Practical session - Modèles de régression linéaire

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IV. Application: GAFAM or BATX dataset

The data set below shows the number of monthly active users (MAU) on Facebook from 2008 to 2021 in millions (source: https://www.statista.com/statistics/264810/number-of-monthly-active-facebook-users-worldwide/). The numbers were taken from Q4 of each year except for the year 2008, whose data is only available in Q3.

```
tab <- read.table("fb_mau.txt", header=TRUE, sep=",")
tab</pre>
```

```
##
      year
            mau
## 1
     2008
            100
      2009
           360
## 3
      2010 608
## 4
      2011 845
## 5
     2012 1056
      2013 1228
## 7
      2014 1393
      2015 1591
## 8
## 9 2016 1860
## 10 2017 2129
## 11 2018 2320
## 12 2019 2498
## 13 2020 2797
## 14 2021 2912
```

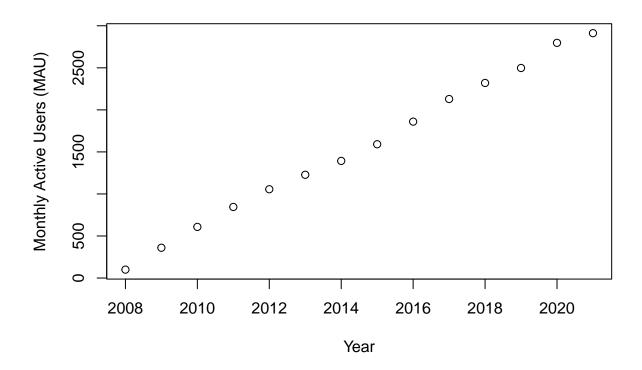
The dimension of the data set:

```
dim(tab)
```

```
## [1] 14 2
```

We then try visualizing the data in order to see if there is an apparent linear relationship between the year and the number of users.

```
plot(tab, xlab="Year", ylab="Monthly Active Users (MAU)")
```



Based on the plotted graph above, we can see that the relationship is fairly linear. Therefore, we can use a linear model to represent the relationship.

```
modreg = lm(mau ~ year, data=tab)
summary(modreg)
```

```
##
## Call:
## lm(formula = mau ~ year, data = tab)
##
##
  Residuals:
##
       Min
                1Q
                    Median
                                3Q
                                       Max
##
   -66.651 -35.664
                    -0.732
                            37.167
                                    60.701
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -4.330e+05
                           5.764e+03
                                      -75.13
                                                <2e-16 ***
                2.157e+02 2.861e+00
                                       75.40
                                               <2e-16 ***
## year
                   0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Signif. codes:
##
## Residual standard error: 43.15 on 12 degrees of freedom
## Multiple R-squared: 0.9979, Adjusted R-squared: 0.9977
## F-statistic: 5685 on 1 and 12 DF, p-value: < 2.2e-16
```

According to the summary of the model, the estimated intercept equals -4.330×10^5 and the estimated

coefficient of the year variable equals 2.157×10^2 . The model can be written in the form:

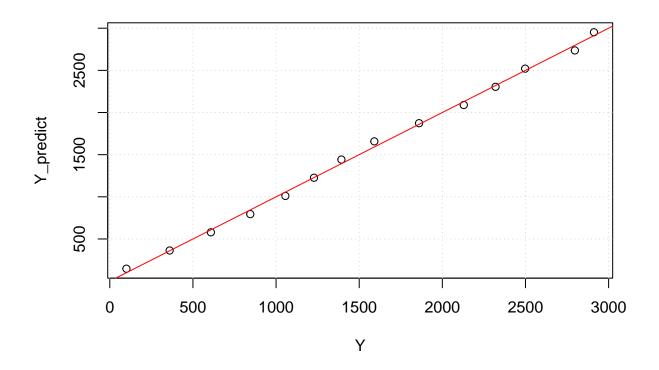
$$\hat{y} = (-4.330 \times 10^5) + (2.157 \times 10^2)x(???? + \hat{\epsilon})$$

where x is the year variable and \hat{y} is the prediction of the MAU.

As for the R^2 , we can see that $R^2 = 0.9979 \approx 1$. It is a great result since the value corresponds to the cosinus of the angle between the vector of predictions and the vector of the target values, and the closer to 0 the angle gets, the better the model becomes.

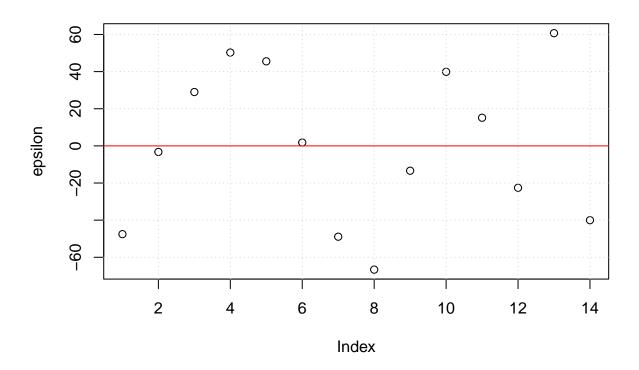
```
Y <- tab$mau
Y_predict <- predict(modreg, tab)

plot(Y, Y_predict)
grid()
abline(a=0, b=1, col="red")</pre>
```



In the graph (y, \hat{y}) above, we can see that the plotted points are fairly close to the bisector, which indicates that the model is acceptable.

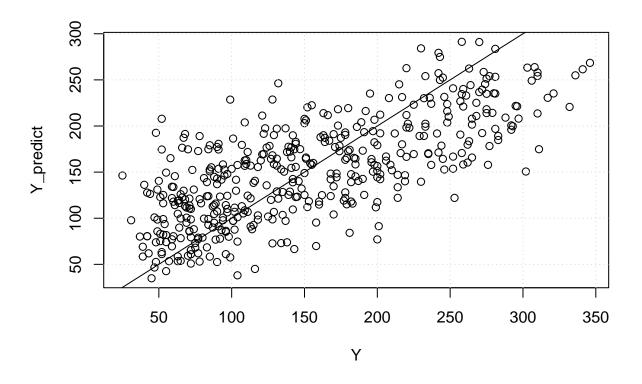
```
epsilon <- Y - Y_predict
plot(epsilon)
grid()
abline(a=0, b=0, col="red")</pre>
```



V. Medical data

```
tab <- read.table("diabetes.txt", header=TRUE, sep="\t")</pre>
modreg = lm(Y~., data=tab)
summary(modreg)
##
## Call:
## lm(formula = Y ~ ., data = tab)
##
## Residuals:
##
        Min
                  1Q
                       Median
                                     ЗQ
                                             Max
## -155.827 -38.536
                       -0.228
                                 37.806 151.353
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -334.56714
                             67.45462 -4.960 1.02e-06 ***
## AGE
                 -0.03636
                              0.21704
                                      -0.168 0.867031
## SEX
                -22.85965
                              5.83582
                                       -3.917 0.000104 ***
## BMI
                  5.60296
                              0.71711
                                        7.813 4.30e-14 ***
## BP
                  1.11681
                              0.22524
                                        4.958 1.02e-06 ***
                 -1.09000
                              0.57333 -1.901 0.057948 .
## S1
```

```
## S2
                  0.74645
                             0.53083
                                       1.406 0.160390
                                       0.475 0.634723
## S3
                  0.37200
                             0.78246
## S4
                  6.53383
                             5.95864
                                        1.097 0.273459
## S5
                 68.48312
                            15.66972
                                        4.370 1.56e-05 ***
## S6
                  0.28012
                             0.27331
                                       1.025 0.305990
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
\#\# Residual standard error: 54.15 on 431 degrees of freedom
## Multiple R-squared: 0.5177, Adjusted R-squared: 0.5066
## F-statistic: 46.27 on 10 and 431 DF, p-value: < 2.2e-16
Y <- tab$Y
Y_predict <- predict(modreg, tab)</pre>
plot(Y, Y_predict)
grid()
abline(a=0, b=1)
```



```
epsilon <- Y - Y_predict
plot(epsilon)
grid()</pre>
```

