# An introduction to parallel computation using foreach package

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## What kind of parallel computation

## Perfectly parallel computation (Embarrassingly parallel computation)

- ► There is little or no effort needed to separate the problem into a number of parallel tasks
- Bootstrapping
- cross-validation
- ▶ grid search, etc

## A basic parctice of parallel computation

#### Hardware requirment

- Single cluster (your own laptop)
- Only requests multiple (cores) CPUs

#### Package requirment

```
library(doParallel)
library(foreach)
library(parallel)
library(iterators)
```

## foreach package

#### An easy and starndard way of parallel comuptation

- Can run a for-loop task as a set of of parallel tasks
- ► Take care of the communication between the tasks (cores)

## Getting start Example

Calculate the sum of the square

$$\sum_{i=1}^{10000} \sum_{j=1}^{i} j^2$$

There is a warning saying the loop ran sequentially To run the loop parallelly, we need to register parallel backends.

```
system.time(foreach(i = 1:10000) %do% sum((1:i)^2))[3]
## elapsed
## 3.789
system.time(foreach(i = 1:10000) %dopar%
         sum((1:i)^2))[3]
## Warning: executing %dopar% sequentially:
## no parallel backend registered
## elapsed
## 4.855
```

#### R foreach

#### Parallel backends

```
registerDoParallel()
```

registerDoParallel() is used to register cores to parallel computation

```
system.time(foreach(i = 1:10000) %do% sum(sqrt(1:i)))[3]
## elapsed
## 3.224
system.time(foreach(i = 1:10000) %dopar%
        sum(sqrt(1:i)))[3]
## elapsed
## 2.43
```

#### R foreach

#### Getting more information of Parallel backends

```
# how many cores are being used
getDoParWorkers()
## [1] 2
# changing back to sequential loop
registerDoSEQ()
getDoParWorkers()
## [1] 1
```

#### R foreach

#### Getting more information of Parallel backends

```
# Check how many cores are avaiable
detectCores()
## [1] 4
# Setting foreach to use 3 cores
registerDoParallel(cores = 3)
getDoParWorkers()
## [1] 3
getDoParName()
## [1] "doParallelMC"
```

 doMC is built on a packaged called multicore (for UNIX-like system), which is used to create a cluster for communication between cores.

## Bootstraping example

```
dim(x)
## [1] 100 2
r <- foreach(i = 1:10000) %dopar% {
    ind <- sample(100, 100, replace = TRUE)</pre>
    result1 <- glm(x[ind, 2] \sim x[ind, 1],
        family = binomial(logit))
    coefficients(result1)
}
r[1]
## [[1]]
## (Intercept) x[ind, 1]
## -11.015560 1.815075
class(r)
## [1] "list"
```

- ► Escaped time for using 3 cores is 14.948 seconds
- ▶ Escaped time for using single core is 27.756 seconds

### .Combine Option

- Specify the way of combining the foreach result
- .combine can also take function like 'c', 'rbind' or even '+'

## Warnings and tips

#### Simple problems may not be benifted by foreach

Communication between cores also takes time

#### Registering too many cores may lead to memory issues

- foreach will copy all the related data for each task
- ▶ Iterator may reduce the usage of memory

#### Use tools to check if foreach runs parallely

- Windows Task Mananger
- Linux top command



Figure 1: Don't waste another second, start parallelizing your computations today

#### Reference

```
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