INFX 573: Problem Set 6 - Regression

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Due: Tuesday, November 15, 2016

Collaborators:

Instructions:

Before beginning this assignment, please ensure you have access to R and RStudio.

- 1. Download the problemset6.Rmd file from Canvas. Open problemset6.Rmd in RStudio and supply your solutions to the assignment by editing problemset6.Rmd.
- 2. Replace the "Insert Your Name Here" text in the author: field with your own full name. Any collaborators must be listed on the top of your assignment.
- 3. Be sure to include well-documented (e.g. commented) code chucks, figures and clearly written text chunk explanations as necessary. Any figures should be clearly labeled and appropriately referenced within the text.
- 4. Collaboration on problem sets is acceptable, and even encouraged, but each student must turn in an individual write-up in his or her own words and his or her own work. The names of all collaborators must be listed on each assignment. Do not copy-and-paste from other students' responses or code.
- 5. When you have completed the assignment and have **checked** that your code both runs in the Console and knits correctly when you click **Knit PDF**, rename the R Markdown file to YourLastName_YourFirstName_ps6.Rmd, knit a PDF and submit the PDF file on Canvas.

Setup:

In this problem set you will need, at minimum, the following R packages.

```
# Load standard libraries
library(tidyverse)

## Warning: package 'tidyverse' was built under R version 3.2.5

## Warning: package 'ggplot2' was built under R version 3.2.5

## Warning: package 'tibble' was built under R version 3.2.5

## Warning: package 'tidyr' was built under R version 3.2.5

## Warning: package 'readr' was built under R version 3.2.5

## Warning: package 'purrr' was built under R version 3.2.5

## Warning: package 'dplyr' was built under R version 3.2.5

library(MASS) # Modern applied statistics functions
```

Housing Values in Suburbs of Boston

In this problem we will use the Boston dataset that is available in the MASS package. This dataset contains information about median house value for 506 neighborhoods in Boston, MA. Load this data and use it to answer the following questions.

```
#data(package = "MASS")
boston_data <- Boston
#?Boston
#str(boston_data)</pre>
```

1. Describe the data and variables that are part of the Boston dataset. Tidy data as necessary.

The data set contains information about varios parameters about Boston's population and locality. For example, the per capita crime rate by town, proportion of residential land, proportion of non-retail business per town, if the tract occupies charles river or not (boolean variable), nitrogen oxide concetration, average numbe of rooms in an apartment, proportion of owner-occupied units, weighted mean of distances to five Boston employment centers, index of accessibility to radial highways, property-tax rate per \$10,000, pupil-teacher ratio by town, proportion of blacks by town, lower status of the population and median value of owner-occupied homes.

2. Consider this data in context, what is the response variable of interest? Discuss how you think some of the possible predictor variables might be associated with this response.

The response variable in the context of this data would be median value of owner-occupied homes in \$1000s (medv). With the help of other variables like proportion of owner-occupied units, weighted mean of distances to 5 Boston employment centres, pupil-teacher ratio by town, proportion of blacks by town, median value of owner occupied homes, lower status of the population (percent), proportion of non-retail business acres per town, per capita crime rate by town. Here, we can find an association of how median value of owner-occupied homes increases or decreases if we consider proportion of owner-occupied units, weighted mean of distances to 5 Boston towns etc as predictor variables.

3. For each predictor, fit a simple linear regression model to predict the response. In which of the models is there a statistically significant association between the predictor and the response? Create some plots to back up your assertions.

Let's consider the association between median value of owner-occupied homes in \$1000s as the response variable and crime rate by town as the predictor variable

```
lm_crim <- lm(medv~crim, data = boston_data)
summary(lm_crim)</pre>
```

```
##
## Call:
## lm(formula = medv ~ crim, data = boston_data)
##
## Residuals:
##
       Min
                    Median
                1Q
                                 ЗQ
                                        Max
## -16.957 -5.449
                    -2.007
                              2.512
                                    29.800
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 24.03311
                           0.40914
                                      58.74
                                              <2e-16 ***
               -0.41519
                                      -9.46
## crim
                           0.04389
                                              <2e-16 ***
## ---
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 8.484 on 504 degrees of freedom
## Multiple R-squared: 0.1508, Adjusted R-squared: 0.1491
## F-statistic: 89.49 on 1 and 504 DF, p-value: < 2.2e-16
```

Residuals vs Fitted 30 20 Residuals 0 9 00 0 0 0 -10 -100 10 20 Fitted values Im(medv ~ crim)

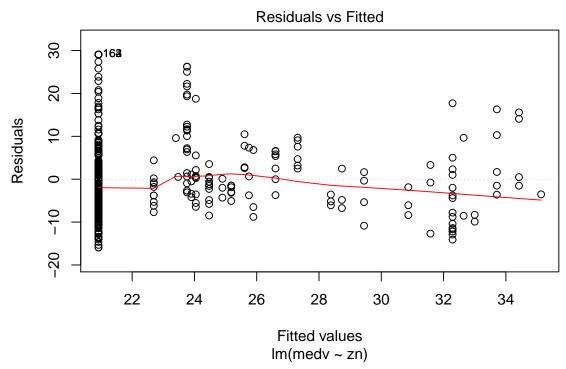
The visualization indicates that there is not a linear relationship between crim and medv. The data points are not bouncing randomly across the linear regression line as there is no sero line due to the uneven nature of the points. But, there is definitely a significant association between median value of homes and crime rate as the p-value is much less than 0.05.

We can also analyze the relationship between median value of homes and proportion of residential land zoned for lots over 25000 sq.ft.

```
lm_zn <- lm(medv~zn, data = boston_data)
summary(lm_zn)</pre>
```

```
##
## Call:
## lm(formula = medv ~ zn, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                                        Max
                    Median
                                 3Q
## -15.918 -5.518
                    -1.006
                              2.757
                                     29.082
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 20.91758
                            0.42474
                                     49.248
                                               <2e-16 ***
## zn
                0.14214
                            0.01638
                                      8.675
                                               <2e-16 ***
## ---
## Signif. codes:
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.587 on 504 degrees of freedom
```

```
## Multiple R-squared: 0.1299, Adjusted R-squared: 0.1282
## F-statistic: 75.26 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_zn, which = c(1))</pre>
```

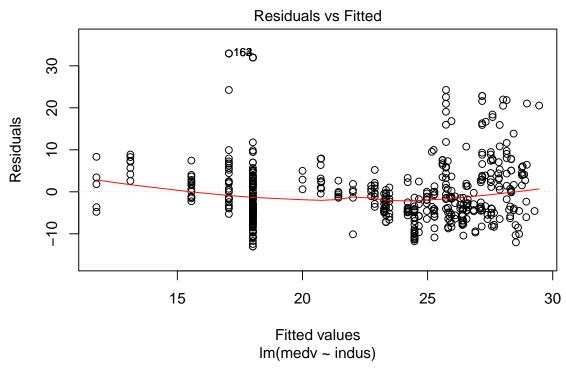


The reiduals vs fitted values graph indicates that the median value of homes and proportion of residential land zoned for lots are not linearly related as for every corresponding fitted value there is not a residual error to nullify the effect of that error. Also, the regression line is not linear. But, the the p-value is much less than 0.05 which indicates that there is a significant relationship between medv and zn but not linear.

```
lm_indus <- lm(medv~indus, data = boston_data)
summary(lm_indus)</pre>
```

```
##
## Call:
## lm(formula = medv ~ indus, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                    Median
                                 3Q
                                        Max
## -13.017 -4.917
                    -1.457
                              3.180
                                     32.943
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 29.75490
                            0.68345
                                      43.54
                                               <2e-16 ***
## indus
               -0.64849
                            0.05226
                                     -12.41
                                               <2e-16 ***
## ---
## Signif. codes:
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.057 on 504 degrees of freedom
```

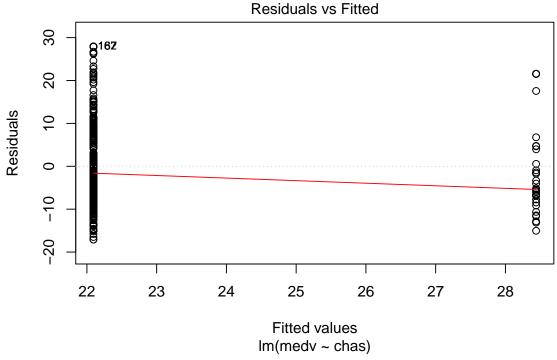
```
## Multiple R-squared: 0.234, Adjusted R-squared: 0.2325
## F-statistic: 154 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_indus, which = c(1))</pre>
```



The shape of the regression line indicates that there is a non-linear relationship between median values of homes and proportion of non-retail business acres. Also, there are some outliers which do not have corresponding residuals below the trend line.

```
lm_chas <- lm(medv~chas, data = boston_data)
summary(lm_chas)</pre>
```

```
##
## Call:
## lm(formula = medv ~ chas, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                    Median
                                ЗQ
                                        Max
## -17.094 -5.894
                    -1.417
                             2.856
                                    27.906
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 22.0938
                            0.4176
                                    52.902 < 2e-16 ***
## chas
                 6.3462
                            1.5880
                                     3.996 7.39e-05 ***
##
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
## Residual standard error: 9.064 on 504 degrees of freedom
## Multiple R-squared: 0.03072,
                                    Adjusted R-squared: 0.02879
## F-statistic: 15.97 on 1 and 504 DF, p-value: 7.391e-05
```



Fitting a linear model for variables with a Bernoulli distribution is viable. For example, one unit increase in a berboulli variable is either a 1 or 0 and there are no continuous values for such variables. So, it is not helpful to analyze such variables using linear regression models, logistic models fit best for Bernoulli distributed variables.

```
lm_nox <- lm(medv~nox, data = boston_data)
summary(lm_nox)</pre>
```

```
##
## Call:
## lm(formula = medv ~ nox, data = boston_data)
## Residuals:
##
       Min
                1Q
                   Median
                                3Q
                                       Max
## -13.691 -5.121
                   -2.161
                             2.959
                                    31.310
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                 41.346
                             1.811
                                     22.83
                                              <2e-16 ***
## nox
                -33.916
                             3.196
                                   -10.61
                                              <2e-16 ***
## ---
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 8.323 on 504 degrees of freedom
## Multiple R-squared: 0.1826, Adjusted R-squared: 0.181
## F-statistic: 112.6 on 1 and 504 DF, p-value: < 2.2e-16
```

Residuals vs Fitted 15 20 25 Fitted values Im(medv ~ nox)

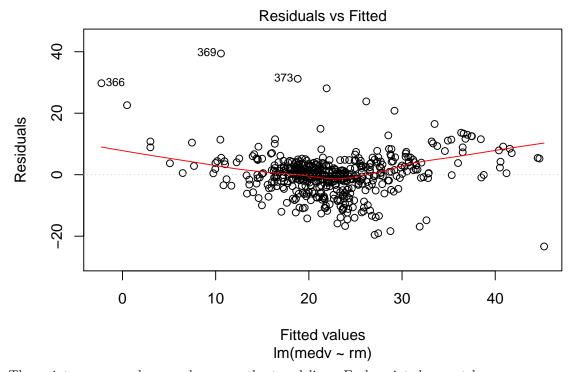
The fitted values are scattered all over the graph and they are not evenly distributed across the trend line. Also, the trend line is not a zero line and hence does not satisfy linear relationship criterion. But, the summary statistics shows there is a significant relationship between median value of homes and nitrogen oxides concentration as the p-value is much less than 0.05 may not be linear.

Let's also consider the relationship between median value of homes and average number of rooms per dwelling.

```
lm_rm <- lm(medv~rm, data = boston_data)
summary(lm_rm)</pre>
```

```
##
## Call:
## lm(formula = medv ~ rm, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                                        Max
                    Median
                                 3Q
##
   -23.346 -2.547
                     0.090
                              2.986
                                     39.433
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                -34.671
                              2.650
                                     -13.08
## (Intercept)
                                               <2e-16 ***
## rm
                  9.102
                              0.419
                                      21.72
                                               <2e-16 ***
##
## Signif. codes:
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.616 on 504 degrees of freedom
```

```
## Multiple R-squared: 0.4835, Adjusted R-squared: 0.4825
## F-statistic: 471.8 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_rm, which = c(1))</pre>
```



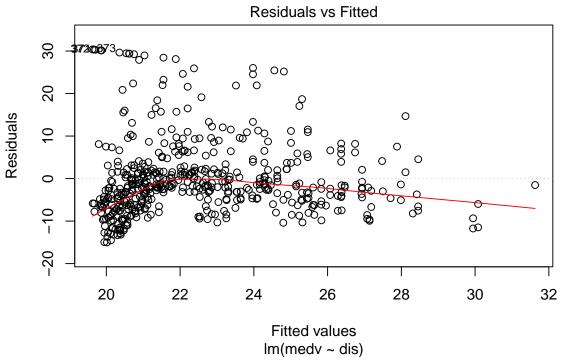
The points are spread unevenly across the trend line. Each point does not have a corresponding residual value to compensate for the error term. Hence, there is not a linear relationship between the median values of homes and average number of rooms per dwelling.

Let's consider another redictor variable dis which is the weighted mean of distances to 5 Boston employment centres.

```
lm_dis <- lm(medv~dis, data = boston_data)
summary(lm_dis)</pre>
```

```
##
## Call:
## lm(formula = medv ~ dis, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                    Median
                                 3Q
                                        Max
## -15.016 -5.556
                    -1.865
                              2.288
                                     30.377
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                                     22.499 < 2e-16 ***
## (Intercept)
               18.3901
                             0.8174
## dis
                 1.0916
                             0.1884
                                      5.795 1.21e-08 ***
## ---
## Signif. codes:
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Residual standard error: 8.914 on 504 degrees of freedom
## Multiple R-squared: 0.06246, Adjusted R-squared: 0.0606
## F-statistic: 33.58 on 1 and 504 DF, p-value: 1.207e-08
plot(lm_dis, which = c(1))
```



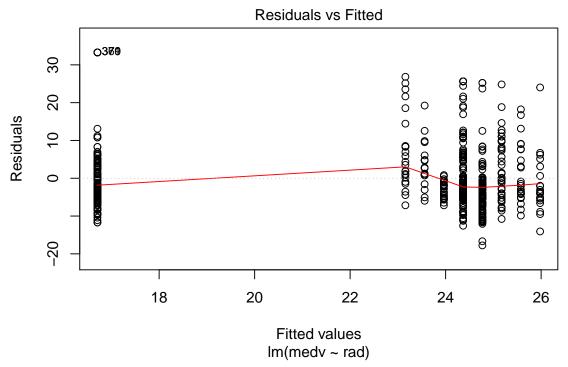
The regression line is not linear and the points are scattered across the line unevenly. Hence, there is ni linear relationship between the median value of homes and weighted mean of distances to 5 Boston employment centres. Also, there are outliers marked by the row numbers in the graph.

Let's also consider the median value of homes to accessibility to radial highways.

```
lm_rad <- lm(medv~rad, data = boston_data)
summary(lm_rad)</pre>
```

```
##
## Call:
## lm(formula = medv ~ rad, data = boston_data)
##
## Residuals:
                                 3Q
       Min
                1Q
                    Median
                                        Max
## -17.770 -5.199
                    -1.967
                              3.321
                                     33.292
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                                     46.964
## (Intercept) 26.38213
                            0.56176
                                               <2e-16 ***
## rad
               -0.40310
                            0.04349
                                     -9.269
                                               <2e-16 ***
## ---
## Signif. codes:
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Residual standard error: 8.509 on 504 degrees of freedom
## Multiple R-squared: 0.1456, Adjusted R-squared: 0.1439
## F-statistic: 85.91 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_rad, which = c(1))</pre>
```



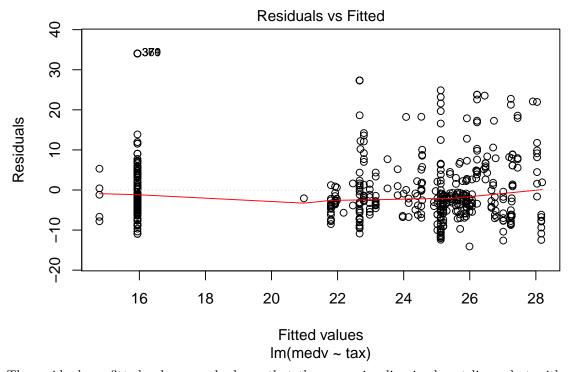
The curve indicates a non-linear relationship between the 2 variables and also the data points are scattered across the curve with no corresponding residuals on both sides of the curve.

We will also consider the effect of full-value property tax on median value of homes.

```
lm_tax <- lm(medv~tax, data = boston_data)
summary(lm_tax)</pre>
```

```
##
## Call:
## lm(formula = medv ~ tax, data = boston_data)
##
## Residuals:
##
      Min
                1Q
                   Median
                                3Q
                                       Max
## -14.091 -5.173
                   -2.085
                             3.158
                                    34.058
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 32.970654
                           0.948296
                                      34.77
                                              <2e-16 ***
## tax
               -0.025568
                           0.002147
                                     -11.91
                                              <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.133 on 504 degrees of freedom
```

```
## Multiple R-squared: 0.2195, Adjusted R-squared: 0.218
## F-statistic: 141.8 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_tax, which = c(1))</pre>
```



The residuals vs fitted values graph shows that the regression line is almost linear but with some amount of non-linearity due to the uneven distribution of data points.

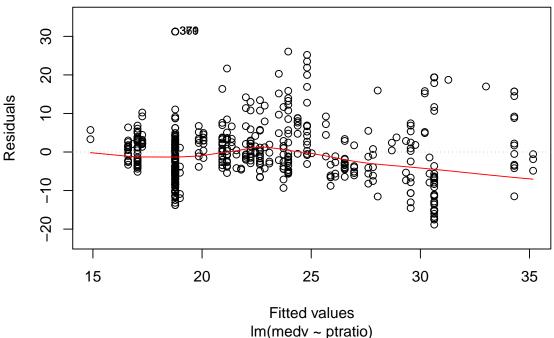
Relationship between pupil-teacher ratio and median value of homes

```
lm_ptratio <- lm(medv~ptratio, data = boston_data)
summary(lm_ptratio)</pre>
```

```
##
## Call:
## lm(formula = medv ~ ptratio, data = boston_data)
##
## Residuals:
##
        Min
                  1Q
                       Median
                                     3Q
                                             Max
                      -0.6426
## -18.8342 -4.8262
                                3.1571 31.2303
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                             3.029
                                     20.58
                                              <2e-16 ***
## (Intercept)
                 62.345
                                              <2e-16 ***
                 -2.157
                             0.163 -13.23
## ptratio
## ---
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 7.931 on 504 degrees of freedom
## Multiple R-squared: 0.2578, Adjusted R-squared: 0.2564
```

```
## F-statistic: 175.1 on 1 and 504 DF, p-value: < 2.2e-16
plot(lm_ptratio, which = c(1))</pre>
```

Residuals vs Fitted

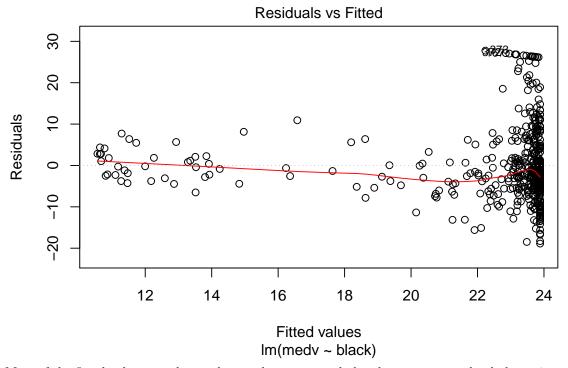


The relationship between median value of homes and pupil-teacher ration is clearly no-linear as the each ftted value does not have a residual value associated with it on the other side of the trend line.

Relationship between proportion of blacks by town and median value of homes

```
lm_black <- lm(medv~black , data = boston_data)
summary(lm_black)</pre>
```

```
##
## Call:
## lm(formula = medv ~ black, data = boston_data)
##
## Residuals:
##
       Min
                                3Q
                1Q
                    Median
                                        Max
## -18.884
           -4.862
                    -1.684
                             2.932
                                    27.763
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 10.551034
                           1.557463
                                       6.775 3.49e-11 ***
                           0.004231
                                       7.941 1.32e-14 ***
## black
                0.033593
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 8.679 on 504 degrees of freedom
## Multiple R-squared: 0.1112, Adjusted R-squared: 0.1094
## F-statistic: 63.05 on 1 and 504 DF, p-value: 1.318e-14
```



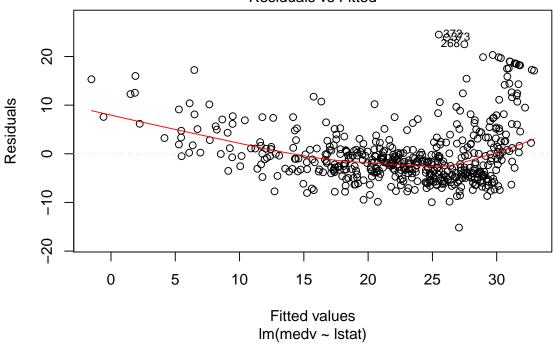
Most of the fitted values are clustered around one area and also they are not randomly bouncing across the trend line. Hence, there is no evidence of a linear relationship between the median value of homes and proportion of black population by town.

Lower status of population and median value of homes

```
lm_lstat <- lm(medv~lstat, data = boston_data)
summary(lm_lstat)</pre>
```

```
##
## Call:
## lm(formula = medv ~ lstat, data = boston data)
##
## Residuals:
##
       Min
                1Q
                    Median
                                ЗQ
                                        Max
## -15.168
           -3.990
                    -1.318
                             2.034
                                     24.500
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 34.55384
                           0.56263
                                      61.41
                                              <2e-16 ***
               -0.95005
                           0.03873
                                    -24.53
                                              <2e-16 ***
## lstat
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 6.216 on 504 degrees of freedom
## Multiple R-squared: 0.5441, Adjusted R-squared: 0.5432
## F-statistic: 601.6 on 1 and 504 DF, p-value: < 2.2e-16
```

Residuals vs Fitted

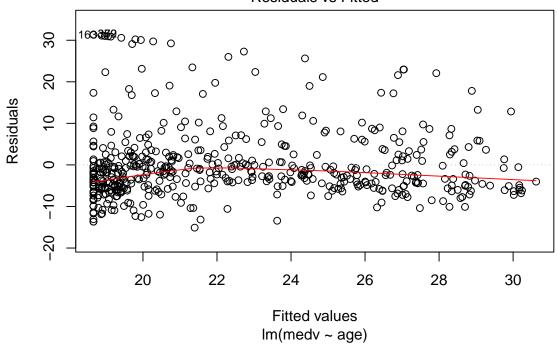


Again, the trend line indicates a non-linear relationship between the 2 variables and also there are some outliers with no corresponding residuals to nulligy the effect of this error.

```
lm_age <- lm(medv~age , data = boston_data)
summary(lm_age)</pre>
```

```
##
## Call:
## lm(formula = medv ~ age, data = boston_data)
##
## Residuals:
##
       Min
                1Q
                    Median
                                3Q
                                        Max
## -15.097 -5.138
                    -1.958
                             2.397
                                    31.338
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
                                    31.006
## (Intercept) 30.97868
                           0.99911
                                              <2e-16 ***
## age
               -0.12316
                           0.01348
                                   -9.137
                                              <2e-16 ***
## ---
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 8.527 on 504 degrees of freedom
## Multiple R-squared: 0.1421, Adjusted R-squared: 0.1404
## F-statistic: 83.48 on 1 and 504 DF, p-value: < 2.2e-16
```

Residuals vs Fitted

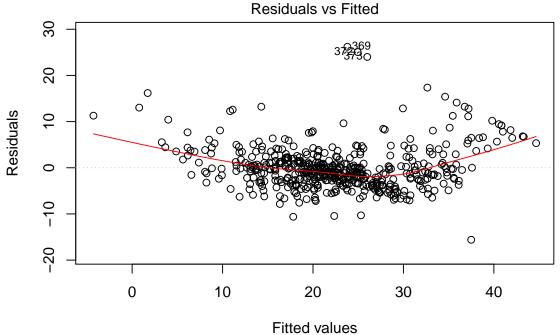


The points are randomly bouncing around the trend line but there are outliers which have no corresponding residual error value on the other side of the trend line. Thus, they do not staisfy the criterion for linear relationship.

4. Fit a multiple regression model to predict the response using all of the predictors. Describe your results. For which predictors can we reject the null hypothesis $H_0: \beta_i = 0$?

```
##
## Call:
## lm(formula = medv ~ crim + zn + indus + chas + nox + rm + rad +
##
       dis + tax + ptratio + lstat + black + age, data = boston_data)
##
## Residuals:
       Min
                1Q
                    Median
                                 3Q
                                        Max
##
  -15.595 -2.730
                    -0.518
                              1.777
                                     26.199
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
                3.646e+01 5.103e+00
                                        7.144 3.28e-12 ***
## (Intercept)
## crim
               -1.080e-01
                           3.286e-02
                                       -3.287 0.001087 **
## zn
                4.642e-02
                           1.373e-02
                                        3.382 0.000778 ***
## indus
                2.056e-02
                           6.150e-02
                                        0.334 0.738288
                2.687e+00 8.616e-01
                                        3.118 0.001925 **
## chas
```

```
## nox
               -1.777e+01
                           3.820e+00
                                       -4.651 4.25e-06 ***
                3.810e+00
                           4.179e-01
                                        9.116 < 2e-16 ***
## rm
## rad
                3.060e-01
                           6.635e-02
                                        4.613 5.07e-06 ***
## dis
               -1.476e+00
                           1.995e-01
                                       -7.398 6.01e-13 ***
## tax
               -1.233e-02
                           3.760e-03
                                       -3.280 0.001112 **
               -9.527e-01
                           1.308e-01
                                       -7.283 1.31e-12 ***
## ptratio
## 1stat
               -5.248e-01
                           5.072e-02 -10.347
                                               < 2e-16 ***
                                        3.467 0.000573 ***
## black
                9.312e-03
                           2.686e-03
## age
                6.922e-04
                           1.321e-02
                                        0.052 0.958229
## ---
## Signif. codes:
                           0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.745 on 492 degrees of freedom
## Multiple R-squared: 0.7406, Adjusted R-squared: 0.7338
## F-statistic: 108.1 on 13 and 492 DF, p-value: < 2.2e-16
plot(lm_multiple, which = c(1))
```



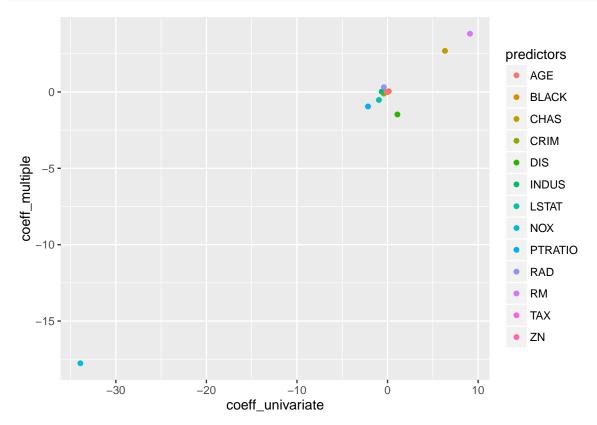
Im(medv ~ crim + zn + indus + chas + nox + rm + rad + dis + tax + ptratio + ...

The multiple regression model shows that we can reject the null hypothesis for all those predictors whose p-value is much less than 0.05 which means there is a significant relationship between the response and predictor. But, the residuals vs fitted values graph shows that there is no linear relationship between the response and all the predictor variables as the trend line is not linear which indicates that there are certain data points which do not have a corresponding match of residuals to nullify the elemnt of error. Also, there are some outliers which refrain the model from following the linearity priciples.

Thus, after analyzing the summary statistics of multiple regression model, we can safely reject the null hypothesis for the following variables: crim, zn, chas, nox, dis, ptratio, rad, rm, tax, lstat and black.

5. How do your results from (3) compare to your results from (4)? Create a plot displaying the univariate regression coefficients from (3) on the x-axis and the multiple regression coefficients from part (4) on

the y-axis. Use this visualization to support your response.



When we observe the coefficients of univariate analysis and multivariate analysis in the intermediate table coeff_df, it is evident that in multiple regression model the coefficient values decrease as compared to univariate regression model. From the graph it is visible that all the coefficients that have higher values in univariate analysis shows reduced coefficient values in the graph.

6. Is there evidence of a non-linear association between any of the predictors and the response? To answer this question, for each predictor X fit a model of the form:

```
Y = \beta_0 + \beta_1 X + \beta_2 X^2 + \beta_3 X^3 + \epsilon
```

```
#Non-linear regression models
nlm_crim <- lm(medv~crim+I(crim^2)+I(crim^3), data = boston_data)</pre>
summary(nlm_crim)
##
## Call:
## lm(formula = medv ~ crim + I(crim^2) + I(crim^3), data = boston_data)
##
## Residuals:
##
      Min
               1Q Median
                              3Q
                                     Max
## -17.983 -4.975 -1.940
                           2.881 33.391
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.519e+01 4.355e-01 57.846 < 2e-16 ***
## crim
              -1.136e+00 1.444e-01 -7.868 2.24e-14 ***
              2.378e-02 6.808e-03 3.494 0.000518 ***
## I(crim^2)
## I(crim^3) -1.489e-04 6.641e-05 -2.242 0.025411 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.159 on 502 degrees of freedom
## Multiple R-squared: 0.2177, Adjusted R-squared: 0.213
## F-statistic: 46.57 on 3 and 502 DF, p-value: < 2.2e-16
nlm_zn \leftarrow lm(medv~zn+I(zn^2)+I(zn^3), data = boston_data)
summary(nlm_zn)
##
## Call:
## lm(formula = medv ~ zn + I(zn^2) + I(zn^3), data = boston_data)
## Residuals:
      Min
               1Q Median
                              3Q
## -15.449 -5.549 -1.049
                           3.225 29.551
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 20.4485972 0.4359536 46.905 < 2e-16 ***
              0.6433652 0.1105611
                                   5.819 1.06e-08 ***
## zn
## I(zn^2)
              ## I(zn^3)
              ## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.43 on 502 degrees of freedom
## Multiple R-squared: 0.1649, Adjusted R-squared: 0.1599
## F-statistic: 33.05 on 3 and 502 DF, p-value: < 2.2e-16
nlm_indus <- lm(medv~indus+I(indus^2)+I(indus^3), data = boston_data)</pre>
summary(nlm_indus)
```

```
## Call:
## lm(formula = medv ~ indus + I(indus^2) + I(indus^3), data = boston_data)
## Residuals:
      Min
               1Q Median
                               3Q
                                       Max
## -15.760 -4.725 -1.009
                             2.932 32.038
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 37.080160 1.663326 22.293 < 2e-16 ***
              -2.806994
                          0.509349 -5.511 5.71e-08 ***
## I(indus^2)
              0.140462
                          0.041554
                                    3.380 0.000781 ***
## I(indus<sup>3</sup>) -0.002399
                         0.001011 -2.373 0.018026 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 7.844 on 502 degrees of freedom
## Multiple R-squared: 0.2768, Adjusted R-squared: 0.2725
## F-statistic: 64.06 on 3 and 502 DF, p-value: < 2.2e-16
nlm_chas <- lm(medv~chas+I(chas^2)+I(chas^3), data = boston_data)</pre>
summary(nlm_chas)
##
## Call:
## lm(formula = medv ~ chas + I(chas^2) + I(chas^3), data = boston_data)
## Residuals:
      Min
               1Q Median
                                3Q
                                      Max
                             2.856 27.906
## -17.094 -5.894 -1.417
##
## Coefficients: (2 not defined because of singularities)
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 22.0938
                            0.4176 52.902 < 2e-16 ***
                                    3.996 7.39e-05 ***
## chas
                 6.3462
                            1.5880
## I(chas^2)
                                                 NA
                    NA
                                NA
                                       NA
## I(chas^3)
                    NA
                                NA
                                       NA
                                                 NA
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 9.064 on 504 degrees of freedom
## Multiple R-squared: 0.03072,
                                   Adjusted R-squared: 0.02879
## F-statistic: 15.97 on 1 and 504 DF, p-value: 7.391e-05
nlm_nox <- lm(medv~nox+I(nox^2)+I(nox^3), data = boston_data)</pre>
summary(nlm_nox)
##
## Call:
## lm(formula = medv ~ nox + I(nox^2) + I(nox^3), data = boston_data)
##
## Residuals:
##
      Min
               1Q Median
                                3Q
                                       Max
## -13.104 -5.020 -2.144
                             2.747 32.416
##
```

```
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -22.49
                            38.52 -0.584
                                            0.5596
                           195.10 1.615
                                            0.1069
## nox
                315.10
## I(nox^2)
               -615.83
                           320.48 -1.922
                                            0.0552 .
## I(nox^3)
                350.19
                           170.92
                                    2.049
                                            0.0410 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.282 on 502 degrees of freedom
## Multiple R-squared: 0.1939, Adjusted R-squared: 0.189
## F-statistic: 40.24 on 3 and 502 DF, p-value: < 2.2e-16
nlm_rm <- lm(medv~rm+I(rm^2)+I(rm^3), data = boston_data)</pre>
summary(nlm_rm)
##
## Call:
## lm(formula = medv ~ rm + I(rm^2) + I(rm^3), data = boston_data)
##
## Residuals:
               1Q Median
      Min
                               3Q
                                      Max
## -29.102 -2.674 0.569
                            3.011 35.911
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 241.3108
                           47.3275 5.099 4.85e-07 ***
## rm
              -109.3906
                           22.9690 -4.763 2.51e-06 ***
## I(rm^2)
                16.4910
                            3.6750 4.487 8.95e-06 ***
## I(rm^3)
                -0.7404
                            0.1935 -3.827 0.000146 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 6.11 on 502 degrees of freedom
## Multiple R-squared: 0.5612, Adjusted R-squared: 0.5586
## F-statistic: 214 on 3 and 502 DF, p-value: < 2.2e-16
nlm_age <- lm(medv~age+I(age^2)+I(age^3), data = boston_data)</pre>
summary(nlm_age)
##
## Call:
## lm(formula = medv ~ age + I(age^2) + I(age^3), data = boston_data)
##
## Residuals:
##
      Min
               1Q Median
                               30
                                      Max
## -16.443 -4.909 -2.234
                            2.185 32.944
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.893e+01 2.992e+00
                                     9.668
                                              <2e-16 ***
## age
              -1.224e-01 2.014e-01 -0.608
                                               0.544
                                     0.599
                                               0.549
## I(age^2)
               2.355e-03 3.930e-03
              -2.318e-05 2.279e-05 -1.017
## I(age^3)
                                             0.310
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.472 on 502 degrees of freedom
## Multiple R-squared: 0.1566, Adjusted R-squared: 0.1515
## F-statistic: 31.06 on 3 and 502 DF, p-value: < 2.2e-16
nlm_dis <- lm(medv~dis+I(dis^2)+I(dis^3), data = boston_data)</pre>
summary(nlm_dis)
##
## Call:
## lm(formula = medv ~ dis + I(dis^2) + I(dis^3), data = boston_data)
##
## Residuals:
      Min
               1Q Median
                              3Q
                                    Max
## -12.571 -5.242 -2.037
                           2.397 34.769
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 7.03789
                         2.91134 2.417 0.01599 *
                         2.06633 4.158 3.77e-05 ***
## dis
              8.59284
                         0.41235 -3.030 0.00257 **
## I(dis^2)
              -1.24953
## I(dis^3)
              0.05602
                         0.02428
                                  2.307 0.02146 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.727 on 502 degrees of freedom
## Multiple R-squared: 0.105, Adjusted R-squared: 0.09968
## F-statistic: 19.64 on 3 and 502 DF, p-value: 4.736e-12
nlm_rad <- lm(medv~rad+I(rad^2)+I(rad^3), data = boston_data)</pre>
summary(nlm rad)
##
## lm(formula = medv ~ rad + I(rad^2) + I(rad^3), data = boston_data)
##
## Residuals:
      Min
               1Q Median
                              3Q
                                    Max
## -16.630 -5.151 -2.017
                           3.169 33.594
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 30.251303 2.567860 11.781 < 2e-16 ***
## rad
             -3.799454 1.307156 -2.907 0.003815 **
## I(rad^2)
              ## I(rad^3)
             ## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
\#\# Residual standard error: 8.37 on 502 degrees of freedom
## Multiple R-squared: 0.1767, Adjusted R-squared: 0.1718
## F-statistic: 35.91 on 3 and 502 DF, p-value: < 2.2e-16
nlm_tax <- lm(medv~tax+I(tax^2)+I(tax^3), data = boston_data)</pre>
summary(nlm_tax)
```

```
##
## Call:
## lm(formula = medv ~ tax + I(tax^2) + I(tax^3), data = boston_data)
## Residuals:
##
      Min
               1Q Median
                               3Q
                                      Max
## -15.109 -4.952 -1.878
                            2.957 33.694
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 5.222e+01 1.397e+01
                                     3.739 0.000206 ***
              -1.635e-01 1.133e-01 -1.443 0.149646
## tax
## I(tax^2)
                                     1.055 0.292004
               3.029e-04 2.872e-04
## I(tax^3)
              -2.079e-07 2.236e-07 -0.930 0.353061
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 8.115 on 502 degrees of freedom
## Multiple R-squared: 0.2261, Adjusted R-squared: 0.2215
## F-statistic: 48.89 on 3 and 502 DF, p-value: < 2.2e-16
nlm_ptratio <- lm(medv~ptratio+I(ptratio^2)+I(ptratio^3), data = boston_data)</pre>
summary(nlm_ptratio)
##
## Call:
## lm(formula = medv ~ ptratio + I(ptratio^2) + I(ptratio^3), data = boston_data)
## Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
## -17.7795 -5.0364 -0.9778
                               3.4766 31.1636
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 312.28642 152.48693
                                     2.048
                                              0.0411 *
               -48.69114
                           26.88441 -1.811
                                              0.0707 .
## ptratio
                                              0.0700 .
## I(ptratio^2)
                 2.83995
                            1.56413
                                     1.816
                            0.03005 -1.892
                                              0.0590 .
## I(ptratio^3) -0.05686
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 7.898 on 502 degrees of freedom
## Multiple R-squared: 0.2669, Adjusted R-squared: 0.2625
## F-statistic: 60.91 on 3 and 502 DF, p-value: < 2.2e-16
nlm_black <- lm(medv~black+I(black^2)+I(black^3), data = boston_data)</pre>
summary(nlm black)
##
## lm(formula = medv ~ black + I(black^2) + I(black^3), data = boston_data)
##
## Residuals:
      Min
               10 Median
                               3Q
                                      Max
## -19.005 -4.802 -1.613
                            2.852 28.051
```

```
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 1.260e+01 2.517e+00
                                       5.006
                                              7.7e-07 ***
## black
               -1.703e-02
                           6.150e-02
                                      -0.277
                                                0.782
                2.036e-04
                                       0.625
                                                0.532
## I(black^2)
                           3.258e-04
## I(black^3)
              -2.224e-07
                           4.765e-07
                                      -0.467
                                                0.641
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.685 on 502 degrees of freedom
## Multiple R-squared: 0.1135, Adjusted R-squared: 0.1082
## F-statistic: 21.43 on 3 and 502 DF, p-value: 4.463e-13
nlm_lstat <- lm(medv~lstat+I(lstat^2)+I(lstat^3), data = boston_data)</pre>
summary(nlm_lstat)
##
## Call:
## lm(formula = medv ~ lstat + I(lstat^2) + I(lstat^3), data = boston_data)
##
## Residuals:
##
       Min
                  10
                       Median
                                    30
                                            Max
## -14.5441 -3.7122 -0.5145
                                2.4846
                                        26.4153
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) 48.6496253
                          1.4347240
                                     33.909
               -3.8655928
## 1stat
                           0.3287861 -11.757
                                              < 2e-16 ***
## I(lstat^2)
                0.1487385
                           0.0212987
                                       6.983 9.18e-12 ***
              -0.0020039
                                     -5.013 7.43e-07 ***
## I(lstat^3)
                           0.0003997
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 5.396 on 502 degrees of freedom
## Multiple R-squared: 0.6578, Adjusted R-squared: 0.6558
## F-statistic: 321.7 on 3 and 502 DF, p-value: < 2.2e-16
```

On analyzing the summary statistics, it is observed that most of the predictors have the cubical coefficients which have significant p-values. Hence, the model cannot ignore these coefficients. Thus, if we retain the cubical coefficients then we have to keep the lower exponential coefficients. Thus, this indictaes that there is non-linear reationship between the reposne that is median values of homes and other predictors. Though there are some predictors for which the quadratic and cubic coefficients are not very significant as their p-value is much higher. Thus, we cannot consider predictors like age, tax, and black to have even a non-linear relationship with the response.

7. Consider performing a stepwise model selection procedure to determine the bets fit model. Discuss your results. How is this model different from the model in (4)?

```
#Stepwise model selection
step_model <- stepAIC(lm_multiple, direction = "both")

## Start: AIC=1589.64

## medv ~ crim + zn + indus + chas + nox + rm + rad + dis + tax +

## ptratio + lstat + black + age
##</pre>
```

```
Df Sum of Sq RSS AIC
          1 0.06 11079 1587.7
## - age
                 2.52 11081 1587.8
## - indus 1
## <none>
                     11079 1589.6
         1
                218.97 11298 1597.5
## - chas
## - tax
          1 242.26 11321 1598.6
## - crim
           1 243.22 11322 1598.6
## - zn
          1 257.49 11336 1599.3
            1 270.63 11349 1599.8
## - black
## - rad 1 479.15 11558 1609.1
## - nox
            1 487.16 11566 1609.4
## - ptratio 1 1194.23 12273 1639.4
            1 1232.41 12311 1641.0
## - dis
## - rm
           1 1871.32 12950 1666.6
## - lstat
          1 2410.84 13490 1687.3
##
## Step: AIC=1587.65
## medv ~ crim + zn + indus + chas + nox + rm + rad + dis + tax +
     ptratio + lstat + black
##
           Df Sum of Sq RSS
                              AIC
## - indus 1 2.52 11081 1585.8
                      11079 1587.7
## <none>
                0.06 11079 1589.6
          1
## + age
           1 219.91 11299 1595.6
## - chas
## - tax
           1
                242.24 11321 1596.6
## - crim
            1
                243.20 11322 1596.6
## - zn
           1
              260.32 11339 1597.4
## - black
          1 272.26 11351 1597.9
## - rad
            1 481.09 11560 1607.2
               520.87 11600 1608.9
## - nox
            1
## - ptratio 1 1200.23 12279 1637.7
## - dis
         1 1352.26 12431 1643.9
## - rm
            1 1959.55 13038 1668.0
            1 2718.88 13798 1696.7
## - lstat
## Step: AIC=1585.76
## medv ~ crim + zn + chas + nox + rm + rad + dis + tax + ptratio +
## lstat + black
##
##
           Df Sum of Sq RSS AIC
## <none>
                     11081 1585.8
## + indus
          1
                 2.52 11079 1587.7
## + age
          1
                 0.06 11081 1587.8
## - chas
           1 227.21 11309 1594.0
## - crim
                 245.37 11327 1594.8
            1
## - zn
                257.82 11339 1595.4
            1
## - black
                270.82 11352 1596.0
          1
                273.62 11355 1596.1
## - tax
            1
## - rad
               500.92 11582 1606.1
            1
## - nox
            1 541.91 11623 1607.9
## - ptratio 1 1206.45 12288 1636.0
## - dis 1 1448.94 12530 1645.9
          1 1963.66 13045 1666.3
## - rm
```

```
step_model$anova
## Stepwise Model Path
## Analysis of Deviance Table
##
## Initial Model:
## medv ~ crim + zn + indus + chas + nox + rm + rad + dis + tax +
##
       ptratio + lstat + black + age
##
## Final Model:
## medv ~ crim + zn + chas + nox + rm + rad + dis + tax + ptratio +
##
       lstat + black
##
##
##
                  Deviance Resid. Df Resid. Dev
                                                      AIC
## 1
                                  492
                                        11078.78 1589.643
       - age 1 0.06183435
                                        11078.85 1587.646
                                  493
             1 2.51754013
## 3 - indus
                                  494
                                        11081.36 1585.761
```

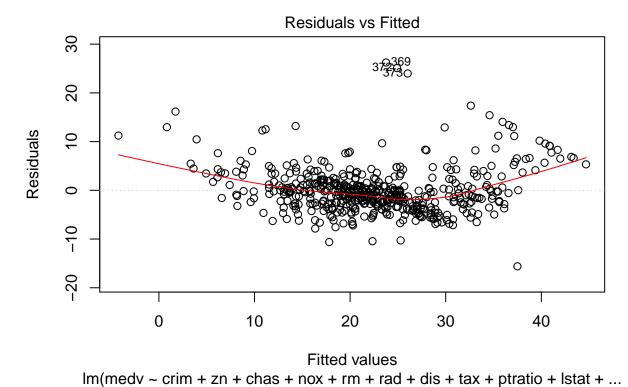
2723.48 13805 1695.0

- 1stat

From the final step of stepwise model selection, does not consider the variables indus and age to generate best fir multiple regression model. This is because we start with all variables and the AIC value is 1589.64. Then we try removing age and the AIC value decreases to 1587. Then we try eliminating indus and the value increases a bit. Hence, in the next step we start with the best least AIC value got so far which is 1587.65 and eliminate age from this step. Now, when we eliminate indus, the value of AIC goes down further to 1585. Also, in the same step when we try to add indus it increases. So, in this step we got the least AIC of 1585. In the final step, we start with this AIC value obtained and try to add the removed values again. BUt, it is observed that AIC values goes on increasing. Hence, the best fit model has the least AIC when we eliminate age and indus.

8. Evaluate the statistical assumptions in your regression analysis from (7) by performing a basic analysis of model residuals and any unusual observations. Discuss any concerns you have about your model.

```
#Plot of residuals vs. fitted for the stepwise model
plot(step_model, which = c(1))
```



The residuals vs. fitted values graph shows that there is a non-linear relationship between the response and the predictors. This is because the regression line is not coinciding with the 0 line in the above plot. This means the fitted values are not randomly bouncing around the 0-line. Also, there are many outliers in the

graph which do not have a corresponding residual value to nullify the effect of error introduced in the model.