

# Package ‘simpleMH’

May 9, 2026

**Title** Simple Metropolis-Hastings MCMC Algorithm

**Version** 0.1.1

**Description** A very bare-bones interface to use the Metropolis-Hastings Monte Carlo Markov Chain algorithm. It is suitable for teaching and testing purposes.

**Imports** mvtnorm

**Suggests** coda, mockery, testthat (>= 3.0.0), knitr, rmarkdown

**License** GPL-3

**Config/testthat/edition** 3

**Encoding** UTF-8

**Language** en-GB

**RoxygenNote** 7.1.1.9001

**VignetteBuilder** knitr

**URL** <https://github.com/Bisaloo/simpleMH>

**BugReports** <https://github.com/Bisaloo/simpleMH/issues>

**NeedsCompilation** no

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**Repository** CRAN

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simpleMH

*Simple Metropolis-Hastings MCMC***Description**

Simple Metropolis-Hastings MCMC

**Usage**

```
simpleMH(f, inits, theta.cov, max.iter, coda = FALSE, ...)
```

**Arguments**

f	function that returns a single scalar value proportional to the log probability density to sample from.
inits	numeric vector with the initial values for the parameters to estimate
theta.cov	covariance matrix of the parameters to estimate.
max.iter	maximum number of function evaluations
coda	logical. Should the samples be returned as <code>coda::mcmc</code> object? (defaults to FALSE)
...	further arguments passed to f

**Value**

- if `coda = FALSE` a list with:
  - *samples*: A two dimensional array of samples with dimensions generation x parameter
  - *log.p*: A numeric vector with the log density evaluate at each generation.
- if `coda = TRUE` a list with:
  - *samples*: A object of class `coda::mcmc` containing all samples.
  - *log.p*: A numeric vector with the log density evaluate at each generation.

**Examples**

```
p.log <- function(x) {
  B <- 0.03
  return(-x[1]^2/200 - 1/2*(x[2]+B*x[1]^2-100*B)^2)
}

simpleMH(p.log, inits=c(0, 0), theta.cov = diag(2), max.iter=3000)
```

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