

# Package ‘parSim’

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**Type** Package

**Title** Parallel Simulation Studies

**Version** 0.1.4

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**Description** Perform flexible simulation studies using one or multiple computer cores.  
The package is set up to be usable on high-performance clusters in addition to being run locally, see examples on <<https://github.com/SachaEpskamp/parSim>>.

**Imports** dplyr, parallel, methods, pbapply, snow

**Suggests** ggplot2, tidyr

**License** GPL-2

**NeedsCompilation** no

**Repository** CRAN

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parSim	<i>Parallel Simulator</i>
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### Description

Takes a set of conditions and an R expression and returns a data frame with simulation results. See details.

**Usage**

```
parSim(..., expression, reps = 1, write = FALSE, name, nCores
       = 1, export, exclude, debug = FALSE, progressbar =
       TRUE)
```

**Arguments**

...	Vectors indicating any number of conditions. For example, if you want to vary sample size between $N = 100, 250,$ and $1000,$ and a regression slope between $\beta = 0, 0.5,$ and $1,$ you can assign as first two arguments <code>sampleSize = c(100, 250, 1000), beta = c(0, 0.5, 1).</code>
expression	An R expression that uses the conditions as object names. For example, if the conditions <code>sampleSize = c(100, 250, 1000)</code> is used, then in the R expression you can use the object <code>sampleSize,</code> which may be $100, 250$ or $1000$ depending on the simulation condition.
reps	Number of times each condition has to be replicated.
write	Logical, should the results be written to a file instead of returned as a dataframe?
name	Name of the file if <code>write = TRUE</code>
nCores	Number of cores to use. NOTE: Only setting <code>nCores</code> to $1$ allows you to see a progress bar and to use <code>browser()</code> in the R expression for debugging.
export	A character string of objects to be exported. Only needed if <code>nCores &gt; 1.</code>
exclude	A list with logical calls to exclude cases. Written as formula.
debug	Allows for some debugging controls and output. Not recommended.
progressbar	Logical: should a progress bar be shown. Setting this to <code>FALSE</code> will make simulations much faster!

**Details**

The R expression should use object names assigned as conditions, and should return a list with single values, or a data frame. If you want to output more than one row of results per condition, you may return a data frame with multiple rows. When using multiple cores, note that all packages should be loaded in the R expression, all objects needed should be exported using the `export` object, and you will not see a progress bar.

**Value**

A data frame with every simulation as a row.

**Author(s)**

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**Examples**

```
# Some function we might use:
bias <- function(x,y){abs(x-y)}

# Run the simulation:
Results <- parSim(
  # Any number of conditions:
  sampleSize = c(50, 100, 250),
  beta = c(0, 0.5, 1),
  sigma = c(0.25, 0.5, 1),

  # Number of repetitions?
  reps = 25, # more is better!

  # Parallel?
  nCores = 1,

  # Write to file?
  write = FALSE,

  # Export objects (only needed when nCores > 1):
  export = c("bias"),

  # R expression:
  expression = {
    # Load all R packages in the expression if needed
    # library(...)

    # Want to debug? Enter browser() and run the function. Only works with nCores = 1!
    # browser()

    # Enter whatever codes you want. I can use the conditions as objects.
    X <- rnorm(sampleSize)
    Y <- beta * X + rnorm(sampleSize, sigma)
    fit <- lm(Y ~ X)
    betaEst <- coef(fit)[2]
    Rsquared <- summary(fit)$r.squared

    # Make a data frame with one row to return results (multiple rows is
    # also possible but not recommended):
    data.frame(
      betaEst = betaEst,
      bias = bias(beta,betaEst),
      Rsquared = Rsquared
    )
  }
)

# Analyze the results:
library("ggplot2")
library("tidyr")
```

```
# We want to plot bias and R-squared. Let's first make it long format:
Long <- gather(Results,metric,value,bias:Rsquared)

# Make factors with nice labels:
Long$sigmaFac <- factor(Long$sigma,levels = c(0.25,0.5,1),
  labels = c("Sigma: 0.025", "Sigma: 0.5", "Sigma: 1"))

# Now let's plot:
g <- ggplot(Long, aes(x = factor(sampleSize), y = value, fill = factor(beta))) +
  facet_grid(metric ~ sigmaFac, scales = "free") +
  geom_boxplot() +
  theme_bw() +
  xlab("Sample size") +
  ylab("") +
  scale_fill_discrete("Beta")
print(g)
```

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