## Question 1:

- a. There are 506 rows and 13 variables in the data set Boston. The variable Istat represents the lower status of the population (percent).
- b.  $B_0^- = -3.33054$ , standard error = 0.69376, p-value = 2.09e-06  $B_1^- = 0.54880$ , standard error = 0.04776, p-value = <2e-16 So the crim rate is calculated by = (-3.3305) + 0.5488(Istat)
- c. zn: if the proportion of residential land zoned for lots overs 25,000 sq.ft increases then the per capita crime rate decreases

$$B_0^* = 4.45369, B_1^* = -0.07393$$

indus: if the proportion of non-retail business acres per town increases, the per capita crime rate increases

$$B_0^- = -2.06374$$
,  $B_1^- = 0.50978$ 

chas: if the Charles River dummy variable increases, the per capita crime rate decreases  $B_0^2 = 3.7444$ ,  $B_1^2 = -1.8928$ 

nox: if the nitrogen oxide concentration increases, the per capita crime rate increases  $B_0^2 = -13.720$ ,  $B_1^2 = 31.249$ 

rm: if the average number of rooms per dwelling increases, the per capita crime rate decreases

$$B_0^2 = 20.482$$
,  $B_1^2 = -2.684$ 

age: if the proportion of owner-occupied units built prior to 1940 increases, the per capita crime rate decreases

$$B_0^- = -3.77791$$
,  $B_1^- = 0.10779$ 

dis: if the weighted mean of the distances to five Boston employment centers increases, the per capita crime rate decreases

$$B_0^- = 9.4993$$
,  $B_1^- = -1.5509$ 

rad: if the index of accessibility to radial highways increases, the per capita crime rate increases

$$B_0^- = -2.28716$$
,  $B_1^- = 0.61791$ 

tax: if the full-value property-tax rate per \$10k increases, the per capita crime rate increases

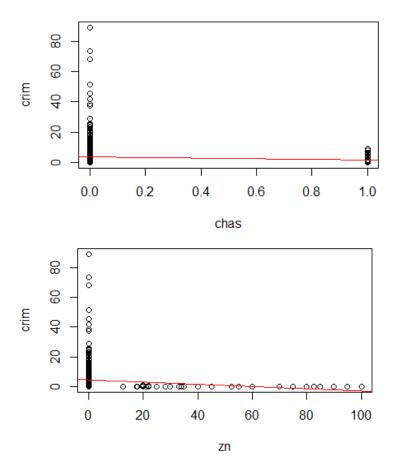
$$B_0^- = -8.528369$$
,  $B_1^- = 0.029742$ 

ptratio: if the pupil-teacher ratio by town increases, the per capita crime rate increases  $B_0^2 = -17.6469$ ,  $B_1^2 = 1.1520$ 

medv: if the median value of owner-occupied homes increases, the per capita crime rate decreases

$$B_0^* = 11.79654$$
,  $B_1^* = -0.36316$ 

We will need to test H0: B1 = 0 to tell which predictors are significant. It can be observed that all predictors have a p-value less than 0.05 except the predictor "chas." Thus, there is a statistically significant association between each predictor and response except for the "chas" predictor. A scatter plot for the chas predictor and zn predictor are down below to show that chas does not have a statistically significant association with the response crim.



d. We can reject the null hypothesis for zn, dis, rad, and medv because they all have values less than 0.05 ( $\alpha = 0.05$ )

```
call:
lm(formula = crim \sim ., data = Boston)
Residuals:
           1Q Median
                          3Q
  Min
                               Max
-8.534 -2.248 -0.348
                      1.087 73.923
Coefficients:
              Estimate Std. Error t value Pr(>|t|)
                                     1.946 0.052271
(Intercept) 13.7783938
                        7.0818258
             0.0457100
                        0.0187903
                                     2.433 0.015344
indus
             -0.0583501
                         0.0836351
                                    -0.698 0.485709
             -0.8253776
                         1.1833963
                                    -0.697 0.485841
chas
             -9.9575865
                         5.2898242
                                    -1.882 0.060370
nox
             0.6289107
                         0.6070924
rm
                                     1.036 0.300738
             -0.0008483
                         0.0179482
                                    -0.047 0.962323
age
dīs
             -1.0122467
                         0.2824676
                                    -3.584 0.000373 ***
             0.6124653
                         0.0875358
                                     6.997 8.59e-12
rad
            -0.0037756
                        0.0051723
                                    -0.730 0.465757
tax
ptratio
            -0.3040728
                         0.1863598
                                    -1.632 0.103393
İstat
             0.1388006
                         0.0757213
                                     1.833 0.067398
             -0.2200564
                        0.0598240
                                    -3.678 0.000261
medv
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
Residual standard error: 6.46 on 493 degrees of freedom
Multiple R-squared: 0.4493,
                                Adjusted R-squared: 0.4359
F-statistic: 33.52 on 12 and 493 DF, p-value: < 2.2e-16
```

e. The table below shows that using multiple simple linear regression models is not sufficient compared to a multiple linear regression model. This is because in a simple

regression model, all other predictors are ignored while in a multiple regression model, the other predictors are fixed.

```
simple
                        multiple
        -0.07393498 0.0457100386
zn
        0.50977633 -0.0583501107
indus
        -1.89277655 -0.8253775522
chas
        31.24853120 -9.9575865471
nox
       -2.68405122 0.6289106622
        0.10778623 -0.0008482791
age
        -1.55090168 -1.0122467382
dis
        0.61791093 0.6124653115
rad
        0.02974225 -0.0037756465
tax
ptratio 1.15198279 -0.3040727572
        0.54880478 0.1388005968
lstat
       -0.36315992 -0.2200563590
medv
```

f. Training MSE = 42.49345

Test MSE = 41.19923

g. Training MSE = 43.97466

Test MSE = 39.62763

We were using less predictors in this model so I expected the training MSE to be higher and the test MSE to be lower than results in part f.

h.  $H_0$ :  $B_j = 0$ ,  $H_1$ :  $B_j \neq 0$ Test statistic: 3.264

Null distribution: t-distribution with 495 degrees of freedom n-p+1 = 506-12+1=495

p-value: 0.001173

The null hypothesis is rejected and the results are statistically significant.  $B_j$  is significantly different from 0 at significance level 0.05. We have to assume  $H_0$  is true for this to work

i.  $H_0$ :  $B_i = 5$ ,  $H_1$ :  $B_i \neq 5$ 

Test statistic: (2.839993 - 5) / 0.870007 = -2.4827

Null distribution: t-distribution with 495 degrees of freedom n-p+1 = 506-12+1=495

p-value: 0.01336767

The null hypothesis is rejected and the results are statistically significant.  $B_j$  is significantly different from 0 at significance level 0.05. We have to assume  $H_0$  is true for this to work

## Question 2:

- a. 1. There is a linear relationship between X1, X2..., Xp and Y
  - 2.  $E(\epsilon i) = 0$
  - 3.  $Var(\epsilon i) = \sigma^2$
  - 4. ci's are uncorrelated
- b. No, this is not a true population regression model. The student used E(Y<sub>i</sub>). They should've used Y<sub>i</sub>. Basically, instead of using a particular Y<sub>i</sub>, they used the true mean of Y<sub>i</sub> in their model.
- c. This is false. The student should just be using  $B_j$  but they used  $B_j^*$  which is an estimator of the parameter  $B_i$ . We do not need to test an estimate.

d. This is true. The training MSE should be smaller than the test MSE because the training MSE is calculated from data used to build the model while test MSE is calculated using data the model has not seen yet.

## **Problem 3:**

```
B0 = 0, B1 = 1, B2 = 2, B3 = 3, B4 = 4, B5 = 5
```

There are 19 individual t-tests that are significant at  $\alpha$  = 0.05. Here, there should only be 5 predictors that are significant but the individual t-tests tell us that there are 19. We should not depend on these individual t-tests to tell us whether there is a relationship between at least one of the predictors and Y. Doing these individual t-tests could result in some small p-values. And if we do a lot of individual tests, there might be a chance of type 1 errors. For example, if we are predicting salary based on job, factors like eye color, and height are unrelated predictors. But we might see that these unrelated predictors are significant due to the multiple testing problem – this basically amplifies the probability of a false-positive result. Using f-tests can solve the issue and make sure that at least one predictor is significant.

```
set.seed(2)
x = matrix(NA, 1000, 200)
n=1000
beta_0 = 0
beta_1 = 1
beta_2 = 2
beta_3 = 3
beta_4 = 4
beta_5 = 5
error = rnorm(n,0,1)
for(i in 1:200){
 x[,i] = rnorm(1000)
Y = beta_0 + beta_1*x[,1] + beta_2*x[,2] + beta_3*x[,3] + beta_4*x[,4] + beta_5*x[,5] + error
data = as.data.frame(cbind(Y,x))
fit = lm(Y\sim., data=data)
summary(fit)
p_values = summary(fit)$coefficients[,4]
length(which(p_values<0.05))
```