HW 3

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Exercise 1

```
library(ISLR2)
## Warning: package 'ISLR2' was built under R version 4.1.3
data("Auto")
library(ggplot2)
(a)
# Perform simple linear regression
model <- lm(mpg ~ horsepower, data = Auto)</pre>
# Print the summary of the regression results
summary(model)
##
## Call:
## lm(formula = mpg ~ horsepower, data = Auto)
## Residuals:
                1Q Median
       Min
                                   3Q
## -13.5710 -3.2592 -0.3435 2.7630 16.9240
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 39.935861 0.717499 55.66 <2e-16 ***
## horsepower -0.157845 0.006446 -24.49 <2e-16 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 4.906 on 390 degrees of freedom
## Multiple R-squared: 0.6059, Adjusted R-squared: 0.6049
## F-statistic: 599.7 on 1 and 390 DF, p-value: < 2.2e-16
```

- i. Yes, there is a relationship between the predictor "horsepower" and the response "mpg" given the p-value is "<2e-16"
- ii. The multiple R-squared value is 0.6059 thus this means that about 60.59% of the variance in "mpg" can be explained by the "horsepower" predictor. This indicates a moderate to strong relationship between the two.
- iii. The coefficient for "horsepower" is -0.157845. Since this is less than zero, it indicates a negative relationship between "horsepower" and "mpg" or in otherwords as "horsepower" increases by 1, "mpg" should decrease by about -0.157845.

(b)

```
new_data <- data.frame(horsepower = 98)

# Use the 'predict' function to make the prediction
predicted_mpg <- predict(model, newdata = new_data)

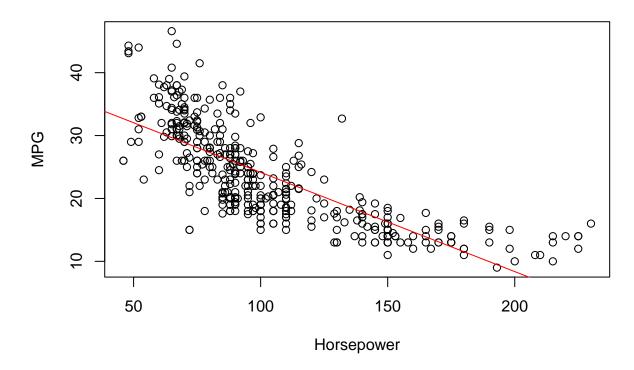
# Print the predicted 'mpg' value
print(predicted_mpg)

## 1
## 24.46708</pre>
```

(c)

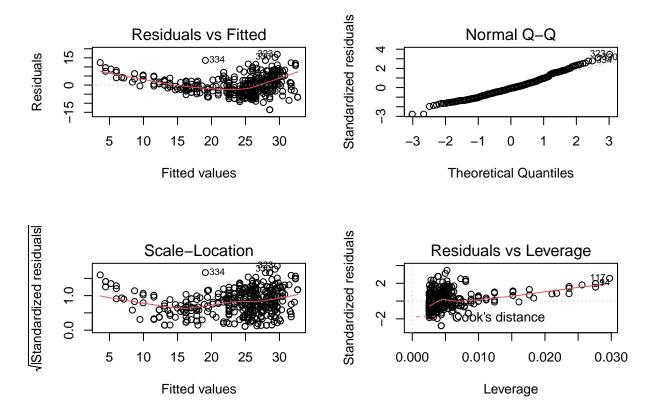
```
# Create a scatterplot of mpg vs. horsepower
plot(Auto$horsepower, Auto$mpg, xlab = "Horsepower", ylab = "MPG", main = "Scatterplot of MPG vs. Horse
# Add the least squares regression line
abline(model, col = "red")
```

Scatterplot of MPG vs. Horsepower



(d)

Create diagnostic plots for the linear regression model
par(mfrow = c(2, 2)) # Set up a 2x2 grid for the plots
plot(model)



Looking at the residuals vs fitted graph, there is a slight curve to the graph thus meaning a more complex model might be better for the data.

Exersize 2

```
set.seed(1)

(a)

X <- rnorm(100, mean = 0, sd = 1)

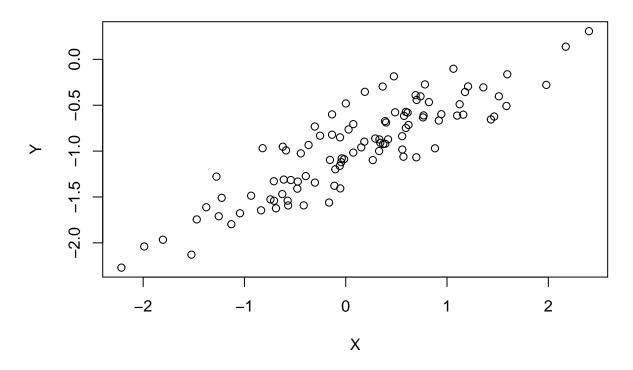
(b)

epsilon <- rnorm(100, mean = 0, sd = 0.25)</pre>
```

(c)

```
# Calculate Y based on the linear model
Y \leftarrow -1 + 0.5 * X + epsilon
# (i) Length of vector Y
length_Y <- length(Y)</pre>
# (ii) Values of beta
beta_0 <- -1
beta_1 <- 0.5
 i.
length_Y
## [1] 100
  ii.
beta_0
## [1] -1
beta_1
## [1] 0.5
(d)
# Create a scatterplot of X vs. Y
plot(X, Y, main = "Scatterplot of X vs. Y", xlab = "X", ylab = "Y")
```

Scatterplot of X vs. Y



Based off the plot above, there seems to be a positive real ationship between X and Y, and the pattern is rather linear.

(e)

```
\# Fit a least squares linear model
model \leftarrow lm(Y \sim X)
# Print the summary of the model
summary(model)
##
## Call:
## lm(formula = Y ~ X)
##
## Residuals:
##
        Min
                  1Q
                       Median
                                              Max
##
   -0.46921 -0.15344 -0.03487
                               0.13485
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.00942
                            0.02425
                                     -41.63
                                               <2e-16 ***
## X
                                      18.56
                0.49973
                            0.02693
                                               <2e-16 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

```
##
## Residual standard error: 0.2407 on 98 degrees of freedom
## Multiple R-squared: 0.7784, Adjusted R-squared: 0.7762
## F-statistic: 344.3 on 1 and 98 DF, p-value: < 2.2e-16</pre>
```

The actual beta 0 was estimated at about -0.9745 while the actual beta 1 was estimated at about 0.5213. This is extremely close to the predicted beta values from earlier. Another thing to note from this model is that p-value for the relationship between x and y is extremely small thus meaning significance.