```
R Homework Number 9
Fisher Ankney
November 13th, 2018
Statistics 5193
*Note: this document was created using R Markdown.
```

Question 1a

```
library(readxl)
StudentData <- read_excel("~/Documents/data_science/r_stat_5193/data/StudentData.xlsx")
boot.mean <- function(my.data, trim.percent = 0.05) {
   dat.sample <- sample(my.data, replace = T)
   mean(dat.sample, trim = trim.percent)
}</pre>
```

Question 1b

```
# the use of attach is not recommended due to the convoluted nature of the
# function, it reduces reproducibility, therefore I will reference the
# dataframe in the suggested code

set.seed(1)
boot.mean(StudentData$HSClass)

## [1] 309
boot.mean(StudentData$HSClass, trim.percent = .10)
```

[1] 344

Question 2a

```
my.boot.sample <- vector('double', length = 10000)

set.seed(1)
for (i in 1:10000) {
   my.boot.sample[i] <- boot.mean(StudentData$HSClass)
}

mean(my.boot.sample)

## [1] 310.1323

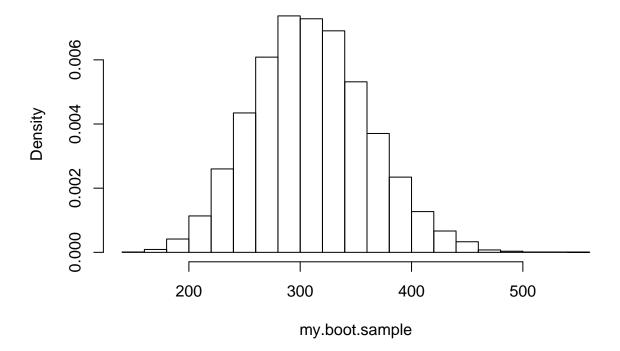
sd(my.boot.sample)

## [1] 51.98851</pre>
```

Question 2b

```
hist(my.boot.sample,
    probability = T,
    main = "Trimmed Mean Sampling Distribution")
```

Trimmed Mean Sampling Distribution

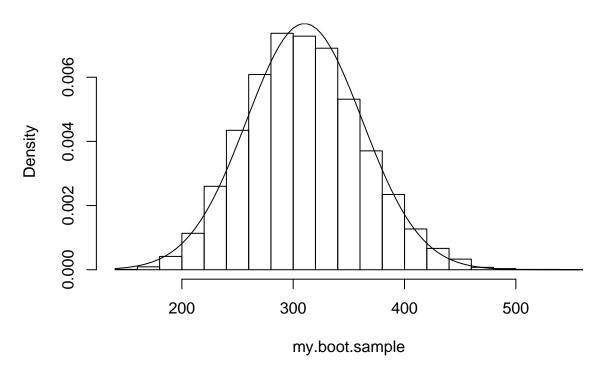


Question 2c

```
hist(my.boot.sample,
    probability = T,
    main = "Trimmed Mean Sampling Distribution")

curve(dnorm(x, mean = mean(my.boot.sample), sd = sd(my.boot.sample)),
    add = T)
```

Trimmed Mean Sampling Distribution



Question 2d

```
quantile(my.boot.sample, c(0.025, 0.975))
## 2.5% 97.5%
## 214.9068 417.5470
```

Question 3a

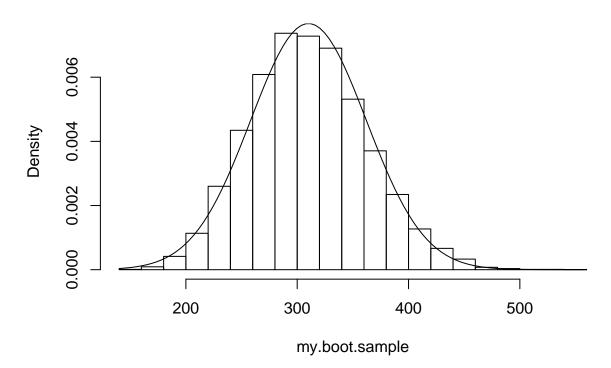
```
boot.ci <- function(my.data, trim.percent = 0.05, plot.it = T) {</pre>
# recreate boot.mean function
  boot.mean <- function(my.data, trim.percent = trim.percent) {</pre>
    dat.sample <- sample(my.data, replace = T)</pre>
    mean(dat.sample, trim = trim.percent)
  }
# create empty vector to store results
  my.boot.sample <- vector('double', length = 10000)</pre>
# loop
  for (i in 1:10000) {
    my.boot.sample[i] <- boot.mean(my.data, trim.percent = trim.percent)</pre>
# save mean and sd for plotting
data.mean <- mean(my.boot.sample)</pre>
data.sd <- sd(my.boot.sample)</pre>
# return confidence interval
  print(quantile(my.boot.sample, c(0.025, 0.975)))
# conditional plot.it
  if (plot.it == T) {
     hist(my.boot.sample,
           probability = T,
           main = "Trimmed Mean Sampling Distribution")
     curve(dnorm(x, mean = data.mean, sd = data.sd), add = T)
  }
}
```

Question 3b

```
# again, I'd rather specify the dataset instead of use attach()
set.seed(1)
boot.ci(StudentData$HSClass)
```

2.5% 97.5% ## 214.9068 417.5470

Trimmed Mean Sampling Distribution



Question 3c

```
# again, I'd rather specify the dataset instead of use attach()
set.seed(1)
boot.ci(StudentData$HSClass, trim.percent=.25)
```

2.5% 97.5% ## 152.2092 382.6329

Trimmed Mean Sampling Distribution

