# Lecture 2: Advanced R

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#### Introduction

In this lecture, we will talk about how to efficiently compute in R.

- R is a vector-operation language. In most case, vectorization speeds up computation.
- When a piece of code is already optimized, we resort to more CPUs for parallel execution.
- Clusters are accessed remotely. Communicating with a remote cluster is different from oprating a local PC. For Windows users, PuTTY and WinSCP help access remote servers.

## **Packages**

The following packages are useful for parallel computing.

```
library(plyr)
library(foreach)
library(doParallel)

## Loading required package: iterators
## Loading required package: parallel
```

#### Speed

Mathematically equivalent though, different calculation methods perform very differently in terms of computational speed.

example: The heteroskedastic-robust variance for the OLS regression is

$$(X'X)^{-1}X'\hat{e}\hat{e}'X(X'X)^{-1}$$

The difficult part is  $X'\hat{e}\hat{e}'X = \sum_{i=1}^{n} \hat{e}_{i}^{2}x_{i}x'_{i}$ . We propose four ways.

- 1. literally do the summation i = 1, ..., n one by one.
- 2.  $X' \operatorname{diag}(\hat{e}^2) X$ , with a dense central matrix.
- 3.  $X' \operatorname{diag}(\hat{e}^2) X$ , with a sparse central matrix.
- 4. Do cross product to X\*ehat. This is different from the matrix formulation. It takes advantage of the element-by-element operation in R.

```
# an example of robust matrix, sparse matrix. Vecteroization.
rm(list = ls())
library(Matrix)

set.seed(111)

# n = 6000; Rep = 10; # 'Matrix' is quick, 'matrix' is slow, adding is OK
n = 50; Rep = 1000; # 'Matrix' is slow, 'matrix' is quick, adding is OK
for (opt in 1:4){
```

```
pts0 = Sys.time()
  for (iter in 1:Rep){
    # set the parameters
    b0 = matrix(c(-1,1), nrow = 2)
    # generate the data
    e = rnorm(n)
    X = cbind( 1, rnorm(n) )
    Y = (X \% *\% b0 + e >=0)
    # note that in this regression b0 is not converge to b0
    # because the model is changed.
    # OLS estimation
    bhat = solve( t(X) \% \% X, t(X)\% \% Y)
    # calculate the t-value
    bhat2 = bhat[2] # parameter we want to test
    e_hat = Y - X %*% bhat
    XXe2 = matrix(0, nrow = 2, ncol = 2)
    if (opt == 1){
      for ( i in 1:n){
       XXe2 = XXe2 + e_{hat[i]^2} * X[i,] %*% t(X[i,])
    } else if (opt == 2) {
      e_hat2_M = matrix(0, nrow = n, ncol = n)
     diag(e_hat2_M) = e_hat^2
     XXe2 = t(X) %*% e_hat2_M %*% X
    } else if (opt == 3) {
      e_hat2_M = Matrix( 0, ncol = n, nrow = n)
      diag(e_hat2_M) = e_hat^2
     XXe2 = t(X) \%*\% e_hat2_M \%*\% X
    } else if (opt == 4) {
     Xe = X * e
     XXe2 = t(Xe) \% \% Xe
   XX_{inv} = solve(t(X) %*% X)
   sig_B = XX_inv %*% XXe2 %*% XX_inv
  cat("n = ", n, ", Rep = ", Rep, ", opt = ", opt, ",
  time = ", Sys.time() - pts0, "\n")
## n = 50 , Rep = 1000 , opt = 1 ,
## time = 1.124578
## n = 50 , Rep = 1000 , opt = 2 ,
## time = 0.2126548
## n = 50 , Rep = 1000 , opt = 3 ,
## time = 1.983931
## n = 50 , Rep = 1000 , opt = 4 ,
```

```
## time = 0.121784
```

# Efficient loop

```
plyr is an R package developed by Hadley Wickham.
example: calculate the empirical coverage probability of a poisson distribution of degree of freedom 2.
CI = function(x){ # construct confidence interval
  # x is a vector of random variables
  n = length(x)
  mu = mean(x)
  sig = sd(x)
  upper = mu + 1.96/sqrt(n) * sig
  lower = mu - 1.96/sqrt(n) * sig
  return( list( lower = lower, upper = upper) )
}
The standard loop
Rep = 10000
sample_size = 1000
# a standard loop
out = rep(0, Rep)
pts0 = Sys.time() # check time
mu = 2
for (i in 1:Rep){
  x = rpois(sample_size, mu)
 bounds = CI(x)
  out[i] = ( ( bounds$lower <= mu  ) & (mu <= bounds$upper) )</pre>
}
cat( "empirical coverage probability = ", mean(out), "\n") # empirical size
## empirical coverage probability = 0.9504
pts1 = Sys.time() - pts0 # check time elapse
print(pts1)
## Time difference of 0.9076478 secs
A plyr loop. It saves book keeping chores, and easier to parallelize.
library(plyr)
capture = function(i){
  x = rpois(sample_size, mu)
  bounds = CI(x)
  return( ( bounds$lower <= mu ) & (mu <= bounds$upper) )</pre>
}
pts0 = Sys.time() # check time
```

```
out = ldply( .data = 1:Rep, .fun = capture )
cat( "empirical coverage probability = ", mean(out$V1), "\n") # empirical size

## empirical coverage probability = 0.9533

pts1 = Sys.time() - pts0 # check time elapse
print(pts1)
```

## Time difference of 1.121264 secs

#### Parallel computing

The basic structure for parallel computing.

In this comparative example, we try

## Time difference of 6.058157 secs

The above block indeed takes more time, because each loop runs very fast.

The code below shows a different story. Each loop takes more time, which dominates the overhead of the CPU communication.

```
Rep = 200
sample_size = 2000000
pts0 = Sys.time() # check time
out = ldply(.data = 1:Rep, .fun = capture, .parallel = FALSE)
cat( "empirical coverage probability = ", mean(out$V1), "\n") # empirical size
pts1 = Sys.time() - pts0 # check time elapse
print(pts1)
```

## **Econ Super**

Try out this script on our econ super computer.

- 1. Log in econsuper;
- 2. Save the code block below as loop\_server.R, and upload it to the server;
- 3. In a terminal, run R --vanilla <loop\_server.R> result\_your\_name.out;
- 4. To run a command in the background, add & at the end of the above command. To keep it running after closing the console, add nohup at the beginning of the command.