# Topic 2: Exercise 1

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## Importing libraries

```
library(dplyr)
library(Rcpp)
```

# Importing data as described by exercise

```
d <- read.csv("../../datasets/Colleges.csv")</pre>
```

### Replacing binary variable Private with 1 and 0

```
d$Private <- ifelse(d$Private == "Yes", 1, 0)
```

## Selecting columns

```
data <- d %>% dplyr::select('Private','Apps','Accept','Enroll','F.Undergrad')
```

#### Calculating covariances

```
cov_matrix <- cov(data)</pre>
cov_matrix
#>
                   Private
                                   Apps
                                               Accept
                                                           Enroll F. Undergrad
                0.1986559 -745.3552
#> Private
                                            -519.2042 -235.1942
                                                                    -1330.764
            -745.3552439 14978459.5301 8949859.8119 3045255.9876 15289702.474
#> Apps
             -519.2042169 8949859.8119 6007959.6988 2076267.7627 10393582.435
#> Accept
#> Enroll
              -235.1942393 3045255.9876 2076267.7627 863368.3923 4347529.884
#> F.Undergrad -1330.7637175 15289702.4742 10393582.4355 4347529.8841 23526579.326
```

#### Calculating correlations

```
corr_matrix <- cov2cor(cov_matrix)</pre>
corr_matrix
#>
               Private
                           Apps
                                 Accept
                                            Enroll F. Undergrad
          1.0000000 -0.4320947 -0.4752520 -0.5679078 -0.6155605
#> Private
            -0.4320947 1.0000000 0.9434506 0.8468221
                                                    0.8144906
#> Apps
           -0.4752520 0.9434506 1.0000000 0.9116367
                                                    0.8742233
#> Accept
#> Enroll
            0.9646397
#> F.Undergrad -0.6155605 0.8144906 0.8742233 0.9646397
                                                    1.0000000
```

### Experimenting a little bit with the private variable

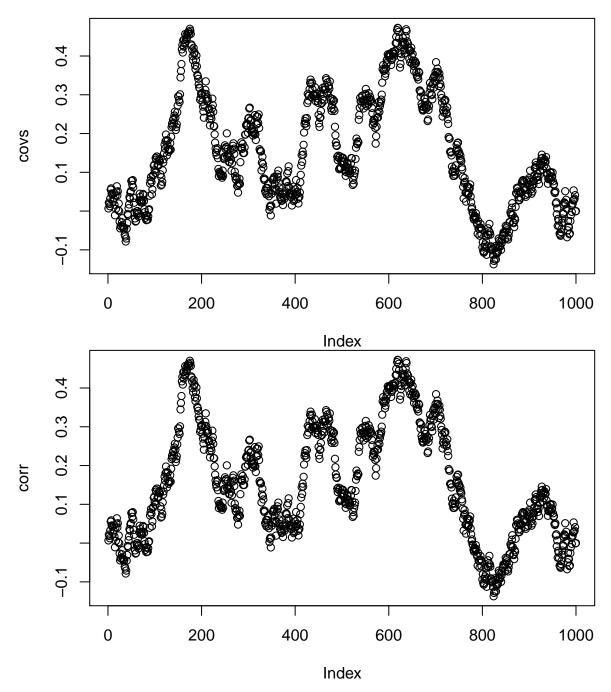
Let's try changing the Yes to 0 and the No to 1 and checking the covariances and correlations

```
d <- read.csv("../../datasets/Colleges.csv")</pre>
d$Private <- ifelse(d$Private == "Yes", 0, 1)</pre>
data <- d %>% dplyr::select('Private','Apps','Accept','Enroll','F.Undergrad')
cov_matrix <- cov(data)</pre>
cov_matrix
#>
                                               Accept
                                                            Enroll F. Undergrad
                   Private
                                    Apps
#> Private
                 0.1986559 7.453552e+02 5.192042e+02
                                                          235.1942
                                                                      1330.764
               745.3552439 1.497846e+07 8.949860e+06 3045255.9876 15289702.474
#> Apps
#> Accept
               519.2042169 8.949860e+06 6.007960e+06 2076267.7627 10393582.435
               235.1942393 3.045256e+06 2.076268e+06 863368.3923 4347529.884
#> Enroll
#> F.Undergrad 1330.7637175 1.528970e+07 1.039358e+07 4347529.8841 23526579.326
corr_matrix <- cov2cor(cov_matrix)</pre>
corr_matrix
#>
                Private
                              Apps
                                     Accept
                                               Enroll F. Undergrad
#> Private 1.0000000 0.4320947 0.4752520 0.5679078 0.6155605
             0.4320947 1.0000000 0.9434506 0.8468221 0.8144906
#> Apps
#> Accept
              0.4752520 0.9434506 1.0000000 0.9116367 0.8742233
#> Enroll
              0.5679078 0.8468221 0.9116367 1.0000000 0.9646397
#> F.Undergrad 0.6155605 0.8144906 0.8742233 0.9646397 1.0000000
```

We get the same numbers with reversed signs.

Let's play with the amount of 1s and 0s in Private and compare it to a simulated variable with only positive values in order to see how the covariance and correlation change and plot it.

```
#include <Rcpp.h>
#include <math.h>
using namespace Rcpp;
// [[Rcpp::export]]
double rcpp_cov(NumericVector v1, NumericVector v2) {
    double vsize = v1.size();
    double cv = 0;
    double v1mean = mean(v1);
    double v2mean = mean(v2);
    double result;
    for (unsigned i=0; i<vsize; i++) {</pre>
        cv = cv + (v1[i] - v1mean)*(v2[i] - v2mean);
    }
    result = cv / (vsize - 1);
    return result;
}
simulate <- function(nrows, simulations, qtvarmin, qtvarmax) {</pre>
    covs <- matrix(rep(0,nrows*simulations), nrow=nrows, byrow=T)</pre>
    corr <- matrix(rep(0,nrows*simulations), nrow=nrows, byrow=T)</pre>
    for (s in 1:simulations) {
        pvtapps <- matrix(rep(0,nrows*2),nrow=nrows,byrow=T)</pre>
        pvtapps[,2] <- runif(nrows, min=qtvarmin, max=qtvarmax)</pre>
        for (i in 1:nrows) {
             pvtapps[,1] \leftarrow c(rep(0,nrows-i), rep(1, i))
             covs[i,s] <- rcpp_cov(pvtapps[,1],pvtapps[,2])</pre>
             corr[i,s] <- rcpp_cov(pvtapps[,1],pvtapps[,2])</pre>
        }
    }
    covs <- rowMeans(covs)</pre>
    corr <- rowMeans(corr)</pre>
    plot(covs)
    plot(corr)
simulate(1000,1000,0,2000)
```



Trying using a variable with both positive and negative values as quantitative variable vs our binary variable simulate(1000,50,-30000,30000)

# What information does the sample covariance provide?

We know that because the Private variable (binary variable) has only 2 possible values, its covariance with other variables is always going to be relatively small and will not provide much information.

What information does the sample correlation provide?