

# Package ‘doParallel’

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**Type** Package

**Title** Foreach Parallel Adaptor for the 'parallel' Package

**Version** 1.0.17

**Description** Provides a parallel backend for the `%dopar%` function using the parallel package.

**Depends** R (>= 2.14.0), foreach (>= 1.2.0), iterators (>= 1.0.0), parallel, utils

**Suggests** caret, mlbench, rpart, RUnit

**Enhances** compiler

**License** GPL-2

**URL** <https://github.com/RevolutionAnalytics/doparallel>

**BugReports** <https://github.com/RevolutionAnalytics/doparallel/issues>

**NeedsCompilation** no

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## Contents

doParallel-package . . . . .	2
registerDoParallel . . . . .	2

<b>Index</b>	<b>4</b>
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doParallel-package      *The doParallel Package*

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### Description

The doParallel package provides a parallel backend for the `foreach/%dopar%` function using the `parallel` package of R 2.14.0 and later.

### Details

Further information is available in the following help topics:

`registerDoParallel`    `register doParallel to be used by foreach/%dopar%`

To see a tutorial introduction to the doParallel package, use `vignette("gettingstartedParallel")`.

To see a tutorial introduction to the `foreach` package, use `vignette("foreach")`.

To see a demo of doParallel computing the `sinc` function, use `demo(sincParallel)`.

Some examples (in addition to those in the help pages) are included in the “examples” directory of the doParallel package. To list the files in the examples directory, use `list.files(system.file("examples", package="doParallel"))`. To run the bootstrap example, use `source(system.file("examples", "bootParallel.R", package="doParallel"))`. This is a simple benchmark, executing both sequentially and in parallel. There are many more examples that come with the `foreach` package, which will work with the doParallel package if it is registered as the parallel backend.

For a complete list of functions with individual help pages, use `library(help="doParallel")`.

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`registerDoParallel`      *registerDoParallel*

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### Description

The `registerDoParallel` function is used to register the parallel backend with the `foreach` package.

### Usage

```
registerDoParallel(cl, cores=NULL, ...)  
stopImplicitCluster()
```

## Arguments

<code>cl</code>	A cluster object as returned by <code>makeCluster</code> , or the number of nodes to be created in the cluster. If not specified, on Windows a three worker cluster is created and used.
<code>cores</code>	The number of cores to use for parallel execution. If not specified, the number of cores is set to the value of <code>options("cores")</code> , if specified, or to one-half the number of cores detected by the <code>parallel</code> package.
<code>...</code>	Package options. Currently, only the <code>nocompile</code> option is supported. If <code>nocompile</code> is set to <code>TRUE</code> , compiler support is disabled.

## Details

The `parallel` package from R 2.14.0 and later provides functions for parallel execution of R code on machines with multiple cores or processors or multiple computers. It is essentially a blend of the `snow` and `multicore` packages. By default, the `doParallel` package uses `snow`-like functionality. The `snow`-like functionality should work fine on Unix-like systems, but the `multicore`-like functionality is limited to a single sequential worker on Windows systems. On workstations with multiple cores running Unix-like operating systems, the system `fork` call is used to spawn copies of the current process.

The `doParallel` backend supports both `multicore` and `snow` options passed through the `foreach` function. The supported `multicore` options are `preschedule`, `set.seed`, `silent`, and `cores`, which are analogous to the similarly named arguments to `mclapply`, and are passed using the `.options.multicore` argument to `foreach`. The supported `snow` options are `preschedule`, which like its `multicore` analog can be used to chunk the tasks so that each worker gets a prescheduled chunk of tasks, and `attachExportEnv`, which can be used to attach the export environment in certain cases where R's lexical scoping is unable to find a needed export. The `snow` options are passed to `foreach` using the `.options.snow` argument.

The function `stopImplicitCluster` can be used in vignettes and other places where it is important to explicitly close the implicitly created cluster.

## Examples

```
cl <- makePSOCKcluster(2)
registerDoParallel(cl)
m <- matrix(rnorm(9), 3, 3)
foreach(i=1:nrow(m), .combine=rbind)
stopCluster(cl)
```

# Index

\* **package**

doParallel-package, 2

\* **utilities**

registerDoParallel, 2

doParallel (doParallel-package), 2

doParallel-package, 2

mclapply, 3

registerDoParallel, 2

stopImplicitCluster

(registerDoParallel), 2