Regression Models: Assignment 1

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Importing libraries

```
library(dplyr)
library(MuMIn)
```

Exercise 1

Simulation

```
sim = list()
for (j in 1:1000) {
    vals = c()
    for (i in 1:100) {
        run = 3 + 3*cos(i/10 + 50) + rnorm(1, mean=0, sd=1)
            vals = c(vals, run)
    }
    sim[[j]] = vals
}
sim
```

Exercise 2

Importing the data

```
d <- data.frame(read.table('../data/index.txt', header=TRUE))

X = d$PovPct
Y = d$Brth15to17
beta1 = cov(X, Y)/var(X)
beta0 = mean(Y) - beta1*mean(X)
beta1

## [1] 1.373345
beta0

## [1] 4.267293</pre>
```

Exercise 3

First we have the log-likelihood function for β and σ^2

$$l(\sigma^{2}|X) = \sum_{i=1}^{n} log(\frac{1}{\sqrt{2\pi\sigma^{2}}} - \frac{(Y_{i} - (\beta_{0} + \beta_{1}x_{ik} + \dots + \beta_{k}x_{ik}))^{2}}{2\sigma^{2}})$$

$$\propto -\frac{n}{2}log(\sigma^2) - \frac{(Y-X\beta)\prime(Y-X\beta)}{2\sigma^2}$$

Differentiating the second expression:

$$\frac{\partial l}{\partial \sigma} (-\frac{n}{2} log(\sigma^2) - \frac{(Y - X\beta)\prime(Y - X\beta)}{2\sigma^2} = 0$$

We get:

$$-\frac{n}{2}(\frac{1}{\sigma^2})(2\sigma) - (Y - X\beta)\prime(Y - X\beta) * (-2)(2\sigma^{-3}) = 0$$

We reduce the expression further:

$$-\frac{n}{\sigma} + \frac{(Y - X\beta)\prime(Y - X\beta)}{\sigma^3} = 0$$

We multiply both sides by σ^3 and we get:

$$-n\sigma^2 + (Y - X\beta)\prime(Y - X\beta) = 0$$

And solving for σ^2 we get:

$$\hat{\sigma^2} = \frac{(Y - X\beta)'(Y - X\beta)}{n}$$

Which is our maximum likelihood estimator for σ^2

Exercise 4

Exercise 5

```
bodyfat <- data.frame(read.table('../data/bodyfat.txt', header=TRUE))</pre>
modall <- lm(hwfat ~., data = bodyfat)</pre>
summary(modall)
##
## Call:
## lm(formula = hwfat ~ ., data = bodyfat)
##
## Residuals:
     Min
              1Q Median
                            3Q
                                   Max
## -6.162 -1.858 -0.464 2.502 8.177
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 13.29370
                           9.63027
                                     1.380
                                              0.1718
               -0.32893
                           0.32158 -1.023
                                              0.3098
## age
               -0.06731
                           0.16051
                                    -0.419
## ht
                                              0.6762
                                    -0.527
## wt
               -0.01365
                           0.02591
                                              0.5999
## abs
                0.37142
                           0.08837
                                      4.203 7.55e-05 ***
## triceps
                                              0.0063 **
                0.38743
                           0.13761
                                      2.815
## subscap
                0.11405
                           0.14193
                                     0.804
                                              0.4243
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 3.028 on 71 degrees of freedom
```

```
## Multiple R-squared: 0.8918, Adjusted R-squared: 0.8827
## F-statistic: 97.54 on 6 and 71 DF, p-value: < 2.2e-16
The sum of residuals is zero:
residuals <- sum(resid(modall))
The sum of the observed data is equal to the sum of the fitted values
Y_hat <- predict(modall, bodyfat[1:length(names(bodyfat))-1])
sum(bodyfat$hwfat) - sum(Y_hat)
## [1] 4.547474e-13
The residuals are orthogonal to the predictors
sum(residuals*bodyfat[1:length(names(bodyfat))-1])
## [1] -3.077268e-10
The residuals are orthogonal to the fitted values
sum(residuals*Y_hat)
## [1] -1.568657e-11
Exercise 6
# cols <- names(bodyfat)[1:length(names(bodyfat))-1]</pre>
\# r_2 \leftarrow c()
\# names(r_2) \leftarrow cols
```

```
# rsq \leftarrow function(x,y) cor(x,y)^2
# for (i in 1:length(cols)) {
      modall <- lm(hwfat ~ cols[i], bodyfat)</pre>
      r_2 \leftarrow rsq(predict(hwfat))
# }
# r2
options(na.action = "na.fail")
modall <- lm(hwfat ~., data = bodyfat)
combs <- dredge(modall, extra = "R^2")</pre>
## Fixed term is "(Intercept)"
print("best model")
## [1] "best model"
combs[combs$"R^2" == max(combs<math>$"R^2")]
## Global model call: lm(formula = hwfat ~ ., data = bodyfat)
## ---
## Model selection table
##
      (Intrc)
                 abs
                          age
                                    ht sbscp trcps
                                                             wt
                                                                   R^2 df logLik
        13.29 0.3714 -0.3289 -0.06731 0.1141 0.3874 -0.01365 0.8918 8 -193.43
## 64
       AICc delta weight
## 64 404.9 5.58
## Models ranked by AICc(x)
```