

# Stat 341 Assignment 1

2022-09-26

## Question 1: Basic R Calculations

1a)

```
3^4
```

```
## [1] 81
```

1b)

```
log(100, base = 7) # 1.b)
```

```
## [1] 2.366589
```

1c)

```
x <- seq(1, 100)
sum(sapply(x, function(x) {1/(x^2)}))
```

```
## [1] 1.634984
```

1d)

```
100 %% 7
```

```
## [1] 2
```

1e)

```
dx_steps <- 0.001
x_val <- seq(0, pi/2, by = dx_steps)
sum(sapply(x_val, function(x){ sin(x) * dx_steps })))
```

```
## [1] 0.9997036
```

1f)

```
dx_steps <- 0.001
x_val <- seq(0, 3, by = dx_steps)
sum(sapply(x_val, function(x){ dexp(x, rate = 1/2) * dx_steps })))
```

```
## [1] 0.7771756
```

1g)

```
f <- function(x) {  
  return (x^2 + 3)  
}  
  
dx_steps <- 0.0001  
x_val <- seq(-2, 2, by = dx_steps)  
sum(sapply(x_val, function(x){ f(x) * dx_steps })))  
  
## [1] 17.33403
```

## Question 2: Comparing Spread Attributes

2a)

$$\begin{aligned}
 SD(\mathcal{P} + b) &= \sqrt{\frac{\sum_{u \in \mathcal{P} + b} (y_u - (\text{mean}(\mathcal{P} + b)))^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} ((y_u + b) - (\bar{y} + b))^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} ((y_u - \bar{y}) + b - b)^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} ((y_u - \bar{y}))^2}{N}} \\
 &= SD(\mathcal{P})
 \end{aligned}$$

Hence, Standard Deviation is location invariant.

$$\begin{aligned}
 a(\mathcal{P} + b) = MAD(\mathcal{P} + b) &= \text{median}_{u \in \mathcal{P} + b} \left| y_u - (\text{median}_{u \in \mathcal{P} + b} y_u) \right|. \\
 &= \text{median}_{u \in \mathcal{P}} \left| y_u + b - (\text{median}_{u \in \mathcal{P}} y_u + b) \right|. \\
 &= \text{median}_{u \in \mathcal{P}} \left| (y_u - \text{median}_{u \in \mathcal{P}} y_u) + b - b \right| \\
 &= MAD(\mathcal{P})
 \end{aligned}$$

Hence, Median Absolute Deviation is location invariant.

2b)

$$\begin{aligned}
 SD(\alpha \times \mathcal{P}) &= \sqrt{\frac{\sum_{u \in \alpha \times \mathcal{P}} (y_u - (\text{mean}(\alpha \times \mathcal{P})))^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} ((\alpha \times y_u) - (\alpha \times \bar{y}))^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} (\alpha \times (y_u - \bar{y}))^2}{N}} \\
 &= \sqrt{\frac{\sum_{u \in \mathcal{P}} \alpha^2 \times ((y_u - \bar{y}))^2}{N}} \\
 &= \alpha \times SD(\mathcal{P})
 \end{aligned}$$

Hence, Standard Deviation is scale equivariant.

$$\begin{aligned}
a(\alpha \times \mathcal{P}) &= MAD(\alpha \times \mathcal{P}) = \operatorname{median}_{u \in \alpha \times \mathcal{P}} \left| y_u - \left( \operatorname{median}_{u \in \alpha \times \mathcal{P}} y_u \right) \right| \\
&= \operatorname{median}_{u \in \mathcal{P}} \left| (\alpha \times y_u - \operatorname{median}_{u \in \mathcal{P}} \alpha \times y_u) \right| \\
&= \operatorname{median}_{u \in \mathcal{P}} \left| (\alpha \times (y_u - \operatorname{median}_{u \in \mathcal{P}} y_u)) \right| \\
&= \alpha \times MAD(\mathcal{P})
\end{aligned}$$

Hence, Median Absolute Deviation is scale equivariant.

2c)

$$\begin{aligned}
SD(\mathcal{P}^k) &= \sqrt{\frac{\sum_{u \in \mathcal{P}^k} (y_u - (\operatorname{mean}(\mathcal{P}^k))^2}{N}} \\
&= \sqrt{\frac{\sum_{u \in \mathcal{P}} (y_u - \bar{y})^2}{N}} \\
&= SD(\mathcal{P})
\end{aligned}$$

Hence, Standard Deviation is replication invariant.

$$\begin{aligned}
MAD(\mathcal{P}^k) &= \operatorname{median}_{u \in \mathcal{P}^k} \left| y_u - \left( \operatorname{median}_{u \in \mathcal{P}^k} y_u \right) \right| \\
&= \operatorname{median}_{u \in \mathcal{P}} \left| (y_u - \operatorname{median}_{u \in \mathcal{P}} y_u) \right| \\
&= MAD(\mathcal{P})
\end{aligned}$$

Hence, Median Absolute Deviation is replication invariant.

2d)

```
SD <- function(y) {
  return (sqrt(sum((y - mean(y))^2 / length(y))))
}

MAD <- function(y) {
  return (median(abs(y - median(y))))
}
```

2e)

```
set.seed(341)
sc <- function(pop, y, attr){
  N <- length(pop) + 1
  sapply (y, function(y.new){ N*(attr(c(y.new, pop)) - attr(pop)) })
}
```

```

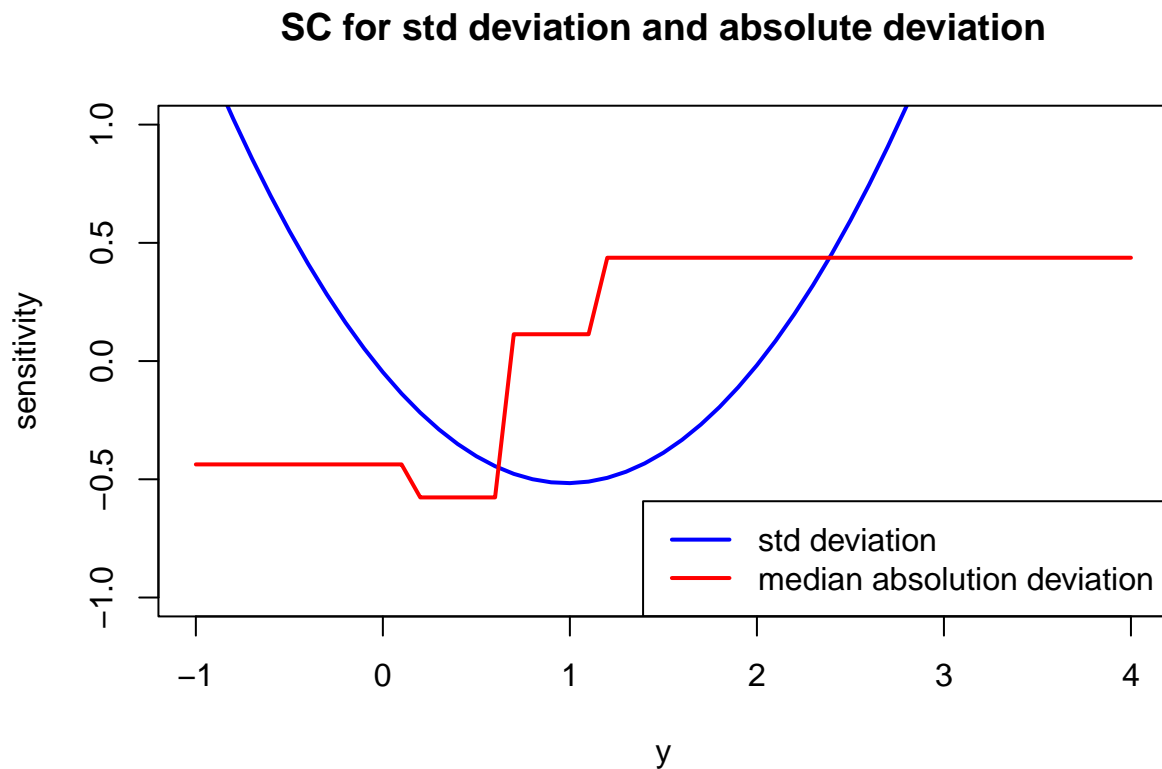
set.seed(341)
pop = rexp(1000)

y_val <- seq(-1, 4, by=0.1)

plot(y_val, sc(pop, y_val, SD), type="l", lwd = 2,
     main="SC for std deviation and absolute deviation", ylab="sensitivity", xlab="y",
     xlim=c(-1,4), ylim=c(-1, 1), col="blue")
lines(y_val, sc(pop, y_val, MAD), type="l", lwd = 2, main="Sensitivity curve for the median absolute deviation", col="red")

legend(x = "bottomright",          # Position
      legend = c("std deviation", "median absolute deviation"), # Legend texts
      col = c("blue", "red"),      # Line colors
      lwd = 2)

```



2f)

SD and MAD are both location invariant, scale equivariant, and replication invariant.

SD is however much more sensitive to extreme values since it's sensitivity curve increases without bounds as  $y \rightarrow \infty$  or  $y \rightarrow -\infty$ .

MAD is not very sensitive to extreme values and it's sensitivity will be bounded and hence, it is a much more robust measure because of its high breakdown point. On the other hand, SD is a fragile attribute and has a very low breakdown point.

It is advantageous to use MAD over SD when we want to limit the effect of outliers on the statistic. It is

advantageous to use SD over MAD to gain clarity over the range of variation within the dataset.

### Question 3: Write a rounded-barplot-making function

3a)

```
rounded.barplot <- function(x, xlab){
  table_x <- table(x)
  categories <- names(table_x)
  categories_frequencies <- as.numeric(table_x)

  plot.new()
  plot(NULL, type="n", xlim=c(0, 10*length(categories_frequencies)), ylim=c(0, max(categories_frequencies)))

  axis(2, at=seq(from=0, to=max(categories_frequencies), by=10))
  mtext(xlab, side=1, line=2)
  mtext("Frequency", side=2, line=3)

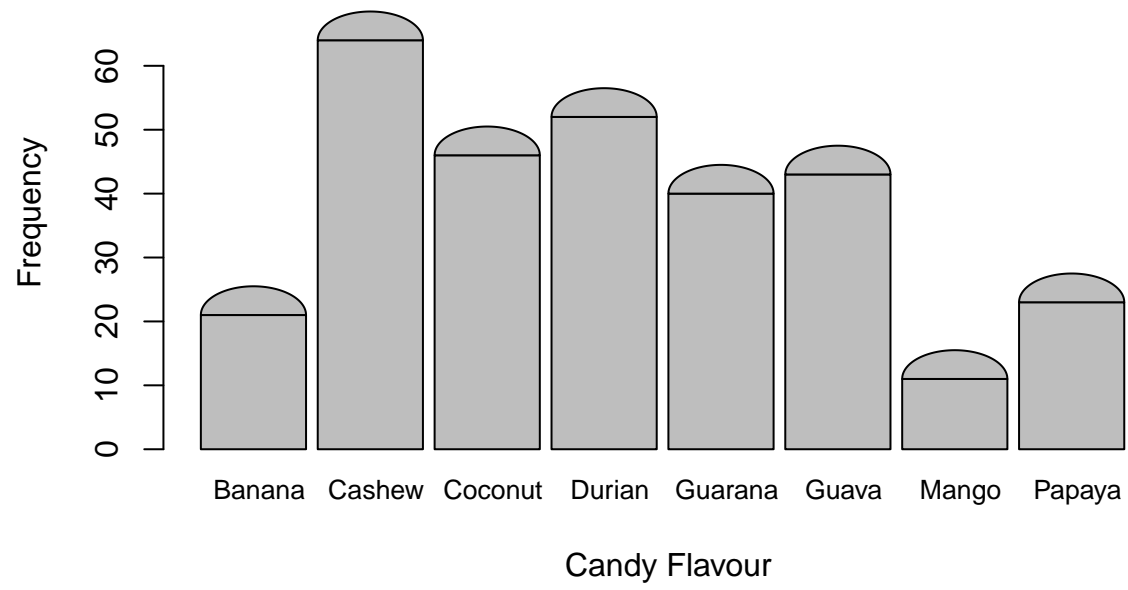
  x_semi <- seq(-4.5, 4.5, by=0.01)
  y_semi <- sqrt(20.25-x_semi^2)

  for (i in c(1: length(categories_frequencies))){
    rect(10*(i-1), 0, 10*i-1, categories_frequencies[i], col = "gray", border = "black")
    mtext(categories[i], 1, at=10*i-5, cex=0.85)
    polygon(x_semi + 4.5 + 10*(i-1), y_semi + categories_frequencies[i], col = "gray")
  }
}
```

3b)

```
set.seed(12345)
flavours = c("Mango", "Papaya", "Banana", "Coconut", "Guava", "Guarana", "Durian", "Cashew")
candies = sample(flavours, size=300, prob=(1:8)/sum(1:8), replace=TRUE)

rounded.barplot(candies, xlab="Candy Flavour")
```





## Question 4: R Analysis Question

4a)

```
setwd("C:/Users/2baja/OneDrive/Desktop/STAT 341/A1")
apartment_eval <- read.csv("Apartment_Building_Evaluation.csv")

score_90 <- apartment_eval[, "SCORE"] >= 90
sum(score_90)
```

```
## [1] 410
```

4b)

```
# 4.b)
davenport <- which(apartment_eval[, "WARDNAME"] == "Davenport")
davenport_apartments <- apartment_eval[davenport,]
davenport_apartments_sorted_addresses <- davenport_apartments[order(-davenport_apartments$SCORE), "SITE_",]
davenport_apartments_sorted_addresses[c(1:5)]
```

```
## [1] "1544 DUNDAS ST W" "1544 DUNDAS ST W" "1289 DUNDAS ST W"
## [4] "19-21 RUSHOLME RD" "410 DOVERCOURT RD"
```

4c)

Scarborough North has the highest score on average: 81.5. River-Black Creek has the lowest score on average: 68.79.

```
unique_wardnames <- unique(apartment_eval[, "WARDNAME"])
sapply(unique_wardnames, function(name) { mean(apartment_eval[which(apartment_eval$WARDNAME == name), "SCORE"]) })
```

|    |                        |                       |                          |
|----|------------------------|-----------------------|--------------------------|
| ## | Scarborough Southwest  | Eglinton-Lawrence     | Scarborough-Agincourt    |
| ## | 72.03354               | 72.17902              | 78.33333                 |
| ## | Beaches-East York      | Davenport             | Spadina-Fort York        |
| ## | 72.44581               | 68.86260              | 75.14400                 |
| ## | Toronto-Danforth       | Toronto Centre        | Toronto-St. Paul's       |
| ## | 73.21563               | 71.90877              | 73.62217                 |
| ## | University-Rosedale    | York South-Weston     | Humber River-Black Creek |
| ## | 71.81912               | 70.28017              | 68.79331                 |
| ## | Willowdale             | Scarborough-Guildwood | Scarborough Centre       |
| ## | 76.86667               | 72.28054              | 74.51587                 |
| ## | Etobicoke Centre       | Don Valley East       | York Centre              |
| ## | 72.14054               | 76.30913              | 71.53305                 |
| ## | Don Valley West        | Parkdale-High Park    | Etobicoke-Lakeshore      |
| ## | 76.69196               | 69.34385              | 71.47331                 |
| ## | Etobicoke North        | Scarborough North     | Don Valley North         |
| ## | 69.30645               | 81.50000              | 79.19310                 |
| ## | Scarborough-Rouge Park |                       |                          |
| ## | 75.05479               |                       |                          |

4d)

```
plot(apartment_eval$YEAR_BUILT, apartment_eval$SCORE, pch = 16, col=adjustcolor("black", alpha = 0.25),
unique_years <- unique(apartment_eval[, "YEAR_BUILT"])
```

```

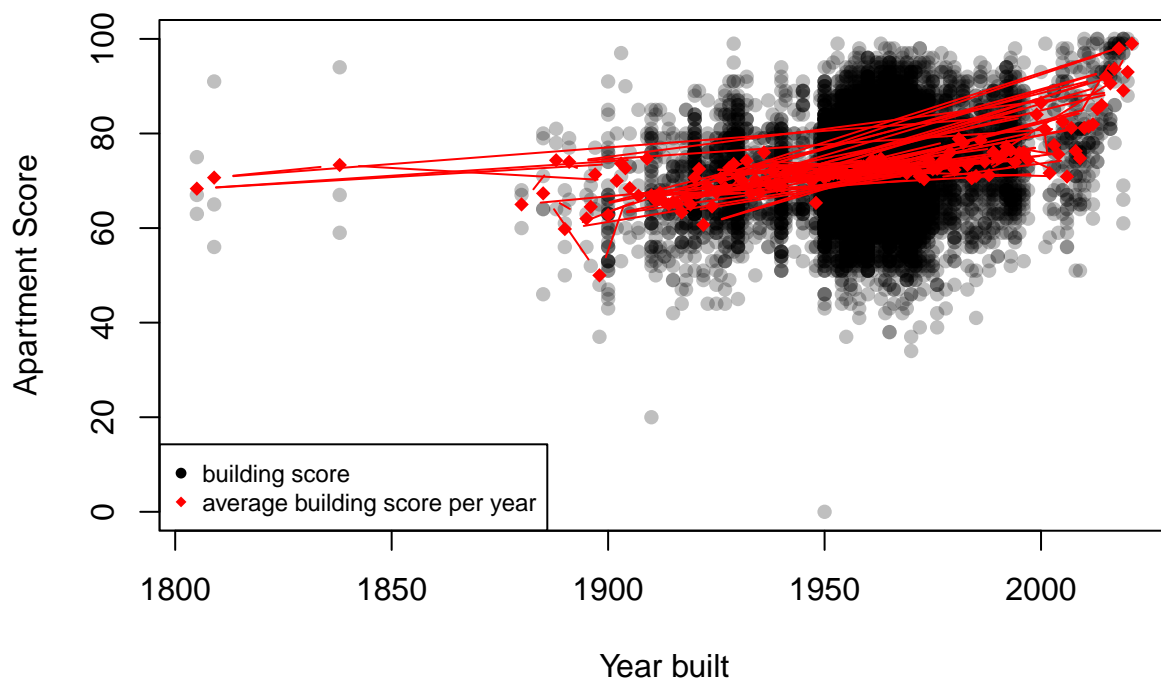
average_score_by_year <- sapply(unique_years, function(year_built) { mean(apartment_eval[which(apartmen

lines(unique_years, average_score_by_year, pch = 18, col="red", type="b")

legend(x = "bottomleft",          # Position
      legend = c("building score", "average building score per year"), # Legend texts
      col = c("black", "red"),    # Line colors
      cex = 0.75,
      pch = c(16, 18))

```

## Appartment scores for buildings vs the year they were built



4e)

```

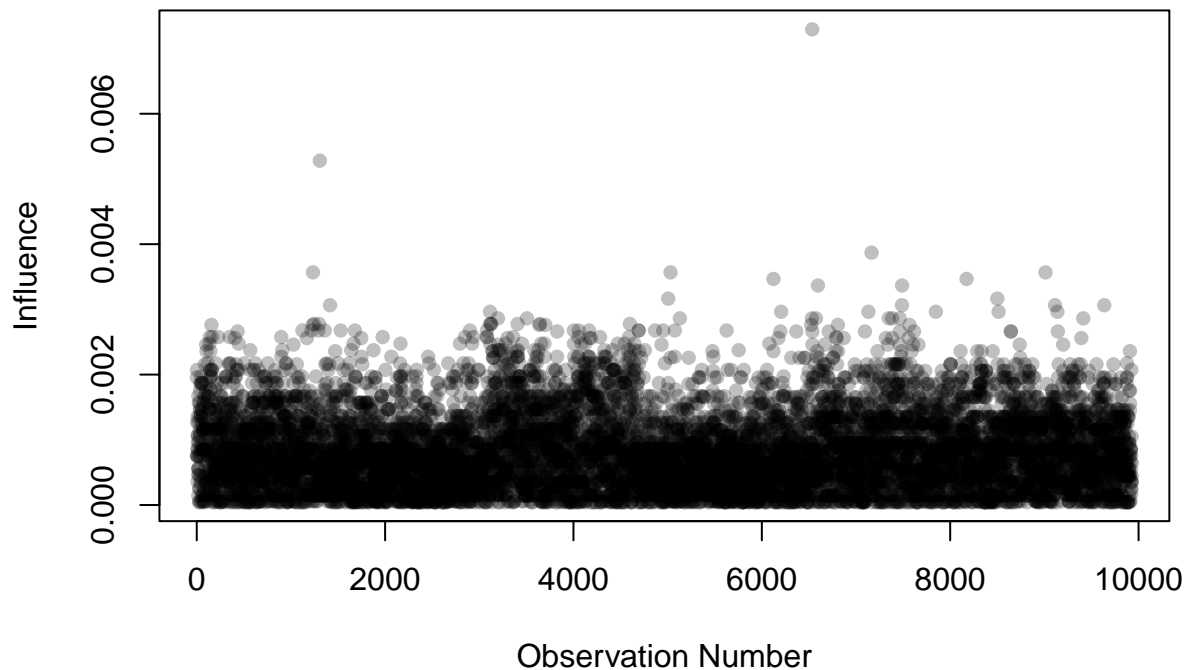
influence <- function(pop, attribute){
  N <- length(pop)
  attribute_total_pop <- attribute(pop)

  return (sapply(1:N, function(x) { abs(attribute_total_pop - attribute(pop[-x])) })))
}

plot(1:length(apartment_eval$SCORE), influence(apartment_eval$SCORE, mean), pch = 16, col=adjustcolor("

```

## Influence of apartment on mean apartment score



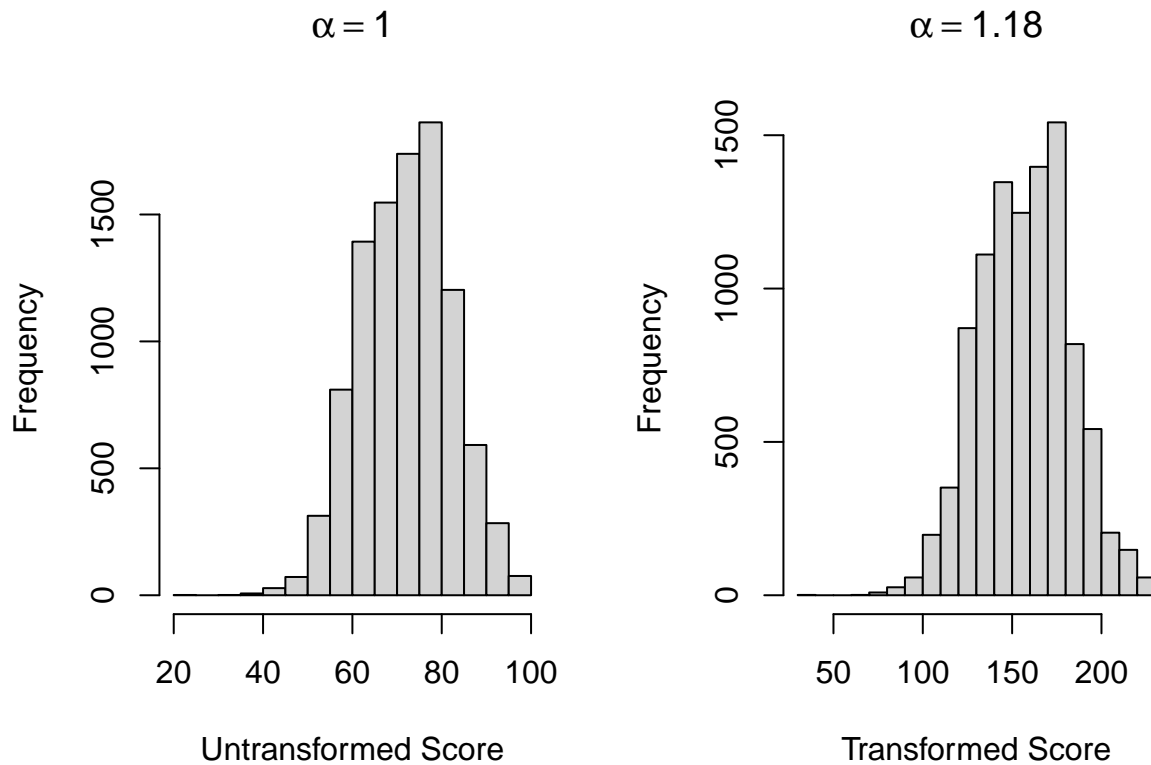
The building with the largest influence has the following observation number:

```
which.max(influence(apartment_eval$SCORE, mean))
```

```
## [1] 6535
```

4.f)

```
apartment_eval_without_outlier <- apartment_eval[-which.max(influence(apartment_eval$SCORE, mean)),]  
  
powerfun <- function(y, alpha) {  
  if(sum(y <= 0) > 0) stop("y must be positive")  
  if (alpha == 0)  
    log(y)  
  else if (alpha > 0) {  
    y^alpha  
  } else -(y^alpha)  
}  
  
par(mfrow=c(1,2))  
hist(powerfun(apartment_eval_without_outlier$SCORE, 1), main=bquote(alpha == .(1)), xlab="Untransformed",  
hist(powerfun(apartment_eval_without_outlier$SCORE, 1.18), main=bquote(alpha == .(1.18)), xlab="Transformed")
```



The  $\alpha$  that makes the SCORE distribution more symmetric is chosen as 1.18 since the transformed histogram has a skewness that is very close to 0.

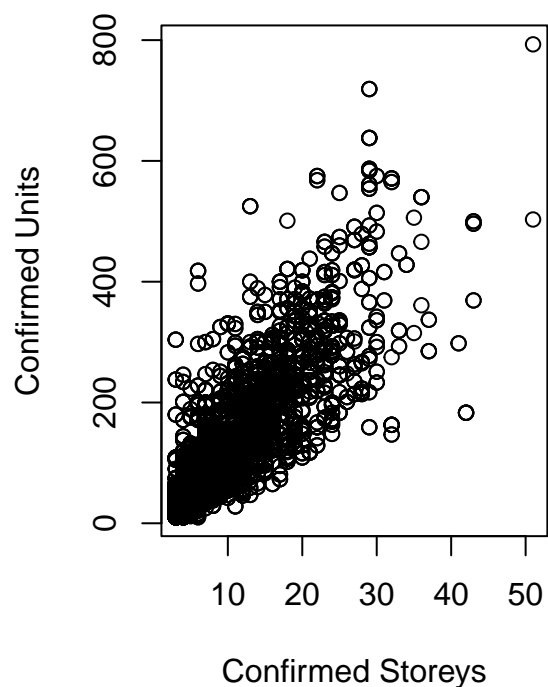
```
library("moments")
skewness(powerfun(apartment_eval_without_outlier$SCORE, 1.18))
```

```
## [1] 0.0006098378
```

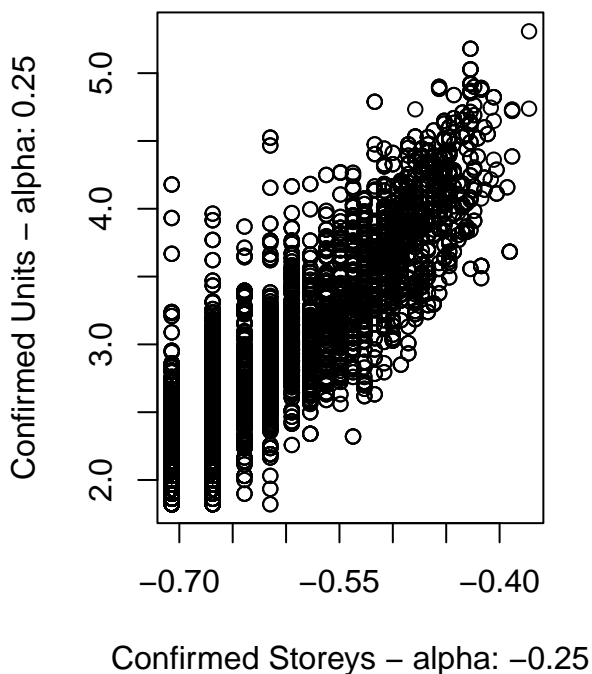
4.g)

```
par(mfrow=c(1,2))
plot(apartment_eval_without_outlier$CONFIRMED_STOREYS, apartment_eval_without_outlier$CONFIRMED_UNITS,
plot(powerfun(apartment_eval_without_outlier$CONFIRMED_STOREYS + 1, -0.25), powerfun(apartment_eval_wit
```

**Untransformed Units vs Storeys**



**Transformed Units vs Storeys**



$\alpha_x$  is chosen as  $-0.25$  and  $\alpha_y$  is chosen as  $0.25$ .

The corresponding linear regression model shows that transformed plot has an  $r^2 = 0.8003$  which depicts a closer linear relationship than the untransformed plot which has an  $r^2 = 0.7469$ .