STAT 420: Exam I Material

Summer 2018, Unger

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When copy and pasting from a code block, or from your local R session, be sure to include all available digits for any numeric answer. It would be best to copy and paste values that were returned using printing methods that do not round results. (Notably the direct output from calling summary().) Also, do not modify the default digits option in the code blocks or your local R session.

All questions are worth 1 point.

Question 1

Variation A

```
# Preamble
```

```
# Starter
set.seed(42)
# Your code here.
# Your code here.
```

After setting the given seed, generate 250 random observations from a normal distribution with a mean of 2 and a variance of 10. What proportion of these observations are larger than 5? (Your answer should be a number between 0 and 1.)

```
# Solution
set.seed(42)
sims = rnorm(n = 250, mean = 2, sd = sqrt(10))
mean(sims > 5)
```

```
## [1] 0.168
```

Variation B

```
# Preamble
```

```
# Starter
set.seed(42)
# Your code here.
# Your code here.
```

After setting the given seed, generate 350 random observations from a normal distribution with a mean of 3 and a variance of 11. What proportion of these observations are larger than 4? (Your answer should be a number between 0 and 1.)

```
# Solution
set.seed(42)
sims = rnorm(n = 350, mean = 3, sd = sqrt(11))
mean(sims > 4)
```

```
## [1] 0.3514286
```

Variation C

```
# Preamble
```

```
# Starter
set.seed(42)
# Your code here.
# Your code here.
```

After setting the given seed, generate 450 random observations from a normal distribution with a mean of 5 and a variance of 13. What proportion of these observations are larger than 6? (Your answer should be a number between 0 and 1.)

```
# Solution
set.seed(42)
sims = rnorm(n = 450, mean = 5, sd = sqrt(13))
mean(sims > 6)
```

```
## [1] 0.36
```

Question 2

Variation A

```
# Preamble
```

```
# Starter
(some fun() + some fun(arg1 = 3, arg2 = 1:10)) / some fun(arg1 = 5, arg2 = 1:5)
```

Write a function named some fun. The function should take two arguments as input:

- arg1, an integer larger than 1, with a default value of 2
- arg2, a vector of real numbers with a default value of 1.

The function should output the average of the elements of arg2 multiplied by arg1. Write and run your function, then report the value from running the starter code.

```
# Solution
some fun = function(arg1 = 2, arg2 = 1) {
 arg1 * mean(arg2)
(some_fun() + some_fun(arg1 = 4, arg2 = 1:10)) / some_fun(arg1 = 5, arg2 = 1:5)
```

```
## [1] 1.6
```

Variation B

```
# Preamble
```

```
# Starter
(some_fun() + some_fun(arg1 = 3, arg2 = 1:10)) / some_fun(arg1 = 5, arg2 = 1:5)
```

Write a function named some fun. The function should take two arguments as input:

- arg1, a vector of real numbers with a default value of 1.
- arg2, an integer larger than 1, with a default value of 2

The function should output the average of the elements of arg1 divided by arg2. Write and run your function, then report the value from running the starter code.

```
# Solution
some_fun = function(arg1 = 1, arg2 = 2) {
  mean(arg1) / arg2
}
(some_fun() + some_fun(arg1 = 1:4, arg2 = 4)) / some_fun(arg1 = 1:9, arg2 = 2)
```

```
## [1] 0.45
```

Question 3

Variation A

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

What proportion of the variance of circumference is explained by a linear relationship with age?

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
summary(orange model)$r.squared
```

```
## [1] 0.8345167
```

Variation B

Preamble

```
# Starter
```

For this question use the iris dataset, which is built into R.

What proportion of the variance of Sepal. Length is explained by a linear relationship with Sepal. Width?

```
# Solution
iris_model = lm(Sepal.Length ~ Sepal.Width, data = iris)
summary(iris_model)$r.squared
```

```
## [1] 0.01382265
```

Variation C

Preamble

```
# Starter
```

For this question use the trees dataset, which is built into R.

What proportion of the variance of Volume is explained by a linear relationship with Height?

```
# Solution
tree_model = lm(Volume ~ Height, data = trees)
summary(tree_model)$r.squared
```

```
## [1] 0.3579026
```

Question 4

Variation A

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create predictions for the circumference of orange trees in millimeters when their age is 400 days and 2500 days. Report the value of the prediction that you feel is more valid.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
# predict(orange_model, newdata = data.frame(age = c(400, 2500)))
# range(Orange$circumference)
predict(orange model, newdata = data.frame(age = 400))
```

```
##
## 60.10778
```

Variation B

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create predictions for the circumference of orange trees in millimeters when their age is 500 days and 2500 days. Report the value of the prediction that you feel is more valid.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
# predict(orange_model, newdata = data.frame(age = c(500, 2500)))
# range(Orange$circumference)
predict(orange_model, newdata = data.frame(age = 500))
```

```
## 70.78481
```

Variation C

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create predictions for the circumference of orange trees in millimeters when their age is 600 days and 2500 days. Report the value of the prediction that you feel is more valid.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
# predict(orange_model, newdata = data.frame(age = c(600, 2500)))
# range(Orange$circumference)
predict(orange_model, newdata = data.frame(age = 600))
```

```
##
          1
## 81.46185
```

Question 5

Variation A

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 90% confidence interval for the change in mean circumference of orange trees in millimeters when age is increased by 1 day. Report the lower bound of this interval.

```
# Solution
orange model = lm(circumference ~ age, data = Orange)
confint(orange_model, parm = "age", level = 0.90)[, 1]
```

```
## [1] 0.0927633
```

Variation B

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 99% confidence interval for the change in mean circumference of orange trees in millimeters when age is increased by 1 day. Report the upper bound of this interval.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
confint(orange_model, parm = "age", level = 0.99)[, 2]
```

```
## [1] 0.1293926
```

Question 6

Variation A

```
# Starter
```

Preamble

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 90% confidence interval for the mean circumference of orange trees in millimeters when the age is 250 days. Report the lower bound of this interval.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
predict(orange_model, interval = "confidence", newdata = data.frame(age = 250), level = 0.90)[,
"lwr"]
```

```
## [1] 32.48418
```

Variation B

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 95% confidence interval for the mean circumference of orange trees in millimeters when the age is 330 days. Report the upper bound of this interval.

```
# Solution
orange model = lm(circumference ~ age, data = Orange)
predict(orange_model, interval = "confidence", newdata = data.frame(age = 330), level = 0.95)[,
```

```
## [1] 65.52032
```

Question 7

Variation A

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 99% prediction interval for an observation of the circumference of an orange tree in millimeters when its age is 400 days. Report the upper bound of this interval.

```
# Solution
orange model = lm(circumference ~ age, data = Orange)
predict(orange_model, interval = "prediction", newdata = data.frame(age = 400), level = 0.99)[,
"upr"]
```

```
## [1] 126.9615
```

Variation B

```
# Preamble
```

```
# Starter
```

For this question use the Orange dataset, which is built into R.

Use a simple linear regression model to create a 95% prediction interval for an observation of the circumference of an orange tree in millimeters when its age is 500 days. Report the lower bound of this interval.

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
predict(orange_model, interval = "prediction", newdata = data.frame(age = 500), level = 0.95)[,
"lwr"]
```

[1] 21.29195

Question 8

Variation A

Preamble

Starter

Consider the SLR model

$$Y = \beta_0 + \beta_1 x + \epsilon$$

where Y is circumference and x is age.

Calculate the p-value of the test $H_0: eta_1 = 0.123 \ \mathrm{vs} \ H_1: eta_1
eq 0.123.$

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
n = length(resid(orange_model))
est = summary(orange_model)$coefficients[2, "Estimate"]
se = summary(orange_model)$coefficients[2, "Std. Error"]
t = (est - 0.123) / se
2 * pt(abs(t), df = n - 2, lower.tail = FALSE)
```

[1] 0.05837492

Variation B

Preamble

Starter

Consider the SLR model

$$Y = \beta_0 + \beta_1 x + \epsilon$$

where Y is circumference and x is age.

Calculate the p-value of the test $H_0: eta_1 = 0.125 \ \mathrm{vs} \ H_1: eta_1
eq 0.125.$

```
# Solution
orange model = lm(circumference ~ age, data = Orange)
n = length(resid(orange_model))
est = summary(orange_model)$coefficients[2, "Estimate"]
se = summary(orange_model)$coefficients[2, "Std. Error"]
t = (est - 0.125) / se
2 * pt(abs(t), df = n - 2, lower.tail = FALSE)
```

[1] 0.03472037

Variation C

Preamble

Starter

Consider the SLR model

$$Y = \beta_0 + \beta_1 x + \epsilon$$

where Y is circumference and x is age.

Calculate the p-value of the test $H_0: eta_1 = 0.127 ext{ vs } H_1: eta_1
eq 0.127.$

```
# Solution
orange_model = lm(circumference ~ age, data = Orange)
n = length(resid(orange_model))
est = summary(orange model)$coefficients[2, "Estimate"]
se = summary(orange_model)$coefficients[2, "Std. Error"]
t = (est - 0.127) / se
2 * pt(abs(t), df = n - 2, lower.tail = FALSE)
```

[1] 0.02002805

Question 9

Variation A

Preamble

Starter

Consider the true model

$$Y = 2 + 1.5x_1 - 2.1x_2 + 3.2x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=4)$$

What is the probability that Y is less than 3 given that:

- $x_1 = 0$
- $x_2 = 0$
- $x_3 = 0$

```
# Solution
mu = 2 + 1.5 * 0 - 2.1 * 0 + 3.2 * 0
pnorm(3, mean = mu, sd = 2)
```

[1] 0.6914625

Variation B

Preamble

Starter

Consider the true model

$$Y = 3 + 1.4x_1 - 2.3x_2 + 3.1x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=9)$$

What is the probability that Y is less than 5 given that:

- $x_1 = 0$
- $x_2 = 0$
- $x_3 = 0$

```
# Solution
mu = 3 + 1.4 * 0 - 2.3 * 0 + 3.1 * 0
pnorm(5, mean = mu, sd = 3)
```

[1] 0.7475075

Variation C

Preamble

Starter

Consider the true model

$$Y = 4 + 1.3x_1 - 2.4x_2 + 3.7x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=25)$$

What is the probability that Y is less than 3 given that:

- $x_1 = 0$
- $x_2 = 0$
- $x_3 = 0$

```
# Solution
mu = 4 + 1.3 * 0 - 2.4 * 0 + 3.7 * 0
pnorm(3, mean = mu, sd = 5)
```

[1] 0.4207403

Question 10

Variation A

Preamble

Starter

Consider the true model

$$Y = 2 + 1.5x_1 - 2.1x_2 + 3.2x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=4)$$

What is the probability that Y is greater than 11 given that:

- $x_1 = 1$
- $x_2 = 2$
- $x_3 = 3$

```
# Solution
mu = 2 + 1.5 * 1 - 2.1 * 2 + 3.2 * 3
pnorm(11, mean = mu, sd = 2, lower.tail = FALSE)
```

[1] 0.1468591

Variation B

Preamble

Starter

Consider the true model

$$Y = 3 + 1.4x_1 - 2.3x_2 + 3.1x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=9)$$

What is the probability that Y is greater than 11 given that:

- $x_1 = 1$
- $x_2 = 2$
- $x_3 = 3$

```
# Solution
mu = 3 + 1.4 * 1 - 2.3 * 2 + 3.1 * 3
pnorm(12, mean = mu, sd = 3, lower.tail = FALSE)
```

[1] 0.1668553

Variation C

Preamble

Starter

Consider the true model

$$Y = 4 + 1.3x_1 - 2.4x_2 + 3.7x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=25)$$

What is the probability that Y is greater than 13 given that:

- $x_1 = 1$
- $x_2 = 2$
- $x_3 = 3$

```
# Solution
mu = 4 + 1.3 * 1 - 2.4 * 2 + 3.7 * 3
pnorm(13, mean = mu, sd = 5, lower.tail = FALSE)
```

```
## [1] 0.3897388
```

Question 11

Variation A

```
# Preamble
```

```
# Starter
set.seed(420)
x_values = data.frame(
  x1 = rnorm(10),
  x2 = runif(10),
 x3 = runif(10),
  x4 = runif(10)
```

Consider the true model

$$Y = 4 + 2.5x_1 + 3x_2 - 5x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=9)$$

What is $SD[\hat{\beta}_2]$ given the values of predictors above?

```
# Solution
X = as.matrix(cbind(rep(1, 10), x_values))
C = solve(t(X) %*% X)
sqrt(9 * C[2 + 1, 2 + 1])
```

```
## [1] 4.45513
```

Variation B

```
# Preamble
```

```
# Starter
set.seed(420)
x_values = data.frame(
  x1 = rnorm(15),
  x2 = runif(15),
  x3 = runif(15),
  x4 = runif(15)
```

Consider the true model

$$Y = 5 + 1.5x_1 + 2x_2 - 3x_3 + \epsilon$$

where

$$\epsilon \sim N(0,\sigma^2=25)$$

What is $SD[\hat{\beta}_1]$ given the values of predictors above?

```
# Solution
X = as.matrix(cbind(rep(1, 15), x_values))
C = solve(t(X) %*% X)
sqrt(25 * C[1 + 1, 1 + 1])
```

```
## [1] 1.679908
```

Question 12

Variation A

```
# Preamble
```

```
# Starter
```

Calculate the critical value used for a 90% confidence interval about a single eta parameter of a multiple linear regression model with 4 predictors that is fit to 15 observations. (Your answer should be a positive value.)

```
# Solution
conf level = 0.90
sig_level = 1 - conf_level
n = 15
p = 4 + 1
abs(qt(sig_level / 2, df = n - p))
```

```
## [1] 1.812461
```

Variation B

```
# Preamble
```

```
# Starter
```

Calculate the critical value used for a 95% confidence interval about a single eta parameter of a multiple linear regression model with 5 predictors that is fit to 16 observations. (Your answer should be a positive value.)

```
# Solution
conf level = 0.95
sig_level = 1 - conf_level
n = 16
p = 5 + 1
abs(qt(sig_level / 2, df = n - p))
```

```
## [1] 2.228139
```

Variation C

```
# Preamble
```

```
# Starter
```

Calculate the critical value used for a 99% confidence interval about a single eta parameter of a multiple linear regression model with 6 predictors that is fit to 17 observations. (Your answer should be a positive value.)

```
# Solution
conf level = 0.99
sig_level = 1 - conf_level
n = 17
p = 6 + 1
abs(qt(sig_level / 2, df = n - p))
```

```
## [1] 3.169273
```

Question 13

Variation A

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

What proportion of the median home value (medv) is explained by a linear relationship with the average number of rooms per dwelling and the per capita crime rate?

```
# Solution
library(MASS)
boston_mod = lm(medv ~ rm + crim, data = Boston)
summary(boston_mod)$r.squared
```

```
## [1] 0.5419592
```

Variation B

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

What proportion of the median home value (medv) is explained by a linear relationship with the average number of rooms per dwelling and the nitrogen oxide concentration in parts per 10 million?

```
# Solution
library(MASS)
boston_mod = lm(medv ~ rm + nox, data = Boston)
summary(boston_mod)$r.squared
```

```
## [1] 0.535438
```

Variation C

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

What proportion of the median home value (medv) is explained by a linear relationship with the average number of rooms per dwelling and the full-value property-tax rate per \$10,000?

```
# Solution
library(MASS)
boston_mod = lm(medv ~ rm + tax, data = Boston)
summary(boston_mod)$r.squared
```

```
## [1] 0.5605639
```

Question 14

Variation A

Preamble

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

Consider the multiple regression model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{rm}} x_{\texttt{rm}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

with the usual assumptions on the error term and Y is $\ensuremath{\mathsf{medv}}$.

Report the p-value for testing $H_0: \beta_{\mathtt{crim}} = 0 \text{ vs } H_1: \beta_{\mathtt{crim}} \neq 0.$

```
# Solution
library(MASS)
boston_mod = lm(medv ~ lstat + rm + crim + tax + nox, data = Boston)
summary(boston_mod)$coefficients["crim", "Pr(>|t|)"]
```

```
## [1] 0.09102573
```

Variation B

Preamble

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

Consider the multiple regression model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{rm}} x_{\texttt{rm}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

with the usual assumptions on the error term and Y is medv.

Report the p-value for testing $H_0: \beta_{\mathtt{tax}} = 0 \text{ vs } H_1: \beta_{\mathtt{tax}} \neq 0.$

```
# Solution
library(MASS)
boston_mod = lm(medv ~ lstat + rm + crim + tax + nox, data = Boston)
summary(boston mod)$coefficients["tax", "Pr(>|t|)"]
```

```
## [1] 0.003344395
```

Variation C

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package.

Consider the multiple regression model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{rm}} x_{\texttt{rm}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

with the usual assumptions on the error term and Y is $\ensuremath{\mathsf{medv}}$.

Report the p-value for testing $H_0: \beta_{nox} = 0 \text{ vs } H_1: \beta_{nox} \neq 0.$

```
# Solution
library(MASS)
boston mod = lm(medv \sim lstat + rm + crim + tax + nox, data = Boston)
summary(boston_mod)$coefficients["nox", "Pr(>|t|)"]
```

```
## [1] 0.1724987
```

Question 15

Variation A

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package. Use medv as the response variable.

Compare the model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{rm}} x_{\texttt{rm}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

to the model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{rm}} x_{\texttt{rm}} + \epsilon$$

using an F test with an α of 0.05. Report the RSS of the model you prefer.

```
# Solution
library(MASS)
full mod = lm(medv ~ lstat + rm + crim + tax + nox, data = Boston)
null_mod = lm(medv ~ lstat + rm, data = Boston)
# anova(null_mod, full_mod)
sum(resid(full_mod) ^ 2)
```

[1] 14869.32

Variation B

```
# Preamble
```

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package. Use medv as the response variable.

Compare the model

$$Y = \beta_0 + \beta_{\texttt{rm}} x_{\texttt{rm}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

to the model

$$Y = eta_0 + eta_{ exttt{rm}} x_{ exttt{rm}} + eta_{ exttt{nox}} x_{ exttt{nox}} + \epsilon$$

using an F test with an α of 0.05. Report the RSS of the model you prefer.

```
# Solution
library(MASS)
full_mod = lm(medv ~ rm + crim + tax + nox, data = Boston)
null\ mod = lm(medv \sim rm + nox, data = Boston)
# anova(null_mod, full_mod)
sum(resid(full mod) ^ 2)
```

```
## [1] 18134.23
```

Variation C

Preamble

```
# Starter
library(MASS)
```

For this question use the Boston dataset from the MASS package. Use medv as the response variable.

Compare the model

$$Y = \beta_0 + \beta_{\texttt{lstat}} x_{\texttt{lstat}} + \beta_{\texttt{crim}} x_{\texttt{crim}} + \beta_{\texttt{tax}} x_{\texttt{tax}} + \beta_{\texttt{nox}} x_{\texttt{nox}} + \epsilon$$

to the model

$$Y = eta_0 + eta_{ exttt{lstat}} x_{ exttt{lstat}} + eta_{ exttt{tax}} x_{ exttt{tax}} + \epsilon$$

using an F test with an α of 0.05. Report the RSS of the model you prefer.

```
# Solution
library(MASS)
full_mod = lm(medv ~ lstat + crim + tax + nox, data = Boston)
null_mod = lm(medv ~ lstat + tax, data = Boston)
# anova(null mod, full mod)
sum(resid(null_mod) ^ 2)
```

```
## [1] 19197.98
```