

# Package ‘BayesianLasso’

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**Type** Package

**Title** Bayesian Lasso Regression and Tools for the Lasso Distribution

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**Description** Implements Bayesian Lasso regression using efficient Gibbs sampling algorithms, including modified versions of the Hans and Park Casella (PC) samplers. Includes functions for working with the Lasso distribution, such as its density, cumulative distribution, quantile, and random generation functions, along with moment calculations. Also includes a function to compute the Mills ratio. Designed for sparse linear models and suitable for high-dimensional regression problems.

**License** GPL-3

**Imports** Rcpp (>= 1.0.12)

**LinkingTo** Rcpp, RcppNumerical, RcppArmadillo, RcppEigen

**RoxygenNote** 7.3.2

**Encoding** UTF-8

**URL** <https://garhtarr.github.io/BayesianLasso/>,  
<https://github.com/garhtarr/BayesianLasso>

**VignetteBuilder** knitr, rmarkdown

**Suggests** knitr, rmarkdown, monomvn, posterior, rstan, bayesreg,  
testthat (>= 3.0.0), MASS

**Config/testthat/edition** 3

**BugReports** <https://github.com/garhtarr/BayesianLasso/issues>

**NeedsCompilation** yes

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## Contents

LassoDistribution . . . . .	2
normalize . . . . .	5
<b>Index</b>	<b>7</b>

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LassoDistribution	<i>The Lasso Distribution</i>
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## Description

Provides functions related to the Lasso distribution, including the normalizing constant, probability density function, cumulative distribution function, quantile function, and random number generation for given parameters  $a$ ,  $b$ , and  $c$ . Additional utilities include the Mills ratio, expected value, and variance of the distribution. The package also implements modified versions of the Hans and Park–Casella Gibbs sampling algorithms for Bayesian Lasso regression.

## Usage

```
zlasso(a, b, c, logarithm)
dlasso(x, a, b, c, logarithm)
plasso(q, a, b, c)
qlasso(p, a, b, c)
rlasso(n, a, b, c)
elasso(a, b, c)
vlasso(a, b, c)
mlasso(a, b, c)
MillsRatio(d)
Modified_Hans_Gibbs(X, y, beta_init, a1, b1, u1, v1,
                    nsamples, lambda_init, sigma2_init, thin, verbose,
                    tune_lambda2, rao_blackwellization)
Modified_PC_Gibbs(X, y, a1, b1, u1, v1,
                  nsamples, lambda_init, sigma2_init, thin, verbose)
```

## Arguments

$x, q$	Vector of quantiles (vectorized).
$p$	Vector of probabilities.
$a$	Vector of precision parameter which must be non-negative.
$b$	Vector of off set parameter.

<code>c</code>	Vector of tuning parameter which must be non-negative values.
<code>n</code>	Number of observations.
<code>logarithm</code>	Logical. If TRUE, probabilities are returned on the log scale.
<code>d</code>	A scalar numeric value. Represents the point at which the Mills ratio is evaluated.
<code>X</code>	Design matrix (numeric matrix).
<code>y</code>	Response vector (numeric vector).
<code>a1</code>	Shape parameter of the prior on $\lambda^2$ .
<code>b1</code>	Rate parameter of the prior on $\lambda^2$ .
<code>u1</code>	Shape parameter of the prior on $\sigma^2$ .
<code>v1</code>	Rate parameter of the prior on $\sigma^2$ .
<code>nsamples</code>	Number of Gibbs samples to draw.
<code>beta_init</code>	Initial value for the model parameter $\beta$ .
<code>lambda_init</code>	Initial value for the shrinkage parameter $\lambda^2$ .
<code>sigma2_init</code>	Initial value for the error variance $\sigma^2$ .
<code>thin</code>	Thinning interval for the MCMC chain. Only every ‘thin’-th draw is stored. Default is 1 (no thinning).
<code>verbose</code>	Integer. If greater than 0, progress is printed every verbose iterations during sampling. Set to 0 to suppress output.
<code>tune_lambda2</code>	Logical; if TRUE (default), the tuning parameter $\lambda^2$ is estimated during sampling.
<code>rao_blackwellization</code>	Logical; if TRUE, Rao–Blackwellization is applied to improve posterior estimation. Default is FALSE.

## Details

If  $X \sim \text{Lasso}(a, b, c)$  then its density function is:

$$p(x; a, b, c) = Z^{-1} \exp\left(-\frac{1}{2}ax^2 + bx - c|x|\right)$$

where  $x \in \mathbb{R}$ ,  $a > 0$ ,  $b \in \mathbb{R}$ ,  $c > 0$ , and  $Z$  is the normalizing constant.

More details are included for the CDF, quantile function, and normalizing constant in the original documentation.

## Value

- `zlasso`, `dlasso`, `plasso`, `qlasso`, `rlasso`, `elasso`, `vlasso`, `mlasso`, `MillsRatio`: return the corresponding scalar or vector values related to the Lasso distribution and a numeric value representing the Mills ratio.
- `Modified_Hans_Gibbs`: returns a list containing:
  - `mBeta` Matrix of MCMC samples for the regression coefficients  $\beta$ , with `nsamples` rows and `p` columns.

- vsigma2 Vector of MCMC samples for the error variance  $\sigma^2$ .
- vlambda2 Vector of MCMC samples for the shrinkage parameter  $\lambda^2$ .
- mA Matrix of sampled values for parameter  $a_j$  of the Lasso distribution for each  $\beta_j$ .
- mB Matrix of sampled values for parameter  $b_j$  of the Lasso distribution for each  $\beta_j$ .
- mC Matrix of sampled values for parameter  $c_j$  of the Lasso distribution for each  $\beta_j$ .
- Modified\_PC\_Gibbs: returns a list containing:
  - mBeta Matrix of MCMC samples for the regression coefficients  $\beta$ .
  - vsigma2 Vector of MCMC samples for the error variance  $\sigma^2$ .
  - vlambda2 Vector of MCMC samples for the shrinkage parameter  $\lambda^2$ .
  - mM Matrix of estimated means of the full conditional distributions of each  $\beta_j$ .
  - mV Matrix of estimated variances of the full conditional distributions of each  $\beta_j$ .
  - va\_til Vector of estimated shape parameters for the full conditional inverse-gamma distribution of  $\sigma^2$ .
  - vb\_til Vector of estimated rate parameters for the full conditional inverse-gamma distribution of  $\sigma^2$ .
  - vu\_til Vector of estimated shape parameters for the full conditional inverse-gamma distribution of  $\lambda^2$ .
  - vv\_til Vector of estimated rate parameters for the full conditional inverse-gamma distribution of  $\lambda^2$ .

### See Also

[normalize](#) for preprocessing input data before applying the samplers.

### Examples

```
a <- 2; b <- 1; c <- 3
x <- seq(-3, 3, length.out = 1000)
plot(x, dlasso(x, a, b, c, logarithm = FALSE), type = 'l')

r <- rlasso(1000, a, b, c)
hist(r, breaks = 50, probability = TRUE, col = "grey", border = "white")
lines(x, dlasso(x, a, b, c, logarithm = FALSE), col = "blue")

plasso(0, a, b, c)
qlasso(0.25, a, b, c)
elasso(a, b, c)
vlasso(a, b, c)
mlasso(a, b, c)
MillsRatio(2)

# The Modified_Hans_Gibbs() function uses the Lasso distribution to draw
# samples from the full conditional distribution of the regression coefficients.

y <- 1:20
```

```
X <- matrix(c(1:20,12:31,7:26),20,3,byrow = TRUE)

a1 <- b1 <- u1 <- v1 <- 0.01
sigma2_init <- 1
lambda_init <- 0.1
beta_init <- rep(1, ncol(X))
nsamples <- 1000
verbose <- 100
tune_lambda2 <- TRUE
rao_blackwellization <- FALSE

Output_Hans <- Modified_Hans_Gibbs(
  X, y, beta_init, a1, b1, u1, v1,
  nsamples, lambda_init, sigma2_init,
  verbose, tune_lambda2, rao_blackwellization
)

colMeans(Output_Hans$mBeta)
mean(Output_Hans$vlambda2)

Output_PC <- Modified_PC_Gibbs(
  X, y, a1, b1, u1, v1,
  nsamples, lambda_init, sigma2_init, verbose)

colMeans(Output_PC$mBeta)
mean(Output_PC$vlambda2)
```

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normalize

*Normalize Response and Covariates*

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### Description

This function centers and (optionally) scales the response vector and each column of the design matrix using the population variance. It is used to prepare data for Bayesian Lasso regression.

### Usage

```
normalize(y, X, scale = TRUE)
```

### Arguments

y	A numeric response vector.
X	A numeric matrix or data frame of covariates (design matrix).
scale	Logical; if TRUE, variables are scaled to have unit population variance (default is TRUE).

**Value**

A list with the following elements:

- `vy`: Normalized response vector.
- `mX`: Normalized design matrix.
- `mu.y`: Mean of the response vector.
- `sigma2.y`: Population variance of the response vector.
- `mu.x`: Vector of column means of  $X$ .
- `sigma2.x`: Vector of population variances for columns of  $X$ .

**Examples**

```
set.seed(1)
X <- matrix(rnorm(100 * 10), 100, 10)
beta <- c(2, -3, rep(0, 8))
y <- as.vector(X %*% beta + rnorm(100))
norm_result <- normalize(y, X)
```

# Index

dlasso (LassoDistribution), 2  
elasso (LassoDistribution), 2  
LassoDistribution, 2  
MillsRatio (LassoDistribution), 2  
mlasso (LassoDistribution), 2  
Modified\_Hans\_Gibbs  
    (LassoDistribution), 2  
Modified\_PC\_Gibbs (LassoDistribution), 2  
normalize, 4, 5  
plasso (LassoDistribution), 2  
qlasso (LassoDistribution), 2  
rlasso (LassoDistribution), 2  
vlasso (LassoDistribution), 2  
zlasso (LassoDistribution), 2