

# Package ‘cPseudoMaRg’

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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) <[doi:10.48550/arXiv.1511.04992](https://doi.org/10.48550/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](https://doi.org/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** CRAN

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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
-----	-----------------------------

**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  
)
```



```

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)

```

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mean.cpmResults	<i>calculates the posterior mean point estimate</i>
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### 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**

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

---

`print.cpmResults`      *prints a cpmResults object*

---

**Description**

prints a cpmResults object

**Usage**

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

**Arguments**

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

**Value**

the same cpmResults object

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