3. (a) Let us assume $\mathbf{y} \sim N(\mathbf{X}\boldsymbol{\beta}, \sigma^2\mathbf{I})$. It is shonw that the maximum likelihood estimator of $\boldsymbol{\beta}$ is $\hat{\boldsymbol{\beta}} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}$. Consider then the estimator $\mathbf{X}\hat{\boldsymbol{\beta}}$. Calculate the expected value $\mathrm{E}(\mathbf{X}\hat{\boldsymbol{\beta}})$ and the covariance matrix $\mathrm{Cov}(\mathbf{X}\hat{\boldsymbol{\beta}})$. What distribution the fitted values $\mathbf{X}\hat{\boldsymbol{\beta}}$ are following?

(2 points)

Solution:

Since $\hat{\beta} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}$ the expected value $\mathbf{E}(\mathbf{X}\hat{\boldsymbol{\beta}})$ is

$$E(\mathbf{X}\hat{\boldsymbol{\beta}}) = E(\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}) = \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}' E(\mathbf{y})$$
$$= \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{X}\boldsymbol{\beta} = \mathbf{X} \cdot \mathbf{I} \cdot \boldsymbol{\beta} = \mathbf{X}\boldsymbol{\beta}.$$

The covariance matrix $\mathrm{Cov}(\mathbf{X}\boldsymbol{\hat{\beta}})$ is

$$Cov(\mathbf{X}\hat{\boldsymbol{\beta}}) = Cov(\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{y}) = \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'Cov(\mathbf{y})(\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}')'$$

$$= \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}' \cdot (\sigma^{2}\mathbf{I}) \cdot \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}' = \sigma^{2}\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'$$

$$= \sigma^{2}\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1} \cdot \mathbf{I} \cdot \mathbf{X}' = \sigma^{2}\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'.$$

Furthermore, since $X\hat{\beta} = Hy$, where $G = X(X'X)^{-1}X'$, is linear transformation of the normally distributed random vector y, it also follows normal distribution $X\hat{\beta} \sim N(X\beta, \sigma^2 X(X'X)^{-1}X')$.

(b) Consider the linear model

$$\mathbf{y} \sim N(\boldsymbol{\mu}, \sigma^2 \mathbf{I}),$$

 $\boldsymbol{\mu} = \mathbf{1}\beta_0,$

where **1** is a vector of ones $\mathbf{1} = (1, 1, \dots, 1)'$. Use the fundamental equation of the BLUE to show that the sample mean

$$\bar{y} = \frac{y_1 + y_2 + \dots + y_n}{n} = \frac{1}{n} \mathbf{1}' \mathbf{y}$$

is the best linear unbiased estimator for the parameter β_0 , i.e., $\hat{\beta}_0 = \bar{y}$. (2 points)

Solution:

In the model ${\bf y}={\bf 1}\beta_0+{m arepsilon}$, the model matrix is ${\bf X}={\bf 1}$ and the matrix ${\bf M}$ is

$$\mathbf{M} = \mathbf{I} - \mathbf{1}(\mathbf{1}'\mathbf{1})^{-1}\mathbf{1}' = \mathbf{I} - \frac{1}{n}\mathbf{1}\mathbf{1}'.$$

With these **X** and **M** matrices, the sample mean $\frac{1}{n}$ **1**'**y** is the BLUE of β_0 if and only if the following BLUE equations are holding:

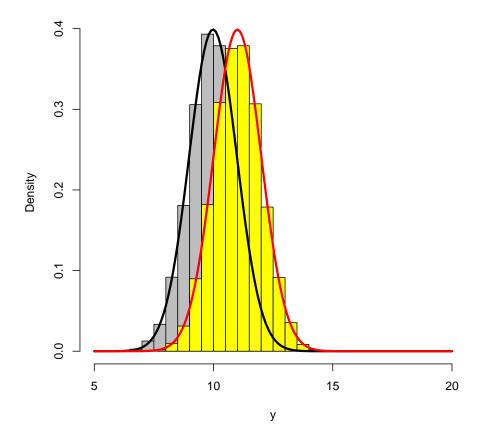
$$\frac{1}{n}\mathbf{1}'(\mathbf{1}:\mathbf{M}) = (1:\mathbf{0}).$$

Now
$$\frac{1}{n}\mathbf{1}'\mathbf{1} = \frac{1}{n} \cdot n = 1$$
 and
$$\frac{1}{n}\mathbf{1}'\mathbf{M} = \frac{1}{n}\mathbf{1}'\left(\mathbf{I} - \frac{1}{n}\mathbf{1}\mathbf{1}'\right) = \frac{1}{n}\mathbf{1}' - \frac{1}{n}\mathbf{1}' \cdot \frac{1}{n}\mathbf{1}\mathbf{1}'$$
$$= \frac{1}{n}\mathbf{1}' - \frac{1}{n^2}\mathbf{1}'\mathbf{1}\mathbf{1}' = \frac{1}{n}\mathbf{1}' - \frac{n}{n^2}\mathbf{1}'$$
$$= \frac{1}{n}\mathbf{1}' - \frac{1}{n}\mathbf{1}' = \mathbf{0}.$$

Thus the fundamental BLUE equations $\frac{1}{n}\mathbf{1}'(\mathbf{1}:\mathbf{M})=(1:\mathbf{0})$ hold and hence the sample mean $\bar{y}=\frac{1}{n}\mathbf{1}'\mathbf{y}$ is the BLUE of β_0 .

(c) Let us consider the large sample situation where we have simulated $n_j = 10000$ observations from the normal distributions $Y_{iA} \sim N(\mu, \sigma^2)$ and $Y_{iB} \sim N(\mu + \delta, \sigma^2)$. The aim is investigate in which values $\delta > 0$ the predictive effect size between Y_{iA} and Y_{iB} is large enough that you would be ready to declare that there is a real effect size between the random variables Y_{iA} and Y_{iB} . Copy the R-code given below. Start changing the value of δ in code and based on the histograms and estimated density curves, decide yourself which values of δ are such that the values of the random variables Y_{iA} and Y_{iB} are "mostly" different. What are your corresponding p-value and d-value when you are feeling that there is a effect size difference?

```
muA<-10
delta<-1
                 # You should change this value
muB<-muA+delta
x<-rep(c("A","B"),each=10000)
yA < -rnorm(10000, mean=muA, sd=1)
yB<-rnorm(10000, mean=muB, sd=1)
y < -c(yA, yB)
model<-lm(y~factor(x))</pre>
betahat<-coef(model)
k1 < -c(0,1)
K<-cbind(k1)
q<-1
Wald < -(t(t(K))*%betahat)%*%solve(t(K))*%vcov(model)%*%K)%*%t(K)%*%betahat)/q
p.value<-pf(Wald, 1, 19998, lower.tail = FALSE)</pre>
p.value
x1 < -cbind(c(1,0))
x2 < -cbind(c(1,1))
pred < -(t(x2)-t(x1))%*%betahat
sigma2<-sigma(model)^2
X<-model.matrix(model)
T<-pred/sqrt(sigma2*(2+(t(x2)-t(x1))%*%solve(t(X)%*%X)%*%(x2-x1)))
d.value<-2*pt(abs(T),df=19998, lower.tail = FALSE)</pre>
d.value
hist(yA, xlim=c(5,20), col="grey", main="", freq=FALSE, xlab="y")
hist(yB, xlim=c(5,20),add=TRUE, col="yellow", freq=FALSE)
lines(seq(5,20,0.1),dnorm(seq(5,20,0.1), mean=betahat[1],sd=sigma(model)),
col="black", lwd=3)
lines(seq(5,20,0.1),dnorm(seq(5,20,0.1), mean=betahat[1]+betahat[2],sd=sigma(model)),
col="red", lwd=3)
```



(2 points)