## PCA Exercises

Part 1 - Descriptive statistics and correlation analysis

## Exercises

## -0.01045241

- 1. Prove that cov(x, x) = var(x)
- 2. Prove that cor(x,x)=1
- 3. In a finite sample, what is the difference between: the average difference between an individual and the mean VS the standard deviation (yes it's not the same)?
- 4. Compute the Pearson and Spearman correlation coefficient on the different variables from the iris dataset. Is there any differences? Why?
- 5. With R, create two samples: one where the Pearson coefficient correlation is equal to 1 (but with 2 distinct variables) and one where it's between [-0.1; 0.1]. Use a sample size >100.
- 6. What is the condition on the type of the variables used in order to compute the Pearson (or Spearman) coefficient correlation (it's implicitly said in the course)?
- 7. What are the condition(s) that make possible that cov(x,y) is equal to cor(x,y)?
- 8. Why the Pearson correlation coefficient isn't enough to say that there is a causal link between the variables?
- 9. What happens when doing cor(X), where X is a matrix of quantitative variables?
- 10. With your own words, make a short description of what correlation is.
- 11. Based on the sample below, explain with your own words why 2 variables created the same way are not correlated.

```
x = rnorm(n=1000)
y = rnorm(n=1000)
cor.test(x,y)

##

## Pearson's product-moment correlation
##

## data: x and y
## t = -0.33022, df = 998, p-value = 0.7413
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.07239858 0.05157409
## sample estimates:
## cor
```

12. Create a scatter plot with the sample size in the x-axis and the Pearson correlation coefficient in the y-axis for a range of 10 to 500 for the sample size. For this, you have to use the generate() function below. Do the same thing with the p-value in the y-axis. What do you see? What are the implications?

```
generate = function(sample_size){

#generate sample
x = rnorm(n=sample_size, sd=15)
y = 1.001*x + rnorm(n=sample_size, sd=30)

#compute correlation for the sample
correlation = cor.test(x,y)
p = correlation$p.value
r = correlation$estimate

return(r) #change it to p for the second plot
}
```