin+500, 6500, 340, 1/J)
thresh <- c(0.1, 0.5, 0.75, 1, 2)
tail.prob.est <- rep(NA, length(thresh))
tail.prob.true <- rep(NA, length(thresh))
for (i in 1:length(thresh)){
  tail.prob.est[i] <- mean(prob.predict(gsm.out$weight[(burnin+1):600,], gsm.out$theta[(burnin+1):600,]))
  tail.prob.true[i] <- sum(y>thresh[i])/length(y)
}
qqplot(tail.prob.true, tail.prob.est, main="Q-Q plot of true vs. estimated tail probability")
abline(0, 1, lty=2)
```

Index

*Topic **distribution**

allcurves.q, 2
gsm, 3
GSM-package, 1
gsm.plot, 4
gsm.theta, 5
GSMDist, 7
prob.predict, 8

*Topic **models**

gsm, 3
GSM-package, 1
gsm.plot, 4
gsm.theta, 5
prob.predict, 8

allcurves.q, 2

dgamma, 7
dgs, 7
dgs(GSMDist), 7

GammaDist, 7
gsm, 2, 3, 4-6, 8
GSM-package, 1
gsm.plot, 2, 4, 4, 6, 8
gsm.theta, 2, 4, 5, 5, 8
GSMDist, 7

pgamma, 7
pgs, 7
pgs(GSMDist), 7
prob.predict, 8

rgamma, 7
rgs, 5, 7
rgs(GSMDist), 7