

# Package: cPseudoMaRg (via r-universe)

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**Type** Package

**Title** Constructs a Correlated Pseudo-Marginal Sampler

**Version** 1.0.1

**Description** The primary function makeCPMSampler() generates a sampler function which performs the correlated pseudo-marginal method of Deligiannidis, Doucet and Pitt (2017) <[arXiv:1511.04992](#)>. If the 'rho=' argument of makeCPMSampler() is set to 0, then the generated sampler function performs the original pseudo-marginal method of Andrieu and Roberts (2009) <[DOI:10.1214/07-AOS574](#)>. The sampler function is constructed with the user's choice of prior, parameter proposal distribution, and the likelihood approximation scheme. Note that this algorithm is not automatically tuned--each one of these arguments must be carefully chosen.

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**RoxygenNote** 7.1.1

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Repository** <https://tbrown122387.r-universe.dev>

**RemoteUrl** <https://github.com/cran/cPseudoMaRg>

**RemoteRef** HEAD

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isBadNum	<i>checks if a log-density evaluation is not a valid number</i>
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**Description**

checks if a log-density evaluation is not a valid number

**Usage**

```
isBadNum(num)
```

**Arguments**

num	evaluation of a log-density
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**Value**

TRUE or FALSE

**Examples**

```
isBadNum(NaN)
```

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makeCPMSampler	<i>correlated pseudo-marginal: generates functions that output a big vector</i>
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**Description**

correlated pseudo-marginal: generates functions that output a big vector

**Usage**

```
makeCPMSampler(
  paramKernSamp,
  logParamKernEval,
  logPriorEval,
  logLikeApproxEval,
  yData,
  numU,
  numIters,
```

```

    rho = 0.99,
    storeEvery = 1,
    nansInLLFatal = TRUE
)

```

## Arguments

paramKernSamp	function(theta) -> theta proposal
logParamKernEval	function(oldTheta, newTheta) -> logDensity.
logPriorEval	function(theta) -> logDensity.
logLikeApproxEval	function(y, thetaProposal, uProposal) -> logApproxDensity.
yData	the observed data
numU	integer number of u samples
numIters	integer number of MCMC iterations
rho	correlation tuning parameter (-1,1)
storeEvery	increase this integer if you want to use thinning
nansInLLFatal	terminate the entire chain on NaNs, or simply disregard sample

## Value

vector of theta samples

## Examples

```

# sim data
realTheta1 <- .2 + .3
realTheta2 <- .2
realParams <- c(realTheta1, realTheta2)
numObs <- 10
realX <- rnorm(numObs, mean = 0, sd = sqrt(realTheta2))
realY <- rnorm(numObs, mean = realX, sd = sqrt(realTheta1 - realTheta2))
# tuning params
numImportanceSamps <- 1000
numMCMCIters <- 1000
randomWalkScale <- 1.5
recordEveryTh <- 1
sampler <- makeCPMSampler(
  paramKernSamp = function(params){
    return(params + rnorm(2)*randomWalkScale)
  },
  logParamKernEval = function(oldTheta, newTheta){
    dnorm(newTheta[1], oldTheta[1], sd = randomWalkScale, log = TRUE)
    + dnorm(newTheta[2], oldTheta[2], sd = randomWalkScale, log = TRUE)
  },
  logPriorEval = function(theta){
    if( (theta[1] > theta[2]) & all(theta > 0)){

```

```

    0
}else{
  -Inf
}
},
logLikeApproxEval = function(y, thetaProposal, uProposal){
  if( (thetaProposal[1] > thetaProposal[2]) & (all(thetaProposal > 0))){
    xSamps <- uProposal*sqrt(thetaProposal[2])
    logCondLikes <- sapply(xSamps,
      function(xsamp) {
        sum(dnorm(y,
          xsamp,
          sqrt(thetaProposal[1] - thetaProposal[2]),
          log = TRUE)) })
    m <- max(logCondLikes)
    log(sum(exp(logCondLikes - m))) + m - log(length(y))
  }else{
    -Inf
  }
},
realY, numImportanceSamps, numMCMCIters, .99, recordEveryTh
)
res <- sampler(realParams)

```

**mean.cpmResults**      *calculates the posterior mean point estimate*

## Description

calculates the posterior mean point estimate

## Usage

```
## S3 method for class 'cpmResults'
mean(x, ...)
```

## Arguments

x	a cpmResults object
...	arguments to be passed to or from methods.

## Value

a vector of parameter estimates (posterior mean)

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`plot.cpmResults`      *plots a cpmResults object*

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**Description**

plots a cpmResults object

**Usage**

```
## S3 method for class 'cpmResults'  
plot(x, ...)
```

**Arguments**

<code>x</code>	a cpmResults object
<code>...</code>	arguments to be passed to or from methods.

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`print.cpmResults`      *prints a cpmResults object*

---

**Description**

prints a cpmResults object

**Usage**

```
## S3 method for class 'cpmResults'  
print(x, ...)
```

**Arguments**

<code>x</code>	a cpmResults object
<code>...</code>	arguments to be passed to or from methods.

**Value**

the same cpmResults object

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