## STAT 600: Homework 1

Hyungjoon Kim, Colorado State University

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## Problem 1, 2

I made an R package named SimpLin, and uploaded to my own github repository. You may check the structure and code of the package this link. You can also install that package including the vignette using the following code:

## Problem 3

I compared the performance of lm with SimpLinR using rbenchmark library. Each simulation data has n=100 observations, and the number of simulations is also 100. I used 7 cores for the parallelization, and SimpLinR is almost twice faster than the built-in lm function. Even though the result of lm function contains much more information than SimpLinR, using Rcpp makes a huge improvement of speed.

```
library(SimpLin)
library(foreach)
library(doParallel)
## Loading required package: iterators
## Loading required package: parallel
library(rbenchmark)
n_iter <- 100
n_obs <- 100
n_cores <- detectCores()</pre>
registerDoParallel(n_cores - 1)
benchmark(
  "SimpLinR" = {
    set.seed(600)
    ans <- foreach(i = 1:n_iter, .combine = "c") %dopar% {
      x <- rnorm(n_obs)
      eps <- rnorm(n_obs)</pre>
      y <- 1 - x + eps
```

```
return(as.list(SimpLinR(x, y)))
    }
  },
  "lm" = {
    set.seed(600)
    ans <- foreach(i = 1:n_iter, .combine = "c") %dopar% {</pre>
      x <- rnorm(n obs)
      eps <- rnorm(n_obs)</pre>
      y <- 1 - x + eps
      return(list(lm(y ~ x)))
    }
  },
  replications = 100,
  columns = c("test", "replications", "elapsed",
               "relative", "user.self", "sys.self")
##
         test replications elapsed relative user.self sys.self
## 2
                        100
                               5.069
                                          2.1
                                                   1.069
                                                            1.924
                               2.414
                        100
                                                   0.747
## 1 SimpLinR
                                          1.0
                                                            1.210
stopImplicitCluster()
```

## Problem 4

I attached three tables which summarize the simulation. According to the simulation, the built-in 1m function and SimpLin give the exactly same results. They both have the same estimates, confidence intervals, bias, and so on. However, despite the simulation was implemented using the same-size data, 1m took much more time for the fit.

```
n_iter <- 100
n_obs <- 100
set.seed(600)
elapsed time <- matrix(0, nrow = n iter, ncol = 2)
bias <- matrix(0, nrow = n_iter, ncol = 4)</pre>
coverage <- matrix(0, nrow = n_iter, ncol = 4)</pre>
mse <- matrix(0, nrow = n_iter, ncol = 2)</pre>
pmse <- matrix(0, nrow = n_iter, ncol = 2)</pre>
beta_est <- matrix(0, nrow = n_iter, ncol = 4)</pre>
true_beta <- c(1, -1)
degrees_of_freedom <- n_obs - length(true_beta)</pre>
for(i in 1:n_iter){
  x <- rnorm(n_obs)
  eps <- rnorm(n_obs)</pre>
  y_true <- true_beta[1] + true_beta[2] * x</pre>
  y <- y_true + eps
  t1 <- Sys.time()</pre>
```

```
fit_lm \leftarrow lm(y \sim x)
t2 <- Sys.time()
fit_rcpp <- SimpLinR(x, y)</pre>
t3 <- Sys.time()
elapsed_time[i, ] \leftarrow c(t2 - t1, t3 - t2)
bias[i, ] <- c(true_beta - fit_lm$coefficients,</pre>
                true beta - c(fit rcpp$coefficients))
coverage[i, ] <- c((confint(fit_lm)[1, 1] < true_beta[1]) *</pre>
                       (confint(fit_lm)[1, 2] > true_beta[1]),
                     (confint(fit_lm)[2, 1] < true_beta[2]) *</pre>
                       (confint(fit_lm)[2, 2] > true_beta[2]),
                     (fit_rcpp$confidence.intervals[1, 1] < true_beta[1]) *</pre>
                       (fit_rcpp$confidence.intervals[1, 2] > true_beta[1]),
                     (fit_rcpp$confidence.intervals[2, 1] < true_beta[2]) *</pre>
                       (fit_rcpp$confidence.intervals[2, 2] > true_beta[2]))
mse[i, ] <- c((t(fit_lm$residuals) %*% fit_lm$residuals)[1, 1] /</pre>
                 degrees_of_freedom,
               (t(fit_rcpp$residuals) %*% fit_rcpp$residuals)[1, 1] /
                 degrees_of_freedom)
pmse[i, ] <- c((t(y_true - fit_lm$fitted.values) %*%</pre>
                    (y_true - fit_lm$fitted.values))[1, 1] / n_obs,
                (t(y_true - fit_rcpp\fitted.values) %*%
                   (y_true - fit_rcpp$fitted.values))[1, 1] / n_obs)
```

Table 1: Average Elapsed Time, MSE, and PMSE

|                 | Elapsed Time | MSE      | PMSE     |
|-----------------|--------------|----------|----------|
| lm              | 0.000259     | 1.001616 | 0.019266 |
| ${\tt SimpLin}$ | 0.000012     | 1.001616 | 0.019266 |

Table 2: Average Coverage Rate of 95% Confidence Interval

| $\beta_0$ | $\beta_1$ |
|-----------|-----------|
| 0.97      | 0.97      |
| 0.97      | 0.97      |
|           | 0.97      |

Table 3: Average Bias of the Estimation

|                 | $\beta_0$ | $\beta_1$ |
|-----------------|-----------|-----------|
| lm              | 0.0143687 | -0.00069  |
| ${\tt SimpLin}$ | 0.0143687 | -0.00069  |