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A01630510

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Name of the course: TC2007

Métodos Cuantitativos y Simulación

The design of experiments DOE constructed

Brief description of the experiment, how was physically executed.

- The experiment was executed by teams conformed of 2 people
- First of all, we only cut and fold every single paper that we had. The results were several paper aircrafts with different features.
- We dropped every single aircraft from the 3rd floor of a building.
- Then we measured 3 different times, the total time that lasts the aircraft to fall down from the 3rd floor.
- Then we only get the average from that 3 different measurements.
- Our time is denominated as our dependent variable, so the independent variables are our features over the aircraft.
- The linear regression model is developed in the next following lines, with python
 & R.
- The results that we obtained were the coefficients and the coefficient determinant, as well as our residuals and our predicted values.
- Finally, we plot every single comparison between the average time & residuals, average time & predicted values, every feature & Time predicted.

Compute the linear regression model (y=B0+B1x1+B2x2.....) using both:

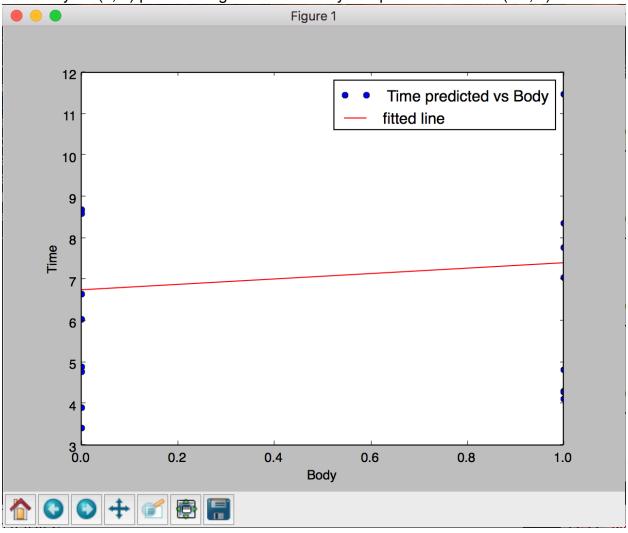
R Code

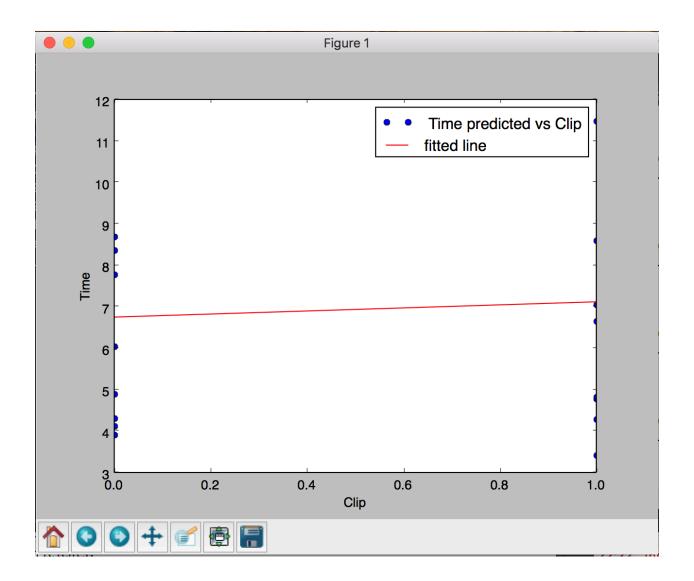
```
setwd("/Users/victormanuel/Downloads") #Set the directory
df <- read.csv("data-set.csv", header = FALSE)</pre>
names(df) <- c('Clip', 'Body', 'Wing', 'Papertype', 'Fly1', 'Fly2',</pre>
'Fly3', 'FlyAvg')
model <- lm(FlyAvg ~ Clip + Body + Wing + Papertype, df)</pre>
model #Define the model & then execute it
Call:
lm(formula = FlyAvg ~ Clip + Body + Wing + Papertype, data = df)
Coefficients:
(Intercept)
                    Clip
                                 Body
                                                      Papertype
                                              Wing
     6.7604
                  0.3708
                               0.6542
                                            1.5792
                                                        -3.7625
summary(model)
Call:
lm(formula = FlyAvg ~ Clip + Body + Wing + Papertype, data = df)
Residuals:
                    Median
     Min
               1Q
                                 3Q
                                         Max
-0.93125 -0.67708 -0.05625 0.33333 2.10208
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
                         0.5099 13.259 4.14e-08 ***
(Intercept)
             6.7604
Clip
              0.3708
                         0.4560
                                  0.813 0.43337
Body
             0.6542
                         0.4560 1.434 0.17926
             1.5792
                                  3.463 0.00531 **
Wing
                         0.4560
                         0.4560 -8.250 4.87e-06 ***
Papertype
             -3.7625
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.9121 on 11 degrees of freedom
Multiple R-squared: 0.8827,
                                Adjusted R-squared:
F-statistic: 20.69 on 4 and 11 DF, p-value: 4.453e-05
df$predicted <- predict(model) #Compare the predicted time vs the</pre>
residuals & plot them
```

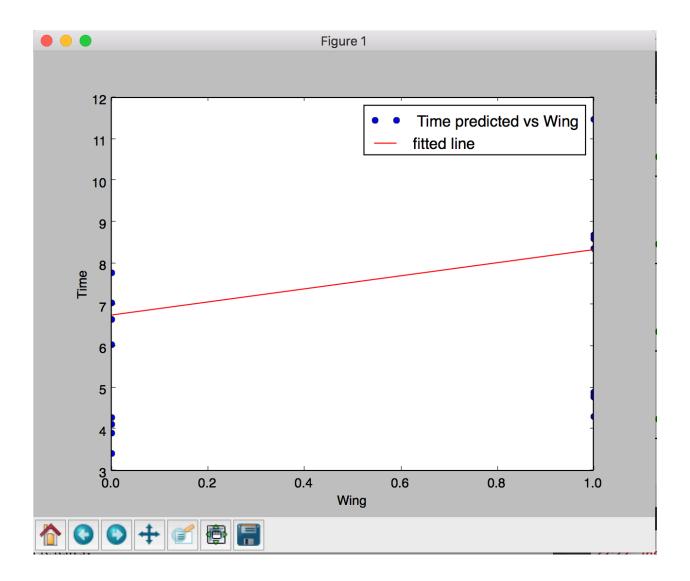
```
df$residuals <- residuals(model)</pre>
install.packages("dplyr")
library(dplyr)
df
plot(df$FlyAvg, df$residuals,ylab="Residuals", xlab="FlyAvg", pch =
16, cex = 1.3, col = "blue", main = "Residuals vs Avg Time")
abline(0,0, col = "red")
plot(df$predicted, df$residuals,ylab="Residuals", xlab="Predicted(Y)",
pch = 16, cex = 1.3, col = "blue", main = "Residuals vs Predicted")
abline(0,0, col = "red")
#Get an histogram of the residuals distribution & then plot it
r <- df$residuals
h <- hist(r, breaks = 10, density = 10, col = "lightgray", xlab =
"Residuals", main = "Histogram of residuals")
 xfit <- seq(min(r), max(r), length = 40)
 yfit <- dnorm(xfit, mean = mean(r), sd = sd(r))</pre>
 yfit <- yfit * diff(h$mids[1:2]) * length(r)</pre>
 lines(xfit, yfit, col = "black", lwd = 2)
 residuals.stdres = rstandard(model)
 qqnorm(residuals.stdres, ylab = "Standardized Residuals", xlab =
"Normal Scores", pch = 16, cex = 1.3, col="blue", main = "" )
 qqline(residuals.stdres)
     Python
import matplotlib.pyplot as plt
import numpy as np
import csv
from sklearn import linear model
# Load dataset, no header
filename = 'data-set.csv'
raw data = open(filename. 'rt')
data = np.loadtxt(raw data, delimiter=",")
# X V
matrix X = data[:,0:4]
matrix Y= data[:,7]
# Create MuLTIPLE linear regression object
modelM = linear model.LinearRegression()
modelM.fit(matrix X,matrix Y)
# The coefficients
print('Coefficients:')
print('B0: \n', modelM.intercept )
print('B1: \n', modelM.coef_[0])
print('B2: \n', modelM.coef_[1])
print('B3: \n', modelM.coef_[2])
```

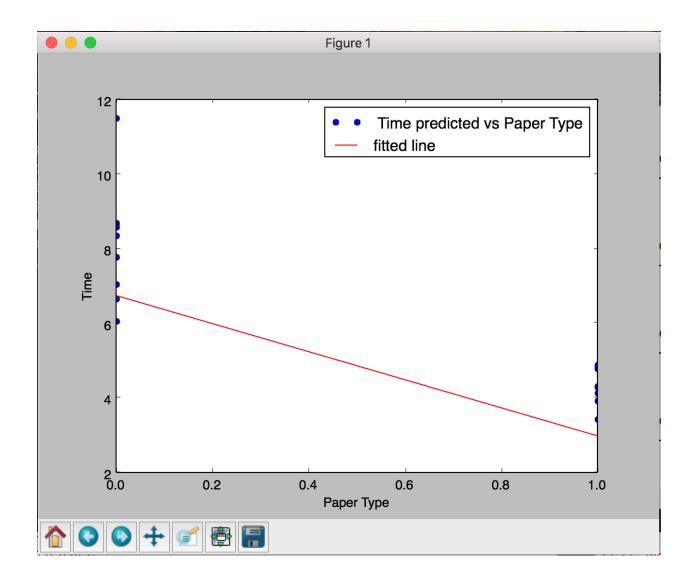
```
print('B4: \n', modelM.coef [3])
# R^2
print('R^2: \n', modelM.score(matrix_X, matrix_Y))
# Different plots of every feature vs the predicted time
plt.plot(data[:,0],matrix_Y, 'o', label = ' Time predicted vs
Clip')
plt.plot(data[:, 0], modelM.intercept +
modelM.coef_[0]*data[:,0], 'r', label = 'fitted line')
plt.xlabel('Clip')
plt.vlabel('Time')
plt.legend()
plt.show()
plt.plot(data[:,1],matrix Y, 'o', label = ' Time predicted vs
Body')
plt.plot(data[:, 1], modelM.intercept_ +
modelM.coef_[1]*data[:,1], 'r', label = 'fitted line')
plt.xlabel('Body')
plt.vlabel('Time')
plt.legend()
plt.show()
plt.plot(data[:,2],matrix_Y, 'o', label = ' Time predicted vs
Wing')
plt.plot(data[:, 2], modelM.intercept_ +
modelM.coef_[2]*data[:,2], 'r', label = 'fitted line')
plt.xlabel('Wing')
plt.ylabel('Time')
plt.legend()
plt.show()
plt.plot(data[:,3],matrix Y, 'o', label = ' Time predicted vs
Paper Type')
plt.plot(data[:, 3], modelM.intercept +
modelM.coef_[3]*data[:,3], 'r', label = 'fitted line')
plt.xlabel('Paper Type')
plt.ylabel('Time')
plt.legend()
plt.show()
```

• In a Layout (2, 2) plot the "Flight time" vs every independent variable (i.e., x)

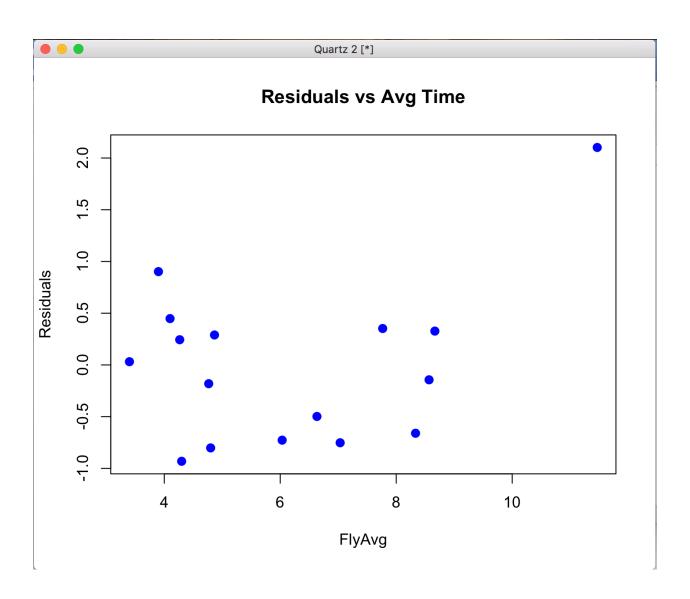






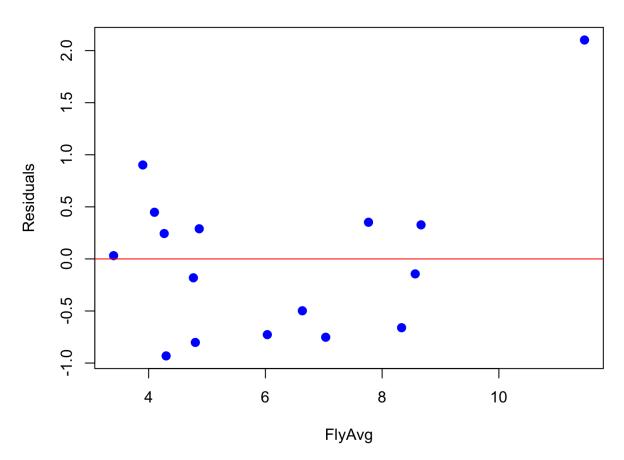


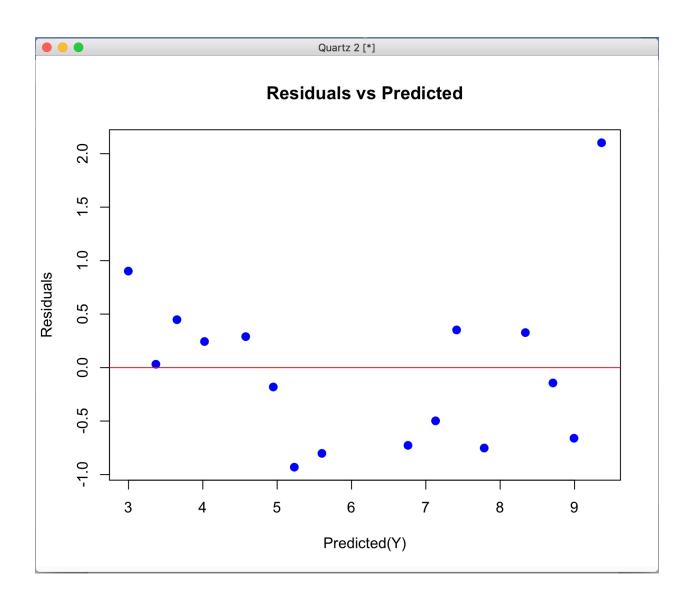
- Add to the data set the "Flight time predicted" and the "Residuals" (as seen in class, MCS_9100_ResidualAnalysisFromLinearRegression.pdf, slide 15)
- Analysis of Residuals as seen in class
 - ∘R²
 - Plot of Residuals vs predicted.

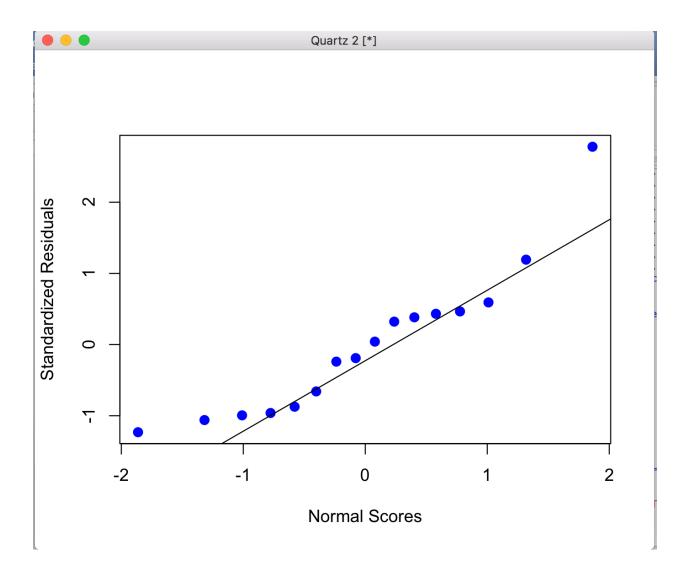


• • Quartz 2 [*]

Residuals vs Avg Time







Explanation of the results of analysis of residuals, and normality of data

What we can say is that our model nearly matches the data, so it's very close to 1 (the coefficient of termination R^2) which means that is not perfect, but is actually a good model. The normality of the data tends to 0 because that's where the actually points (data), trend to be in.

Our residuals (errors) are minimal in our sample, maybe there are 2 or 3 exceptions, but the actual standard is nearly close to 0. So it's good in this case.