

# Homework 2

*Louis Bensard*

*February 22, 2018*

## Problem 1

The method used to sample the points is the Inversion Sampling Method. I used it because I know that the cdf of the exponential density is easily invertible, as shown below:

If  $X \sim \exp(2)$ , then  $F_X(x) = 1 - e^{-x/2} \Rightarrow x = 1 - e^{-F_X^{-1}(x)/2}$ . Thus,

$$F_X^{-1}(x) = -2\ln(1 - x), \quad 0 < x < 1$$

Therefore, if we sample 1000 observations from the *uniform*[0,1] distribution, such that  $U_i \sim \text{unif}[0,1]$  (*iid*), then  $F_X^{-1}(U_i) \sim f_X(x) = \exp(2)$ .

```
n=1000

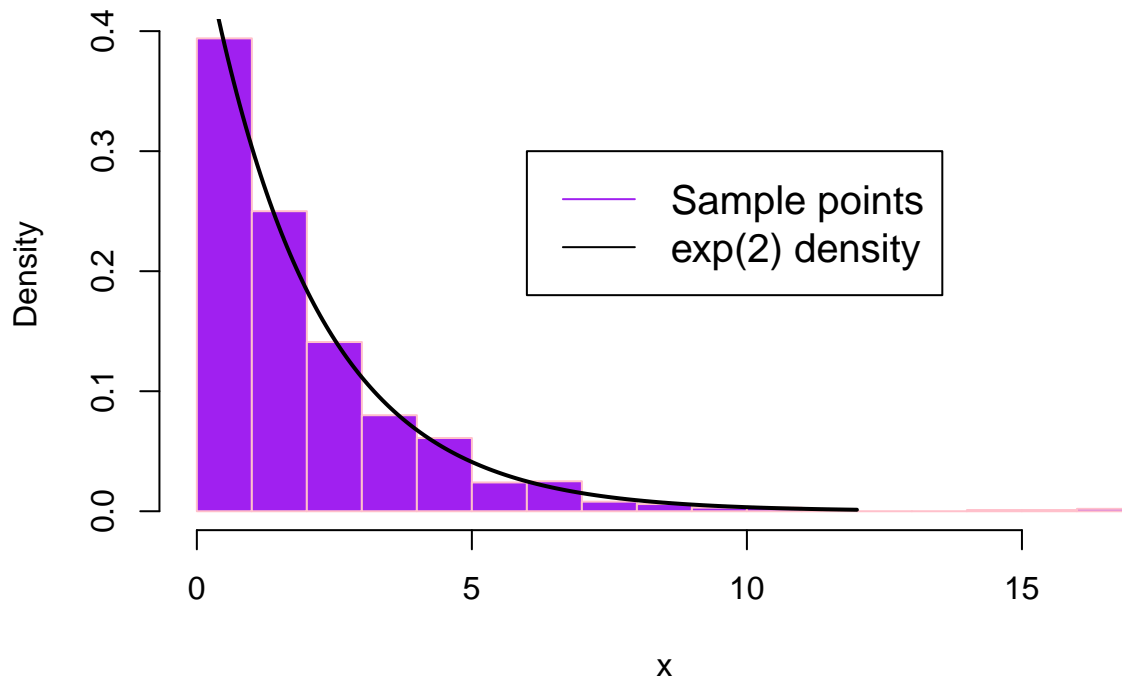
u_vect = runif(n)
F_inv_vect = -2*log(1-u_vect)

x = seq(0,12, length=n)
rexp_vect = rexp(n, rate=1/2)

hist(F_inv_vect, freq=F, col = "purple", border = "pink", breaks = 20,
     main = paste("Histogram of the 1000 points from exp(2) distribution"),
     xlab="x")

lines(x, (1/2)*exp(-x/2), lwd=2)
legend(6,0.3,c("Sample points", "exp(2) density"), col=c("purple", "black"),
      lty=1, cex= 1.25)
```

**Histogram of the 1000 points from exp(2) distribution**



## Problem 2

(i)

PROVE the least squares estimate of  $\beta$  is given by:

$$\hat{\beta} = \frac{\sum_{i=1}^n x_i y_i}{\sum_{i=1}^n x_i^2}$$

We want to find the  $\hat{\beta}$  that minimizes  $SSE = \sum_{i=1}^n \epsilon_i^2 = \sum_{i=1}^n (y_i - \hat{\beta} x_i)^2$ . Now let's set the derivative of SSE with respect to  $\hat{\beta}$  equal to 0, and solve for  $\hat{\beta}$ :

$$\frac{\partial SSE}{\partial \hat{\beta}} = \sum_{i=1}^n (-2x_i)(y_i - \hat{\beta} x_i) = 0$$

$$\Rightarrow \hat{\beta} = \frac{\sum_{i=1}^n x_i y_i}{\sum_{i=1}^n x_i^2}$$

Moreover,  $\frac{\partial^2 SSE}{\partial \hat{\beta}^2} = \sum_{i=1}^n 2x_i^2 > 0$ . Therefore  $\hat{\beta}$  is a minimum.

(ii)

PROVE under the above assumptions that...

(a)

$$E(\hat{\beta}|X) = \beta$$

$$E(\hat{\beta}|X) = E\left(\frac{\sum_{i=1}^n x_i y_i}{\sum_{i=1}^n x_i^2} | X\right) = E((X^T X)^{-1} X^T Y | X) = (x^T x)^{-1} x^T E(Y|X) = (x^T x)^{-1} x^T \beta x = \beta$$

(b)

$$Var(\hat{\beta}|X) = \frac{\sigma_{err}^2}{\sum_{i=1}^n x_i^2}$$

$$\begin{aligned} Var(\hat{\beta}|X) &= [(x^T x)^{-1} x^T] [(x^T x)^{-1} x^T]^T Var(Y|X) \\ &= (x^T x)^{-1} x^T x [(x^T x)^{-1}]^T Var(Y|X) = (x^T x)^{-1} Var(Y|X) \\ &= (x^T x)^{-1} Var(\beta X + \epsilon|X) = (x^T x)^{-1} Var(\epsilon|X) \\ &= (x^T x)^{-1} \sigma_{err}^2 \quad (\text{by assumption}) \\ &= \frac{\sigma_{err}^2}{\sum_{i=1}^n x_i^2} \end{aligned}$$

(c)

$\hat{\beta}|X$  is normal.

$\hat{\beta}|X = (x^T x)^{-1} x^T Y|X$ , therefore  $\hat{\beta}|X$  is a linear combination of  $Y|X = \beta x + \epsilon|X$ . But we know by assumption that  $\epsilon$  is normally distributed, therefore  $\beta x + \epsilon|X = Y|X$  is normally distributed and thus  $\hat{\beta}|X$  is normal.

### Problem 3

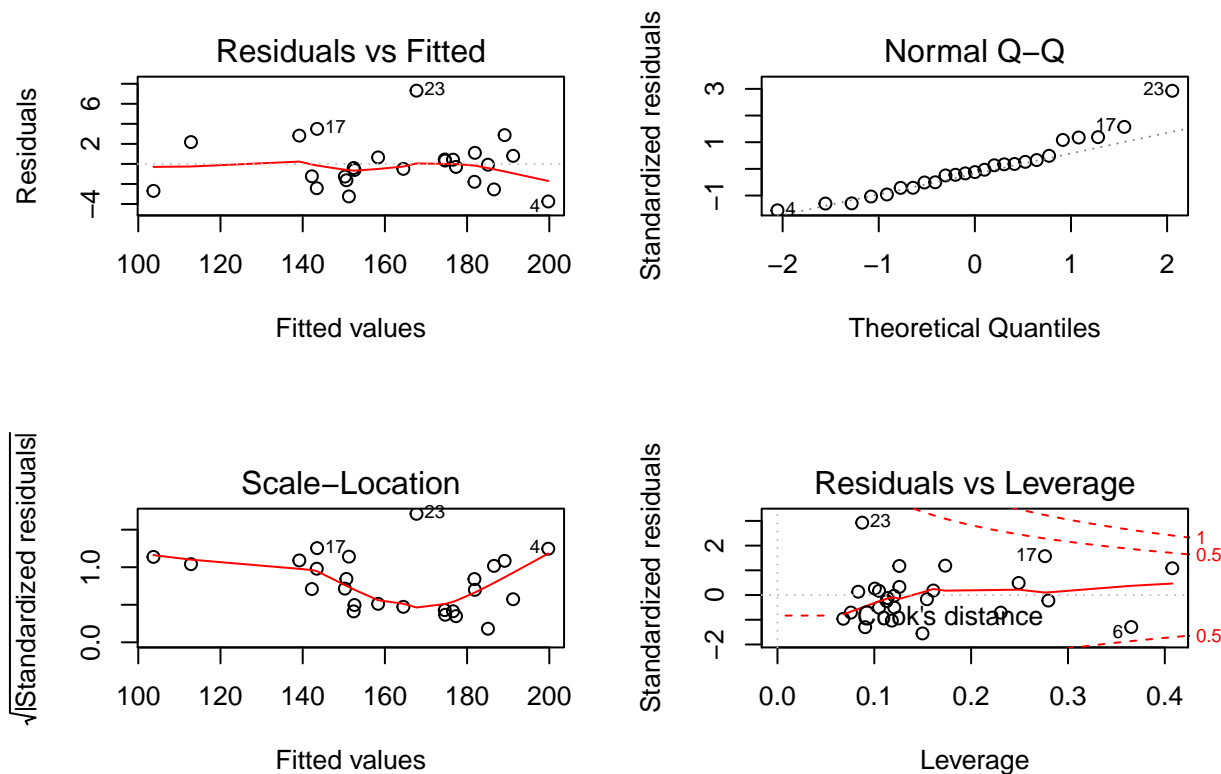
(a)

```
data = read.csv("C:/Users/Louis/Documents/UPMC/M1/Spring 2018/MATH 535/Homework 2/hw2p3.csv")

attach(data)

model1 = lm(FINAL~EXAM1+EXAM2+EXAM3)

par(mfrow=c(2,2))
plot(model1)
```



- The Residual vs Fitted values looks good, even though a small pattern can be observed, therefore the assumption  $E(\hat{\beta}|X) = \beta$  seem valid.
- The Scale-Location graph does not look good, there is a clear pattern observed, therefore the constant variance assumption is invalid.
- The Normal Q-Q plot does not look good as the points are clearly not aligned. Therefore the normality assumption is invalid.
- The assumption of independence is considered always met.

(b)

Since the summary of the model gives us the p-value for  $\beta_{EXAM1} = 0$  by default, then we need to compute it ourselves. We simply need to compute another  $t^*$  adapted to the new problem and then compute the new p-value. We know that for a null hypothesis:  $H_0 : \beta_{EXAM1} = c$ , then:

$$t^* = \frac{\hat{\beta}_{EXAM1} - c}{Std.Error(\hat{\beta}_{EXAM1})}$$

The following code computes this new p-value:

```
#just getting the values we need in the summary table
new_t_star = (coef(summary(model1))[2,1] - 0.2)/coef(summary(model1))[2,2]

new_p_value = 2*pt(abs(new_t_star), length(FINAL)-3-1, lower.tail = FALSE)

print(new_p_value)
```

```
## [1] 0.2129166
```

( $p$  - value > 0.05, then we do not reject the null hypothesis).

(c)

```
predict(model1,newdata=data.frame(EXAM1=70,EXAM2=70,EXAM3=70),
        interval="confidence")
```

```
##          fit          lwr          upr
## 1 140.277 138.7054 141.8485
```

Therefore, a 95% confidence interval for the average of all individuals who score 70's on all three previous exams is [138.7054, 141.8485].

(d)

```
predict(model1,newdata=data.frame(EXAM1=70,EXAM2=70,EXAM3=70),
        interval="prediction")
```

```
##          fit          lwr          upr
## 1 140.277 134.6191 145.9348
```

Therefore, a 95% prediction interval for an individual who scores 70 on their first three exams is [134.6191, 145.9348].