Week 3 - Homework

STAT 420, Summer 2018, Panda5

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Exercise 1 (Using lm for Inference)

For this exercise we will use the cats dataset from the MASS package. You should use ?cats to learn about the background of this dataset.

(a) Fit the following simple linear regression model in $\,\mathbb{R}$. Use heart weight as the response and body weight as the predictor.

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$

Store the results in a variable called cat_{model} . Use a t test to test the significance of the regression. Report the following:

```
library(MASS)
cat_model = lm(Hwt~Bwt,data=cats)
summary(cat_model)$coefficient[,"t value"]
```

```
## (Intercept) Bwt
## -0.5152019 16.1193908
```

```
summary(cat_model)$coefficient[,"Pr(>|t|)"]
```

```
## (Intercept) Bwt
## 6.072131e-01 6.969045e-34
```

```
alpha = 0.05
level = 1- alpha
confint(cat_model,level=level)
```

```
## 2.5 % 97.5 %

## (Intercept) -1.725163 1.011838

## Bwt 3.539343 4.528782
```

```
##
        5 %
                95 %
## 3.619716 4.448409
confint(cat_model,interval="confidence",level=0.99)["(Intercept)",]
##
       0.5 %
                99.5 %
## -2.164125 1.450800
body_weight1 = data.frame(Bwt=c(2.1,2.8))
predict_newdata1=predict(cat_model,newdata=body_weight1,interval="confidence",leve
1=0.99)
body weight2 = data.frame(Bwt=c(2.8,4.2))
predict_newdata2=predict(cat_model,newdata=body_weight2,interval="prediction",leve
1=0.99)
beta_0_hat = summary(cat_model)$coefficient[1,1]
beta_1_hat = summary(cat_model)$coefficient[2,1]
beta_0_hat_se = summary(cat_model)$coefficient[1,2]
beta_1_hat_se = summary(cat_model)$coefficient[2,2]
beta_0_hat_t = (beta_0_hat - 4) / beta_0_hat_se
beta_1_hat_t = (beta_1_hat - 4) / beta_1_hat_se
beta_0_hat_p = 2* pt(abs(beta_0_hat_t),nrow(cars)-2,lower.tail = FALSE)
beta_1_hat_p = 2* pt(abs(beta_1_hat_t),nrow(cars)-2,lower.tail = FALSE)
#Creating Confidence and Prediction Bands
bwt_grid = seq(min(cats$Bwt), max(cats$Bwt), by = 0.01)
hwt ci band = predict(cat model,
                      newdata = data.frame(Bwt = bwt grid),
                      interval = "confidence",
                      level = 0.90)
hwt_pi_band = predict(cat_model,
                      newdata = data.frame(Bwt = bwt_grid),
                       interval = "prediction",
                      level = 0.90)

    The null and alternative hypotheses

      a. Null Hypothesis H_0: \beta_1 = 0
```

confint(cat_model,interval="confidence",level=0.90)["Bwt",]

b. Alternate Hypothesis $H_1:\beta_1\neq 0$

t statistics for β_0 : **-0.5152019** and β_1 : **16.1193908**

The value of the test statistic:

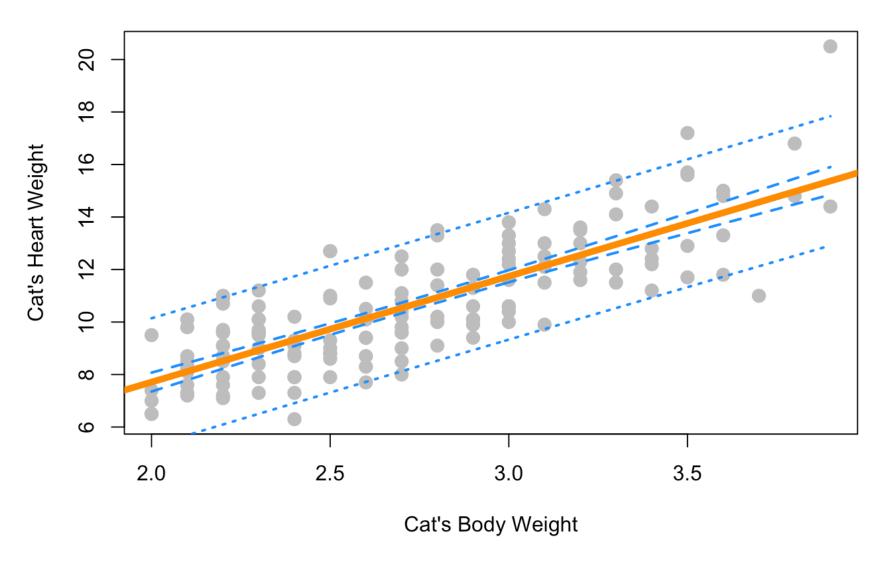
- The p-value of the test : p value for β_0 :0.6072131 and for β_1 : 6.969044610^{-34}
- A statistical decision at $\alpha = 0.05$

The P value for β_1 which is: **6.969044610^{-34}** < **0.05** (α). Since P is smaller than α , we **reject the null Hypothesis**

- A conclusion in the context of the problem:
 - Since we **reject the null hypothesis**, hence β_1 is non zero, so there is **significant linear relationship** Cat's Body Weight and Cat's Heart Weight
- **(b)** Calculate a 90% confidence interval for β_1 . Give an interpretation of the interval in the context of the problem.
- 90% confident interval for β_1 is between **3.619716** and **4.4484094**
- **Interpretation:** We are 90% confident that true change in mean of Cat's Heart Weight for an increase Cat's Body Weight by 1 kg is between **3.619716** and **4.4484094**
- (c) Calculate a 99% confidence interval for β_0 . Give an interpretation of the interval in the context of the problem.
- 99% confidence interval for eta_0 is between **-2.1641253** and **1.4508004**
- **Interpretation:** We are 99% confident that the true change in mean of Cat's Heart Weight for zero Cat's Body is between **-1.5028345** and **0.7895096**
- (d) Use a 99% confidence interval to estimate the mean heart weight for body weights of 2.1 and 2.8 kilograms. Which of the two intervals is wider? Why?
- 99% confidence interval of mean Body weight of 2.1 is between 7.5992252 and 8.6305133
- 99% confidence interval of mean Body weight of 2.8 is between 10.6187959 and 11.2586303
- **2.1** have wider interval than that of **2.8**, becase the range of **2.1** is **1.0312881** > **0.6398344** of **2.8**
- (e) Use a 99% prediction interval to predict the heart weight for body weights of 2.8 and 4.2 kilograms.
- 99% confidence interval of mean Body weight of 2.8 is between lower:7.1332472 and 14.7441791
- 99% confidence interval of mean Body weight of 4.2 is between lower:12.6608829 and 20.5119189
- **(f)** Create a scatterplot of the data. Add the regression line, 90% confidence bands, and 90% prediction bands.

```
plot(Hwt~Bwt,
    data=cats,
    xlab = "Cat's Body Weight",
    ylab = "Cat's Heart Weight",
    main = "Scatter Plot Cat's Body Weight Vs Cat's Heart Weight",
    col = "grey",
    pch = 20,
    cex = 2
    )
abline(cat_model,lwd = 5, col = "darkorange")
lines(bwt_grid,hwt_ci_band[,"lwr"],col="dodgerblue",lwd=2,lty=2)
lines(bwt_grid,hwt_ci_band[,"upr"],col="dodgerblue",lwd=2,lty=2)
lines(bwt_grid,hwt_pi_band[,"lwr"],col="dodgerblue",lwd=2,lty=3)
lines(bwt_grid,hwt_pi_band[,"upr"],col="dodgerblue",lwd=2,lty=3)
```

Scatter Plot Cat's Body Weight Vs Cat's Heart Weight



(g) Use a *t* test to test:

• $H_0: \beta_1 = 4$

• $H_1: \beta_1 \neq 4$

Report the following:

• The value of the test statistic:

test statistics for β_1 : **0.1361084**

• The p-value of the test:

The p-value for β_1 :**0.8923048**

• A statistical decision at $\alpha = 0.05$

Since the p-value of the test for β_1 is : **0.8923048** > 0.05 (= α), hence we **failed to reject the Hypothesis** of β_1 = 4, where the slope (β_1) is close to the value of 4

Exercise 2 (More 1m for Inference)

For this exercise we will use the Ozone dataset from the mlbench package. You should use ?Ozone to learn about the background of this dataset. You may need to install the mlbench package. If you do so, do not include code to install the package in your R Markdown document.

For simplicity, we will re-perform the data cleaning done in the previous homework.

```
data(Ozone, package = "mlbench")
Ozone = Ozone[, c(4, 6, 7, 8)]
colnames(Ozone) = c("ozone", "wind", "humidity", "temp")
Ozone = Ozone[complete.cases(Ozone), ]

ozone_wind_model = lm(ozone~wind,data=Ozone)
ozone_temp_model = lm(ozone~temp,data=Ozone)
```

(a) Fit the following simple linear regression model in $\,\mathbb{R}$. Use the ozone measurement as the response and wind speed as the predictor.

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$

Store the results in a variable called $ozone_wind_model$. Use a t test to test the significance of the regression. Report the following:

- The null and alternative hypotheses
 - a. Null Hypothesis $H_0: \beta_1 = 0$
 - b. Alternate Hypothesis $H_1: \beta_1 \neq 0$
- The value of the test statistic

The t statistics for β_0 : **10.9278455** and β_1 : **-0.2189811**

The p-value of the test

The p value for β_0 :4.818462110^{-24} and for β_1 : 0.8267954

• A statistical decision at $\alpha = 0.01$

The P value for β_1 which is: **0.8267954** > **0.01** (α). Since P is larger than α , we **failed to reject the null Hypothesis**

A conclusion in the context of the problem

Since we failed to reject the null hypothesis, β_1 close to be 0, hence there is no significant linear relationship exists between ozone and wind speed of Ozone's dataset

(b) Fit the following simple linear regression model in R. Use the ozone measurement as the response and temperature as the predictor.

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$

Store the results in a variable called ozone temp model. Use a t test to test the significance of the regression. Report the following:

The null and alternative hypotheses

a. Null Hypothesis $H_0: \beta_1 = 0$

b. Alternate Hypothesis $H_1: \beta_1 \neq 0$

The value of the test statistic

The t statistics for β_0 : -12.4842524 and β_1 : 22.848962

The p-value of the test

The p value for β_0 :9.947454510^{-30} and for β_1 : 8.153763610^{-71}

• A statistical decision at $\alpha=0.01$

The P value for β_1 which is: **8.153763610^{-71}** < **0.01** (α). Since P is larger than α , we **reject the** null Hypothesis

A conclusion in the context of the problem

Since we reject the null hypothesis, $\beta_1 \neq 0$, hence there is significance linear relationship exists between ozone and temperature of the Ozone's dataset

Exercise 3 (Simulating Sampling Distributions)

For this exercise we will simulate data from the following model:

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$

Where $\epsilon_i \sim N(0, \sigma^2)$. Also, the parameters are known to be:

- $\beta_0 = -5$
- $\beta_1 = 3.25$ $\sigma^2 = 16$

We will use samples of size n = 50.

(a) Simulate this model 2000 times. Each time use lm() to fit a simple linear regression model, then store the value of $\hat{\beta}_0$ and $\hat{\beta}_1$. Set a seed using **your** birthday before performing the simulation. Note, we are simulating the x values once, and then they remain fixed for the remainder of the exercise.

```
beta 0 = -5
beta_1 = 3.25
sigma 2 = 16
sigma = sqrt(sigma 2)
birthday = 19770411
set.seed(birthday)
n = 50
x = seq(0, 10, length = n)
sim_data = function(x,beta0,beta1,n){
  epsilon = rnorm(n,mean=0,sd=sigma)
 y = beta0 + beta1 * x + epsilon
  data.frame(predictor = x, response = y)
}
loop\_times = 2000
beta 0 hat = rep(0,loop times)
beta_1_hat = rep(0,loop_times)
for(i in 1:loop_times){
  epsilon = rnorm(n,mean=0,sd=sigma)
 y = beta_0 + beta_1 * x + epsilon
  data = sim_data(x,beta_0,beta_1,n)
  data_model = lm(response~predictor,data=data)
 beta_0_hat[i] = summary(data_model)$coefficient[1,1]
 beta 1 hat[i] = summary(data model)$coefficient[2,1]
}
Sxx = sum((x - mean(x))^2)
sd_beta_1_hat = sqrt(sigma_2 / Sxx)
sd_beta_0_hat = sqrt(sigma_2 * ((1/n) + (mean(x)^2/Sxx)))
```

(b) Create a table that summarizes the results of the simulations. The table should have two columns, one for $\hat{\beta}_0$ and one for $\hat{\beta}_1$. The table should have four rows:

- A row for the true expected value given the known values of x
- A row for the mean of the simulated values
- A row for the true standard deviation given the known values of x
- A row for the standard deviation of the simulated values

```
library(knitr)
beta0_beta_1_summary_report = data.frame(
   attributes = c("True Exepected Value","Mean Simulated Value","True Standard Devi
ation","Standard Deviation of Simulated Value"),
   beta_0 = c(beta_0,mean(beta_0_hat),sd_beta_0_hat,sd(beta_0_hat)),
   beta_1 = c(beta_1,mean(beta_1_hat),sd_beta_1_hat,sd(beta_1_hat))
)

kable(beta0_beta_1_summary_report, format = "pandoc",padding = 2,caption = "Summar izes the results of the Simulations")
```

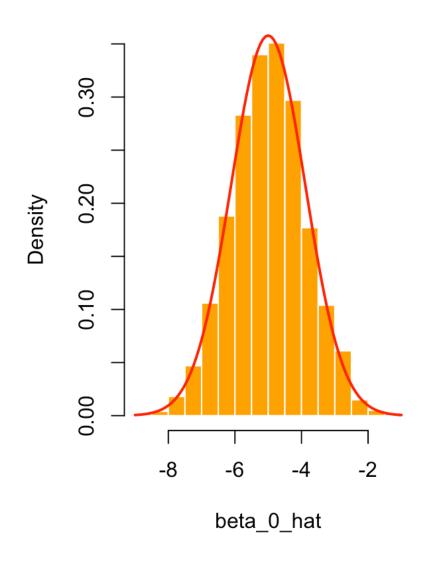
attributes	beta_0	beta_1
True Exepected Value	-5.000000	3.2500000
Mean Simulated Value	-4.989433	3.2473950
True Standard Deviation	1.114609	0.1920784
Standard Deviation of Simulated Value	1.114203	0.1897273

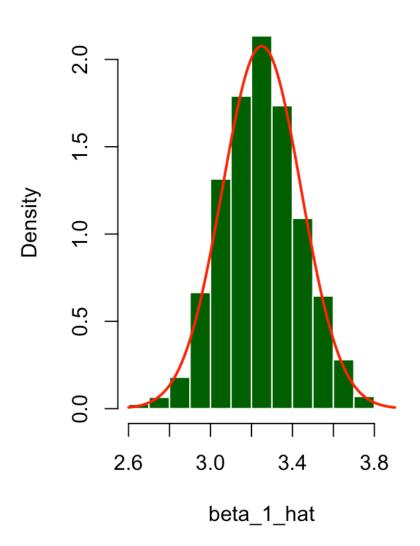
(c) Plot two histograms side-by-side:

- A histogram of your simulated values for $\hat{\beta}_0$. Add the normal curve for the true sampling distribution of $\hat{\beta}_0$.
- A histogram of your simulated values for $\hat{\beta}_1$. Add the normal curve for the true sampling distribution of $\hat{\beta}_1$.

Histogram of beta_0_hat

Histogram of beta_1_hat





Exercise 4 (Simulating Confidence Intervals)

For this exercise we will simulate data from the following model:

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$

Where $\epsilon_i \sim N(0, \sigma^2)$. Also, the parameters are known to be:

- $\beta_0 = 5$ $\beta_1 = 2$ $\sigma^2 = 9$

We will use samples of size n=25.

Our goal here is to use simulation to verify that the confidence intervals really do have their stated confidence level. Do **not** use the confint() function for this entire exercise.

(a) Simulate this model 2500 times. Each time use lm() to fit a simple linear regression model, then store the value of $\hat{\beta}_1$ and s_e . Set a seed using **your** birthday before performing the simulation. Note, we are simulating the x values once, and then they remain fixed for the remainder of the exercise.

```
beta 0 = 5
beta 1 = 2
sigma 2 = 9
sigma = sqrt(sigma 2)
```

```
birthday = 19770411
set.seed(birthday)
n = 25
x = seq(0, 2.5, length = n)
sim data = function(x,beta0,beta1,n){
 epsilon = rnorm(n,mean=0,sd=sigma)
 y = beta0 + beta1 * x + epsilon
 data.frame(predictor = x, response = y)
}
loop times = 2500
se_sim = rep(0,loop_times)
beta_1_hat = rep(0,loop_times)
for(i in 1:loop times){
 epsilon = rnorm(n,mean=0,sd=sigma)
 data = sim data(x, beta 0, beta 1, n)
 data model = lm(response~predictor,data=data)
 se = sqrt((sum((predict(data_model) - data$response)^2))/(length(data$response)
- 2))
 beta_1_hat[i] = summary(data model)$coefficient[2,1]
  se sim[i] = se
}
alpha = (1-0.95)
alpha 2 = alpha/2
crit_95 = qt(1-alpha_2, df = nrow(data) - 2)
alpha = (1-0.99)
alpha 2 = alpha/2
crit_99 = qt(1-alpha_2, df = nrow(data) - 2)
Sxx = sum((x - mean(x))^2)
lower 95 = beta 1 hat - crit 95 * se sim/sqrt(Sxx)
upper_95 = beta_1_hat + crit_95 * se_sim/sqrt(Sxx)
lower_99 = beta_1_hat - crit_99 * se_sim/sqrt(Sxx)
upper_99 = beta_1_hat + crit_99 * se_sim/sqrt(Sxx)
beta_hat_1_t = (beta_1_hat - 0) / (se_sim/sqrt(Sxx))
beta hat 1 p = 2* pt(beta hat 1 t,df=length(beta hat 1 t) - 2, lower.tail = FALSE)
mean(beta 1 > lower 95 & beta 1 < upper 95)</pre>
```

```
## [1] 0.9896

mean(beta_hat_1_p < 0.05)
```

```
## [1] 0.7072
```

```
mean(beta_hat_1_p < 0.01)
```

```
## [1] 0.486
```

- **(b)** For each of the $\hat{\beta}_1$ that you simulated, calculate a 95% confidence interval. Store the lower limits in a vector lower 95 and the upper limits in a vector upper 95. Some hints:
 - You will need to use qt() to calculate the critical value, which will be the same for each interval.
 - Remember that x is fixed, so S_{xx} will be the same for each interval.
 - You could, but do not need to write a for loop. Remember vectorized operations.

lower_95 and upper_95 is created in the above R chunk and Length of lower_95: **2500** and length of upper_95: **2500**

(c) What proportion of these intervals contains the true value of β_1 ?

mean(beta 1 > lower 99 & beta 1 < upper 99)</pre>

- **0.948** is the proportion of these intervals (95% CI) contains the true value of β_1
- (d) Based on these intervals, what proportion of the simulations would reject the test $H_0: \beta_1 = 0$ vs $H_1: \beta_1 \neq 0$ at $\alpha = 0.05$?
- **0.7072** proportion of the simulations would reject the test $H_0: \beta_1=0$ vs $H_1: \beta_1\neq 0$ at $\alpha=0.05$
- (e) For each of the $\hat{\beta}_1$ that you simulated, calculate a 99% confidence interval. Store the lower limits in a vector <code>lower_99</code> and the upper limits in a vector <code>upper_99</code>.

Length of lower_99 2500 and length of upper_99 2500

- **(f)** What proportion of these intervals contains the true value of β_1 ?
- **0.9896** is the proportion of these intervals (99% CI) contains the true value of β_1
- (g) Based on these intervals, what proportion of the simulations would reject the test $H_0: \beta_1=0$ vs $H_1: \beta_1\neq 0$ at $\alpha=0.01$?
- **0.486** proportion of the simulations would reject the test $H_0: \beta_1=0$ vs $H_1: \beta_1 \neq 0$ at $\alpha=0.05$

Exercise 5 (Prediction Intervals "without" predict)

Write a function named calc pred int that performs calculates prediction intervals:

$$\hat{y}(x) \pm t_{\alpha/2, n-2} \cdot s_e \sqrt{1 + \frac{1}{n} + \frac{(x - \bar{x})^2}{S_{xx}}}.$$

for the linear model

$$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$
.

(a) Write this function. You may use the predict() function, but you may **not** supply a value for the level argument of predict(). (You can certainly use predict() any way you would like in order to check your work.)

The function should take three inputs:

- model, a model object that is the result of fitting the SLR model with lm()
- newdata, a data frame with a single observation (row)
 - This data frame will need to have a variable (column) with the same name as the data used to fit model.
- level, the level (0.90, 0.95, etc) for the interval with a default value of 0.95

The function should return a named vector with three elements:

- estimate, the midpoint of the interval
- lower, the lower bound of the interval
- upper, the upper bound of the interval

```
calc_pred_int = function (model, newdata, level = c(0.95)) {
    #Derive the Critical Value/Y-hat and Lower Bound for the default CI - 95%
    crit_95 = qt(1-(1 - 0.95) / 2,length(predict(cat_model)) - 2)
    y_hat = predict(model,newdata=newdata)
    lwr = predict(cat_model,newdata=newdata,interval="prediction")[2]

#Calculate the function
    X_FIND = (y_hat - lwr) / crit_95

#Derive the Critical Value for the default CI - 95%
    crit = qt(1-(1 - level) / 2,length(predict(cat_model)) - 2)
    lower = (y_hat - crit * X_FIND)
    upper = (y_hat + crit * X_FIND)

results <- c(y_hat,lower,upper)
    names(results) <- c("estimate","lower","upper")
    results
}</pre>
```

(b) After writing the function, run this code:

```
newcat_1 = data.frame(Bwt = 4.0)
calc_pred_int(cat_model, newcat_1)
```

```
## estimate lower upper
## 15.77959 12.83018 18.72900
```

(c) After writing the function, run this code:

```
newcat_2 = data.frame(Bwt = 3.3)
calc_pred_int(cat_model, newcat_2, level = 0.99)
```

```
## estimate lower upper
## 12.955744 9.132013 16.779476
```