

CS498 AML,AMO HW6

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TOTAL POINTS

100 / 100

QUESTION 1

1 code for regression and resulting model.

0 / 0

✓ - **0 pts** Correct

QUESTION 2

2 a screenshot of your diagnostic plot and a few sentences of your explanation. **50 / 50**

+ **10 pts** Correct

+ **50** Point adjustment

QUESTION 3

3 a screenshot of your new diagnostic plot.

20 / 20

+ **0 pts** Correct

+ **20** Point adjustment

QUESTION 4

4 a screenshot of your code for subproblem 2. **10 / 10**

+ **0 pts** Correct

+ **0 pts** Incorrect

+ **10** Point adjustment

QUESTION 5

5 a screenshot of Box-Cox transformation plot and the best value you chose. **10 / 10**

✓ - **0 pts** Correct

- **10 pts** Incorrect

QUESTION 6

6 result of the standardized residuals of the regression after Box-Cox transformation and a plot of fitted house price against true house price. **10 / 10**

✓ - **0 pts** Correct

- **10 pts** Incorrect

QUESTION 7

7 code for subproblems 3 and 4. **0 / 0**

✓ - **0 pts** Correct

- **5 pts** Click here to replace this description.

QUESTION 8

8 late penalty **0 / 0**

✓ - **0 pts** Correct

- **5 pts** 1 day

- **10 pts** 2 days

- **15 pts** 3 days

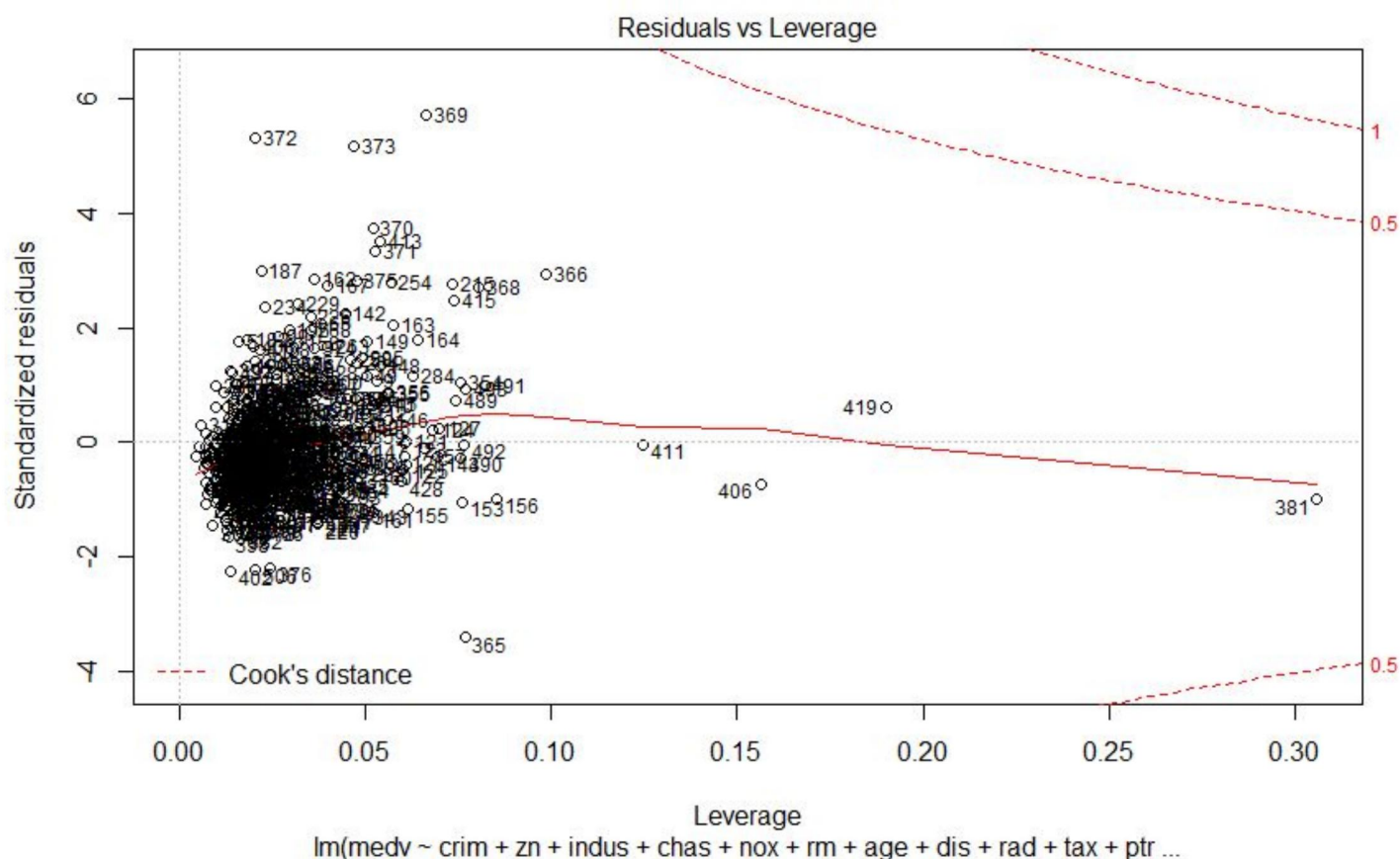
- **20 pts** 4 days

- **30 pts** max

(0 points) Page 1: code for regression and resulting model.

```
library(MASS)
data(Boston)
fit1 <- lm(medv ~ crim + zn + indus + chas + nox + rm + age + dis + rad + tax +
ptratio + black + lstat, data=Boston)
par(mfrow=c(1,1)) # Change the panel layout to 2 x 2
plot(fit1, id.n = 3)
```

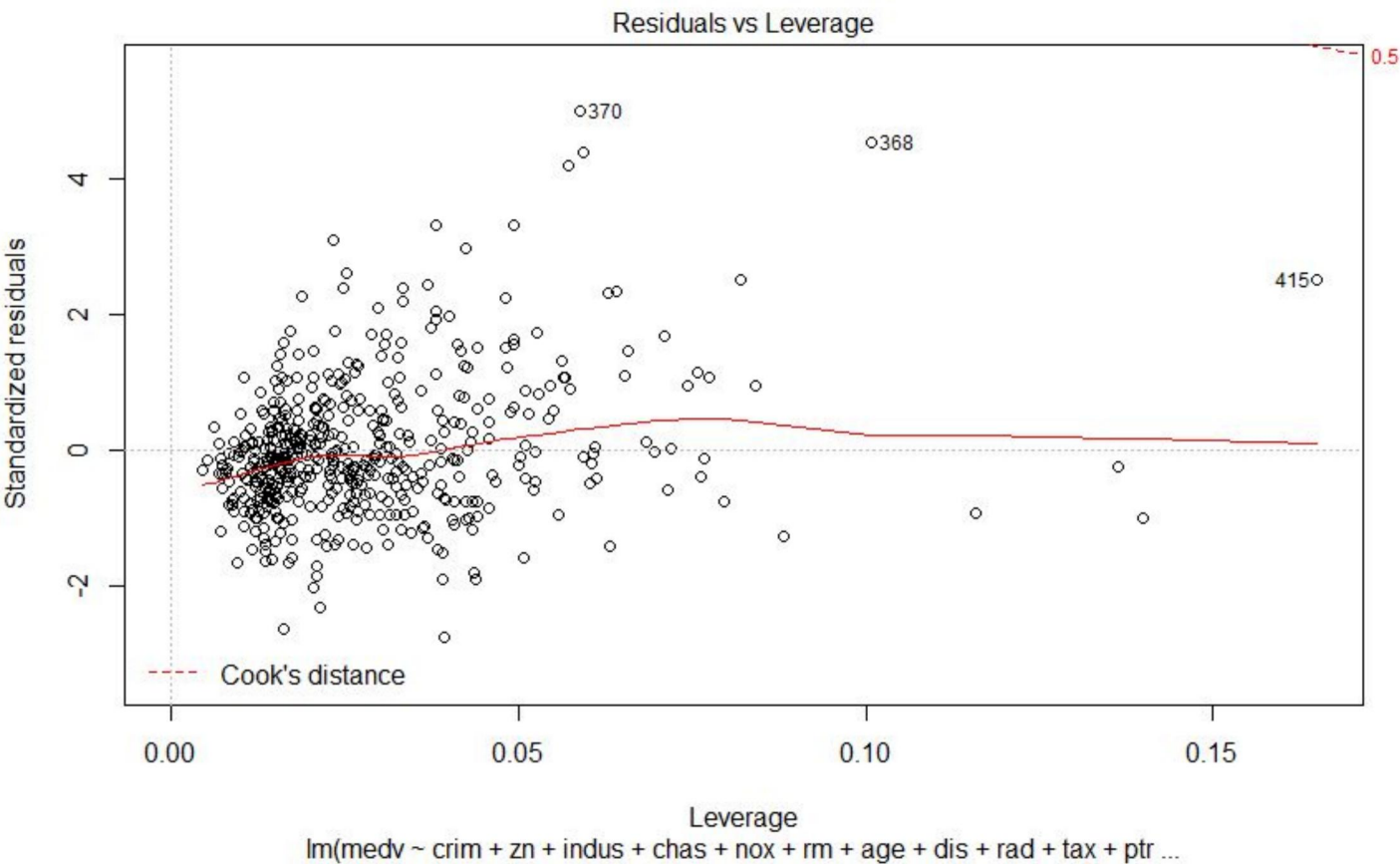
(50 points) **Page 2:** a screenshot of your diagnostic plot and a few sentences of your explanation.



The index 381, 419, 406, 411, 365, 369, 373, 372, and 366 have data that can be considered outliers.

- Index 381, 419, 406, 366, and 411 have high leverage (>0.1), which would make the prediction model depends heavily on these incorrect values.
- Index 365, 369, 373, and 372 have high standardized residual (>3), which mean they are very unlikely to be measured (less than 1% chance).

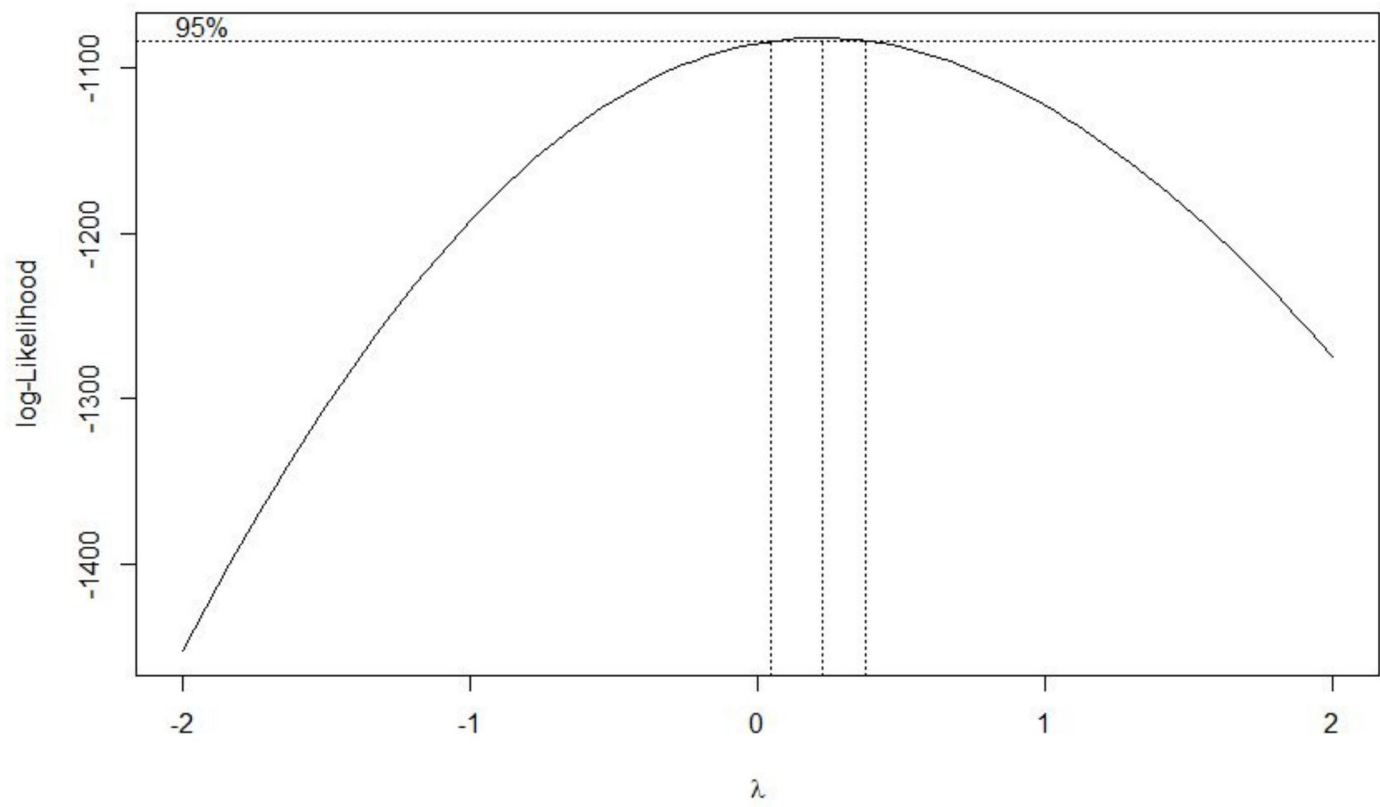
(20 points) Page 3: a screenshot of your new diagnostic plot.



(10 points) **Page 4:** a screenshot of your code for subproblem 2.

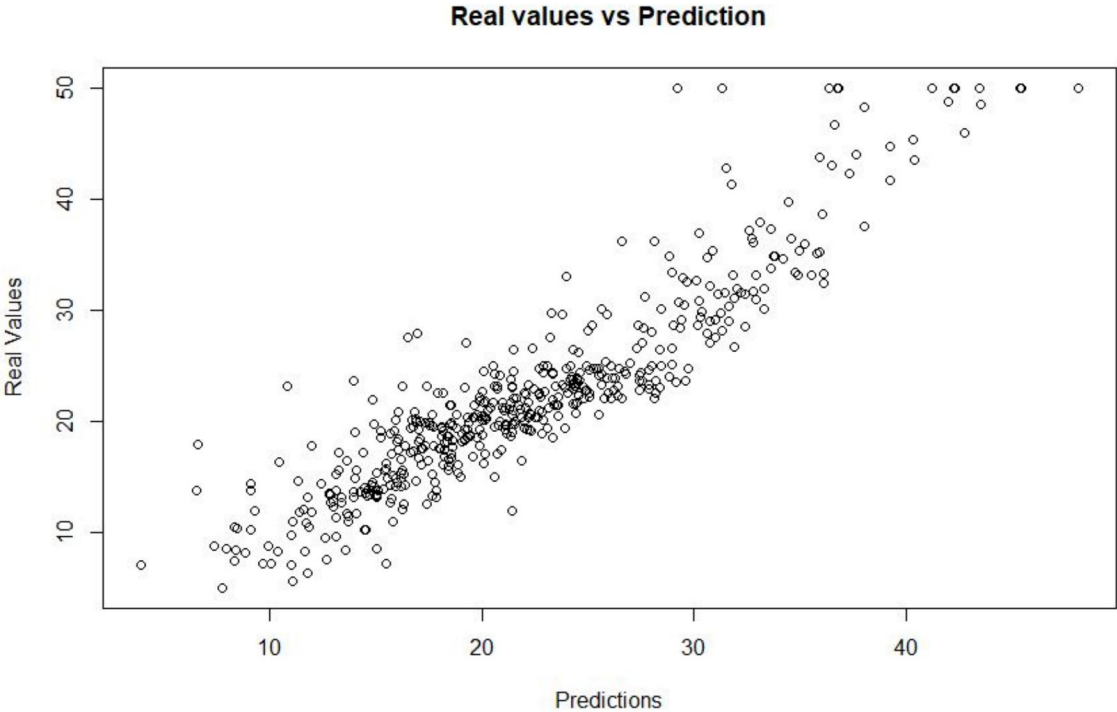
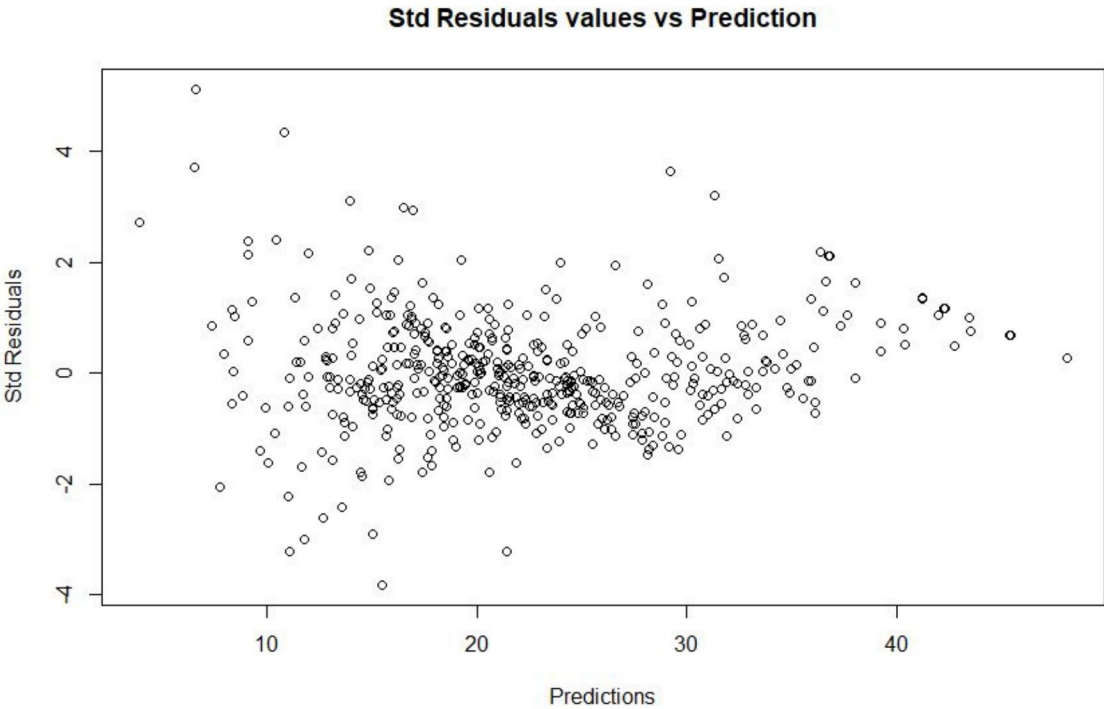
```
Boston2 <- Boston[-c(381,419,406,411,365,369,373,372,366),]  
fit2 <- lm(medv ~ crim + zn + indus + chas + nox + rm + age + dis + rad + tax +  
ptratio + black + lstat, data=Boston2)  
par(mfrow=c(1,1))  
plot(fit2, id.n = 3)
```

(10 points) **Page 5:** a screenshot of Box-Cox transformation plot and the best value you chose.



We chose 0.222222 for lambda.

(10 points) **Page 6:** result of the standardized residuals of the regression after Box-Cox transformation and a plot of fitted house price against true house price.



(0 points) Page 7: code for subproblems 3 and 4.

```
# Apply Box-Cox transformation
par(mfrow=c(1,1))
Box = boxcox(Boston2$medv ~ 1)
Cox = data.frame(Box$x, Box$y) # Create a data frame with the results
Cox2 = Cox[with(Cox, order(-Cox$Box.y)),] # Order the new data frame by
decreasing y
Cox2[1,] # Display the lambda with the greatest
log likelihood
lambda = Cox2[1, "Box.x"] # Extract that lambda
T_box = (Boston2$medv ^ lambda - 1)/lambda # Transform the original data
Boston2$boxcox <- T_box

# Fit the Box-Cox transformed data
fit3 <- lm(boxcox ~ crim + zn + indus + chas + nox + rm + age + dis + rad + tax
+ ptratio + black + lstat, data=Boston2)
Boston2.stdres <- rstandard(fit3)
par(mfrow=c(1,1))

# Plot "Real values vs Prediction"
plot((((fit3$fitted.values)*0.222222222)+1)^(1/0.222222222), Boston2$medv,
ylab = "Real Values", xlab = "Predictions", main = "Real values vs Prediction")

# Plot "Std Residuals values vs Prediction"
plot((((fit3$fitted.values)*0.222222222)+1)^(1/0.222222222), Boston2.stdres,
ylab = "Std Residuals", xlab = "Predictions", main = "Std Residuals values vs
Prediction")
```