

group6_project630

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1) Load the data and check for outliers

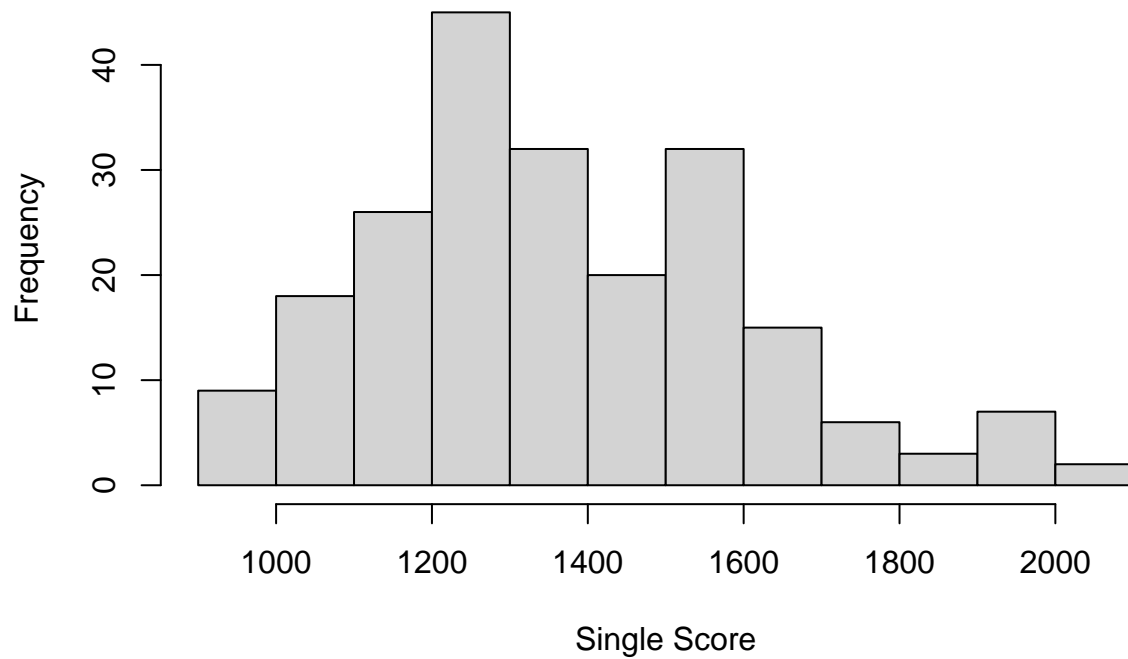
```
project <- read.csv("/Users/atishpanday/Downloads/CPU_r23_v2.csv")
project$manufacturer <- as.factor(project$manufacturer)
summary(project)
```

```
## manufacturer   cpuName          singleScore    multiScore
## AMD   : 86      Length:215      Min.    : 903    Min.    : 5423
## Apple:  4      Class :character  1st Qu.:1207    1st Qu.: 8186
## Intel:125      Mode  :character  Median :1312    Median :10890
##                                     Mean   :1368    Mean   :12980
##                                     3rd Qu.:1534    3rd Qu.:14394
##                                     Max.    :2082    Max.    :75671
##      cores      threads      baseClock      turboClock
## Min.    : 4.000    Min.    : 6.00    Min.    :1.100    Min.    :3.200
## 1st Qu.: 6.000    1st Qu.: 12.00    1st Qu.:2.500    1st Qu.:4.200
## Median : 8.000    Median : 16.00    Median :3.200    Median :4.500
## Mean    : 9.367    Mean    : 17.74    Mean    :3.054    Mean    :4.515
## 3rd Qu.: 9.000    3rd Qu.: 16.00    3rd Qu.:3.600    3rd Qu.:4.800
## Max.    :64.000    Max.    :128.00    Max.    :4.200    Max.    :5.500
##      type
## Length:215
## Class :character
## Mode  :character
##
##
##
```

2) Linear regression model

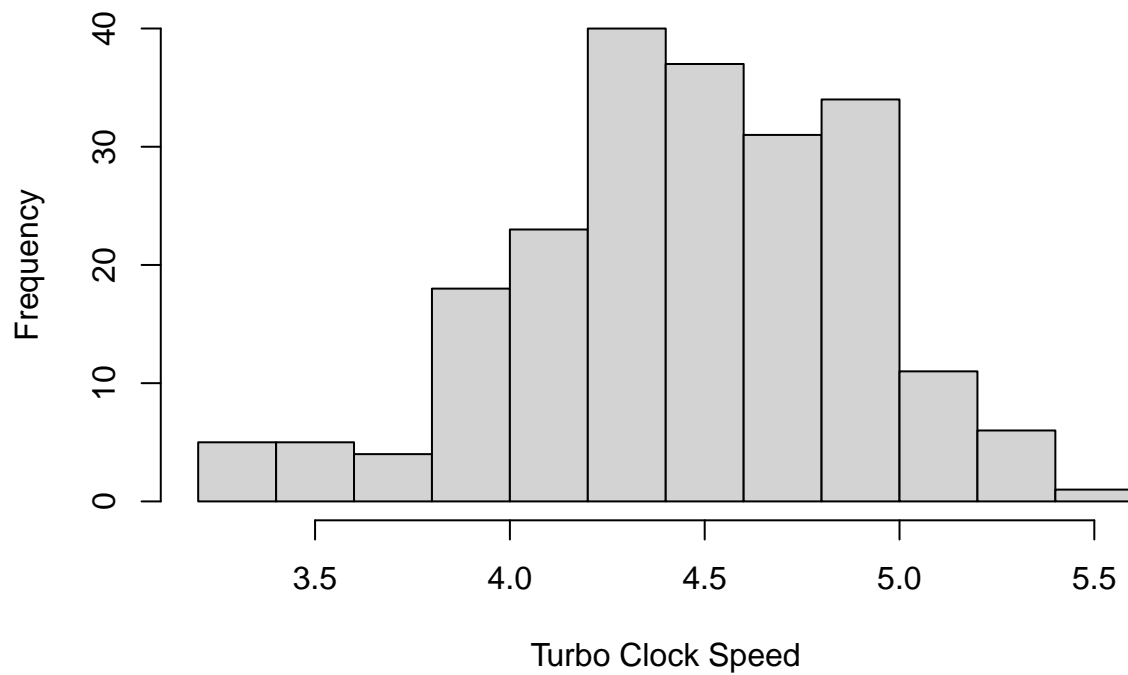
```
mod <- lm(singleScore ~ turboClock, data = project)
hist(project$singleScore, xlab = "Single Score", ylab = "Frequency", main = "Histogram of Single Score")
```

Histogram of Single Score



```
hist(project$turboClock, xlab = "Turbo Clock Speed", ylab = "Frequency", main = "Histogram of Turbo Clock Speed")
```

Histogram of Turbo Clock



3) Check Conditions

Linearity check

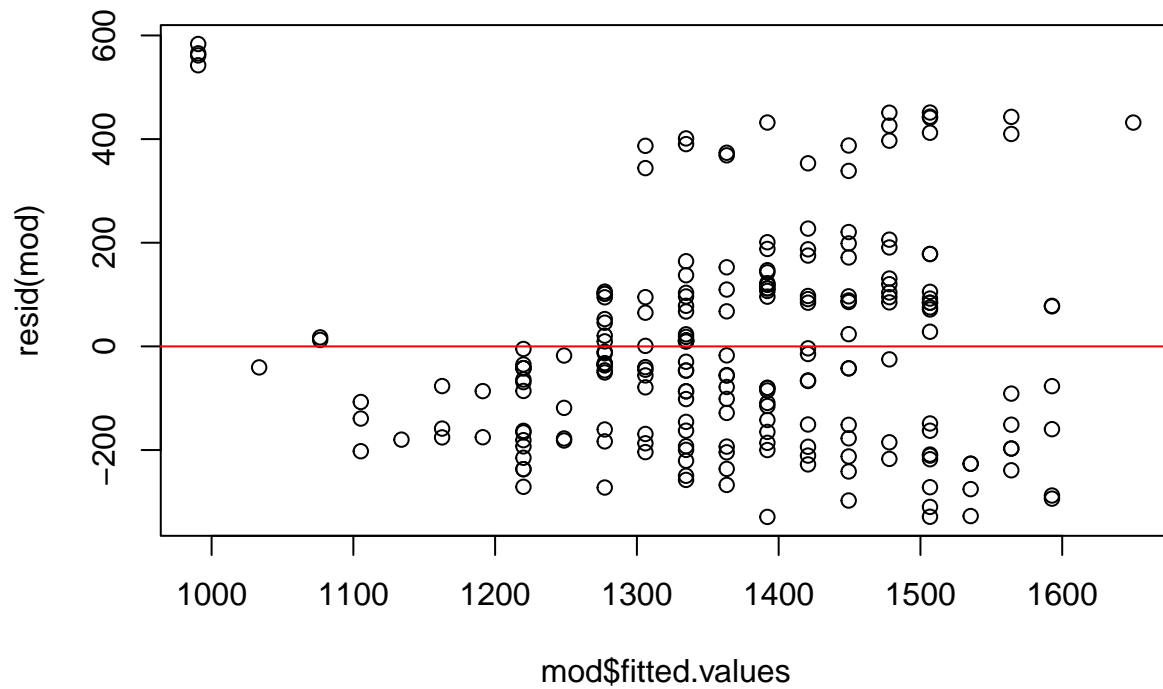
```
library(ggplot2)
ggplot(project, aes(x = turboClock, y = singleScore)) +
  geom_point() +
  geom_smooth(method = "lm",
              se = FALSE,
              col = "red") +
  ggtitle(" Performance vs Boost speed") +
  labs(x = "Boost speed", y = "Performance")
```

```
## `geom_smooth()` using formula 'y ~ x'
```



Constant variance check

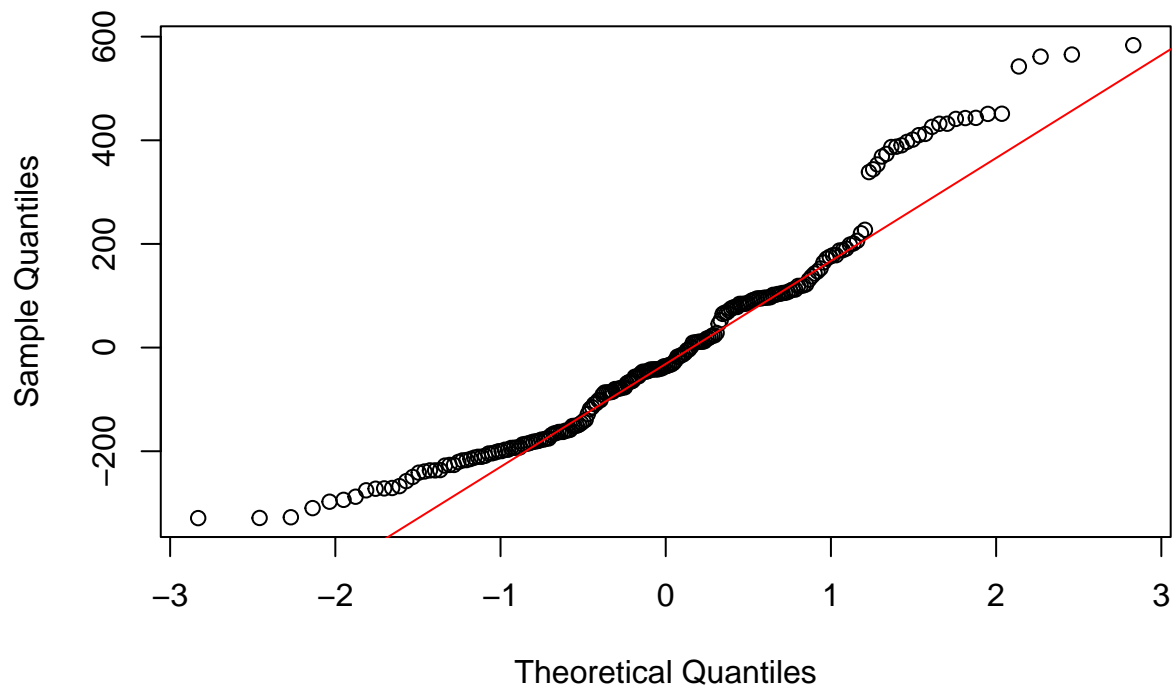
```
mod <- lm(singleScore ~ turboClock, data = project)
plot(resid(mod) ~ mod$fitted.values)
abline(h = 0, col = "red")
```



Normality check

```
qqnorm(resid(mod))
qqline(resid(mod), col = "red")
```

Normal Q-Q Plot



Secondary Discovery

```
ggplot(project, aes(x = turboClock, y = singleScore, col = manufacturer)) +
  geom_point() +
```

```
ggtitle(" Performance vs Boost speed") +
labs(x = "Boost speed", y = "Performance") + scale_color_manual(breaks = c("AMD", "Apple", "Intel"),v
```

