

Fezeka Nzama NZMFEZ001 Project 1

Project Aims & Objectives

In this study we aim explore samples, bootstrapping, hypothesis testing as well as confidence interval construction. In question 1, we conduct an exploratory analysis of the data, perform a 1-sample hypothesis test and construct a confidence interval for both a population mean and a population median. Question 2 focuses on 2-sample hypothesis testing and confidence interval construction, whilst also comparing the results obtained from bootstrapping to those obtained from the use of normal theory. Question 3 explores 4-sample ANOVA testing using bootstrapping.

**Please notes that comprehensive code for all 3 questions as well as the answer for Question 3B are provided in the Appendix.*

Question 1

The following is an analysis of the Income and Expenditure survey conducted by Stats SA, with a focus on the age of the head of the household, and the total monthly income of the household (denoted by the hhinc variable).

Question 1A) The following is an exploratory analysis of the sample age.

#The five number summary of the sample age.

`summary(age)`

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   21.00   33.00   41.00   42.58   52.00   65.00
```

#The variance and standard deviation of the sample age.

`var(age)`

```
## [1] 147.6804
```

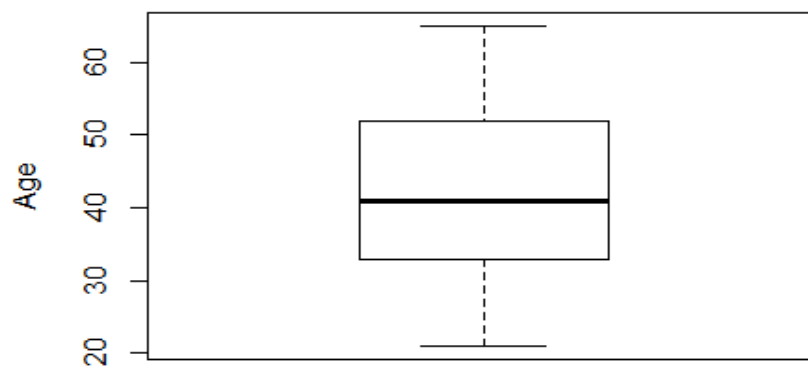
`sd(age)`

```
## [1] 12.15238
```

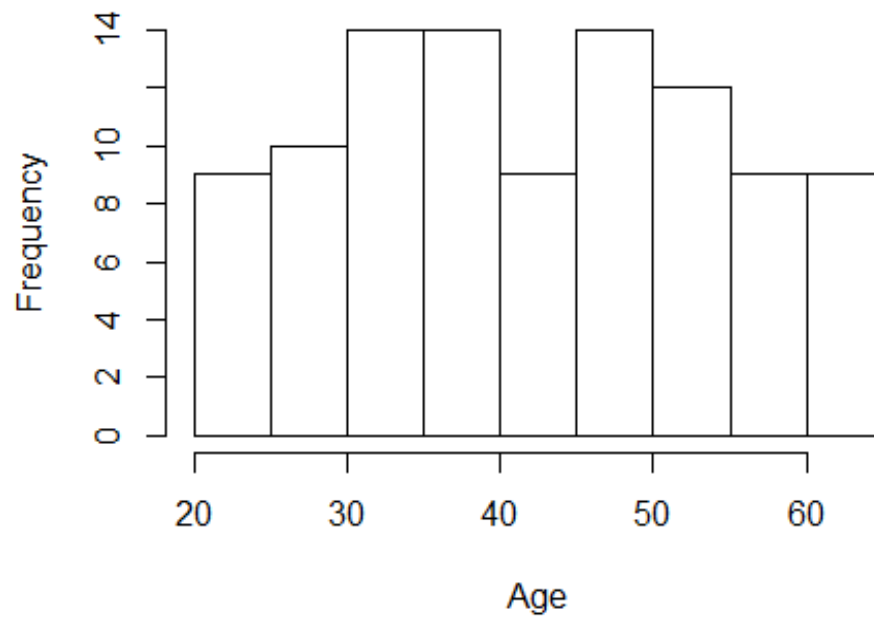
`boxplot(age, ylab = "Age", main="Box-and-Whisker-Plot: Sample age")`

`hist(age, xlab = "Age", main = "Histogram showing the sample age of the head of the household")`

Box-and-Whisker-Plot: Sample age



"Histogram showing the sample age of the head of the household



The following is an exploratory analysis of the sample hhinc.

```
#The five number summary of the sample total monthly income of the household.
summary(hhinc)

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  17372   20752   22912   22649   24182   27362

#The variance and standard deviation of the sample total monthly income of the household.
var(hhinc)

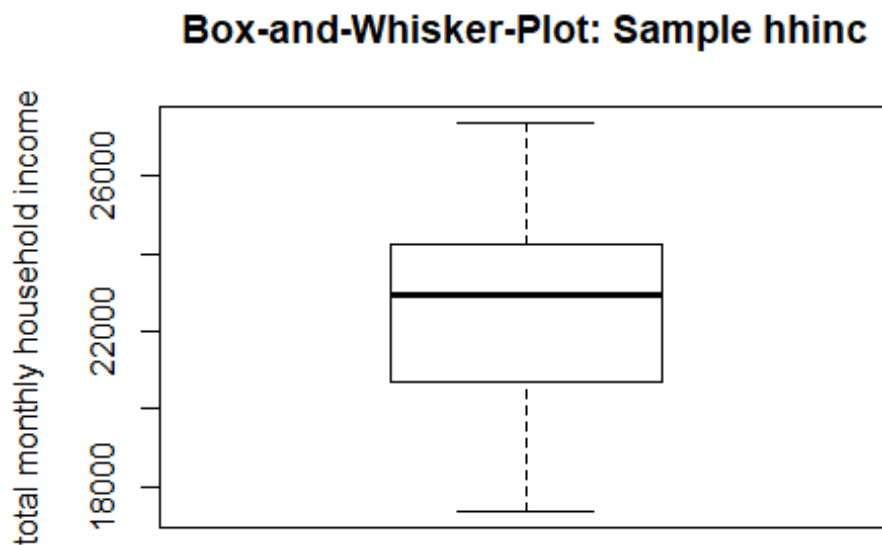
## [1] 6200217

sd(hhinc)

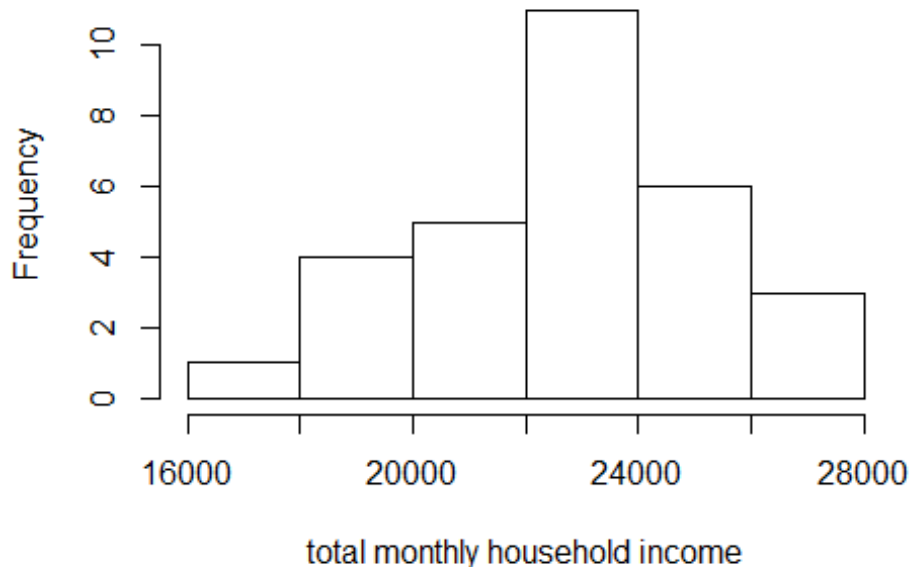
## [1] 2490.024

#The boxplot for the sample total monthly income of the household.
boxplot(hhinc, ylab="total monthly household income", main="Box-and-Whisker-Plot: Sample hhinc")

hist(hhinc, xlab = "Household monthly income", main = "Histogram showing the sample total monthly income in a household")
```



Histogram showing the sample total monthly income of the household



Question 1B)

In the following section we construct a 95% confidence interval for the sample age. To do this we will use bootstrapping to simulate the sampling process. Additionally, we make use of the bootstrap assumption: $\bar{X} - \mu \sim \bar{X}_b - \bar{X}$

#Using the vector of sorted bootstrap means we find the upper and lower bound of the set.

```
lowBound = agesortm[4000*.025]
```

```
uppBound = agesortm[4000*.975]
```

```
truLow=(uppBound*-1)+(2*mean(age))
```

```
truUpp =(-1*lowBound)+(2*mean(age))
```

```
cat("The 95% confidence interval of the mean age is (",truLow,",",truUpp,")")
```

```
## The 95% confidence interval of the mean age is ( 40.14 , 44.98 )
```

The 95% confidence interval can be interpreted as being the probability of stating the correct bounds. Thus, in repeated sampling, bounds between 40.14 and 44.98 were

observed 95% of the time, hence the sample mean age can be said to sit within those bounds in 95% of observations.

Question 1C)

In the following section we use bootstrapping to test the hypothesis that the population mean age for the head of the household is less than or equal to 43. We will make use of the bootstrap assumption: $\bar{X} - \mu \sim \bar{X}_b - \bar{X}$

$H_0: \mu \leq 43$

$H_1: \mu > 43$

```
#calculate the sampling error assuming the null hypothesis is true  
SE = mean(age) - 43
```

Therefore assuming $\mu \leq 43$:

$\bar{X} - \mu \leq -0.42$.

Applying the bootstrap assumption :

$\bar{X}_b - \bar{X} \leq -0.42$

$\bar{X}_b \leq -0.42 + \bar{X}$

```
bMean = SE + mean(age)  
index = match(bMean, agesortm)
```

```
#The pvalue for our hypothesis test is be calculated by summing up the number  
of bootstrap means that are greater than bMean and dividing that by the total  
number of bootstrap means
```

```
pvalue = (4000 - index)/4000  
pvalue
```

```
## [1] 0.6375
```

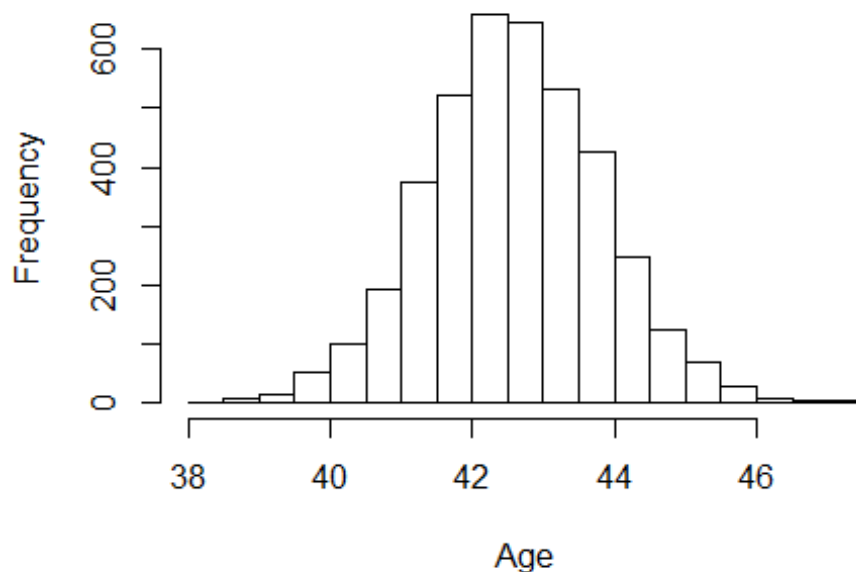
Our observed p-value of 0.6375 does not indicate that there is a significant difference in our assumed value for μ and the true population mean age. Thus, assuming the null hypothesis is true there isn't significant evidence that it is unlikely that the sample we drew would come from a population with that value for μ . As such we cannot reject H_0 .

Question 1D)

In the following section we construct a histogram for the bootstrap means for the age of the head of the household.

```
hist(agesortm, xlab = "Age", main = "Histogram showing the distribution of the  
bootstrap means for the age of the head of the household variable")
```

Histogram showing the distribution of the bootstrap means for the age of the head of the household



The histogram takes on a shape that is very similar to that of the normal distribution. However, it is visibly more peaked, unlike the normal distribution. Thus one can assume that the bootstrap means follow a t-distribution. Additionally, the histogram indicates that the sample mean age for the head of the household lies between 42 and 43 years of age, this is the interval at which the histogram reaches its peak. The histogram also shows that the range of ages for head of the household extends from about 38 to 48, with most of the bootstrap means concentrated around the range from 41 to 44.

Question 1E)

In the following section we construct a 90% confidence interval for the median household income ($\varepsilon_{0.5}$). We make use of bootstrapping to do this, as well as the bootstrap assumption applied to medians rather than means in this case.

```
#Lower and upper median bound
lowMed = hhincM[4000*.05]
uppMed = hhincM[4000*.95]
popLow = -(uppMed - median(hhinc))+median(hhinc)
popUpp = - (lowMed - median(hhinc))+median(hhinc)
cat("(",popLow,",",popUpp,")")

## ( 22171.43 , 23771.28 )
```

The median is often used instead of the mean when estimating the probable value of a random variable when the data tends to exhibit outliers which may skew the mean as the median is a more robust measure. The 90% confidence interval for the population household income median is the frequency in repeated sampling of stating correct bounds

for the population household income median. In this case the confidence interval is (22171.43, 23771.28). Thus in repeated samples, the sample median lies within that range 90% of the time.

Question 2

In this question we compare the marketing strategies to sell a new Apple juice concentrate employed in two cities with the aim of identifying which (if any) is more effective. In the first city the emphasis of the marketing communication is on convenience, whilst in the second city the emphasis is on price. The number of packages sold per week has been recorded for each city, and these numbers are used as an indicator of the success of each marketing strategy.

Question 2A)

In this section we test whether there is a significant difference between the means in the city where convenience is emphasised versus the city where the price is the focal point of the marketing material. To do this we will use bootstrapping and the bootstrap assumption: $\bar{X} - \mu \sim \bar{X}_b - \bar{X}$

(Variances are assumed to be equal)

H₀: $\mu_1 = \mu_2$

H₁: $\mu_1 \neq \mu_2$

```
samplingErr = mean(city1)-mean(city2)-0
samplingErr

## -20.4

uppTail = -samplingErr
```

Therefore assuming **H₀** is true:

$$-20.4 < \bar{X}_1 - \bar{X}_2 - (\mu_1 - \mu_2) < 20.4$$

$$-20.4 < \bar{X}_1 - \bar{X}_2 - 0 < 20.4$$

Applying the bootstrap assumption :

$$-20.4 < \bar{X}_{b1} - \bar{X}_{b2} - 0 < 20.4$$

$$P\text{-val} = \Pr[(\bar{X}_{b1} - \bar{X}_{b2}) < -20.4] + \Pr[(\bar{X}_{b1} - \bar{X}_{b2}) > 20.4]$$

```
greaterThan = match(uppTail, sortCityDiff)
lessThan = match(samplingErr, sortCityDiff)
pval = ((5000-greaterThan)+(lessThan))/5000
pval

## 0.6014
```

Our observed p-value of 0.6014 does not indicate that there is a significant difference in the mean number of packages sold in City 1 (where convenience was emphasised) as opposed to those sold in City 2 (where the emphasis was on the price). Thus we cannot reject the

null hypothesis, meaning that from this result we cannot assume that the marketing employed in each city made a significant difference to the number of units of Apple juice sold.

Question 2B)

In the following section we construct a confidence interval for the difference in population means of packages of Apple Juice sold in City 1 as opposed to City 2.

```
lowBound = sortCityDiff[5000*.025]
uppBound = sortCityDiff[5000*.975]

#calculation of the true population mean difference bounds
PlowBound = (uppBound*-1)+(mean(city1)-mean(city2))
PuppBound = (lowBound*-1)+(mean(city1)-mean(city2))
confi = cat("(",PlowBound,",",PuppBound,")")

## ( -98.1 , 59.6 )
```

The 95% confidence interval can be interpreted as being the probability of stating the correct bounds of the difference in means between city1 and city2. Thus, in repeated sampling, bounds between -98.1 and 59.6 were observed 95% of the time, hence the sample mean difference in packages sold of Apple Juice can be said to sit within those bounds in 95% of observations.

Question 2C) In this section we will be testing the samples from our two cities for equality of variance using bootstrapping.

H₀: $\sigma_1 = \sigma_2$

H₁: $\sigma_1 \neq \sigma_2$

```
sampleFstat = (var(city1))/(var(city2))

#cityFs is the bootstrapped Fstats calculated from the data - see appendix for detail
pvalV= length(cityFs[cityFs>=sampleFstat])/5000
pvalV

## [1] 0.4974
```

We found an observed p-value of 0.4974 which does not indicate a significant difference between the variance in packages of Apple juice sold in City 1 as opposed to City 2. Thus we cannot reject the null hypothesis, as there isn't significant evidence to suggest that the two variances are different.

Question 2D)

In this section we calculate the results obtained from normal theory and bootstrapping.

```
#the p-value calculated using normal theory for the equality of means test
Tp_value = t.test(city1,city2)
Tp_value$p.value
```



```
## [1] 0.6286176
```

```
#the p-value calculated using normal theory for the equality of variance test  
pf(sampleFstat, 9,9)
```

```
## [1] 0.5064688
```

Using normal theory we found that for the equality of means test p-value = 0.6286176 as opposed to the result obtained from bootstrapping of p-value = 0.6014. This shows the p-value obtained via bootstrapping to be very close to that obtained using normal theory. Similarly, when testing for equality of variance the bootstrapped p-value = 0.4974, whilst normal theory gave a p-value = 0.5064688 again showing similarity in the p-value obtained by bootstrapping to normal theory. Thus both sets of results show that normal theory is consistent to results obtained by repeated sampling and observation and as such validate the use of normal theory in hypothesis testing as a means to estimate the p-value when repeated sampling is not possible.

Question 3

In this section we will review data provided by the Internal Revenue Service. The IRS wishes to improve the wording and format of it's tax return form so as to make them easier to fill out. As such the IRS conducted an experiment where 120 individuals were grouped into 4 groups of 30 and asked to fill out tax return forms whilst being timed. Each group received a different form (3 groups getting new forms, whilst one used the old form).

We wish to discern whether there is a significant difference in the mean times of each group and thus determine if there is significant evidence to motivate a change in the wording and format of the forms. We will do this using the bootstrapped ANOVA method.

H₀: $\mu_i = \mu$

H₁: $\mu_i \neq \mu$

The following statistics were obtained from the provided sample data:

SSE = 1.186726710^{5}

SST = 1.244403310^{4}

F-stat = 4.1594525

```
p_value=length(Ratios[Ratios>Forig])/B  
p_value
```

```
## [1] 0.0075
```

Our Bootstrapping process resulted in a p-value of 0.0075. This p-value is significantly small and thus suggests that there is significant difference in mean times for filling in the form. As such we can reject **H₀** at a significance level of 1%.

From this test we can see that there may be value in altering the wording and formatting of the tax return forms, as at least one of the forms had a mean fill out time which was different to the others. This thus provides motivation for further study to be done around this data to determine where the differences actually lie, ie. which form or forms had a different mean time. From this further investigation we would then be able to determine

which form has the best wording and formatting and advise the IRS to adopt that form for use.

Plagiarism Statement: The work presented is all my own.

Appendix

#Question 1 code & Logic

#reading in data

```
mydata = read.table("F:\\sta3030 - Project 1\\incexp.txt", header = TRUE)
age = mydata$AGE[118:(118+99)]
hhinc = mydata$HHINCOME[118:(118+29)]
```

#Question 1B - Bootstrapping

#The 'agebstr' variable is the matrix in which we will store bootstrap sample age.

```
agebstr<-matrix(0,nrow=4000, ncol=100)
for(i in 1:4000){
  samp = sample(age, size=100, replace = TRUE)
  agebstr[i,] = samp
}
```

#The 'agebstrm' is a vector containing the mean of each bootstrap sample, whilst 'agesortm' contains the means sorted in ascending order.

```
agebstrm = apply(agebstr, 1, mean)
agesortm = sort(agebstrm)
```

#the following is the logic used in calculating the population confidence interval

```
#xb = bootstrap mean, ageMean = sample mean, popMean = population meanS
#Pr[LowBound<Xb<uppBound]
#Pr[LowBound-ageMean<Xb-ageMean<uppBound-ageMean]
#Pr[LowBound-ageMean<agemean-popMean<uppBound-ageBound]
#Pr[-(uppBound-ageMean)<popMean-ageMean<-(LowBound-ageMean)]
#Pr[-(uppBound-ageMean)+ageMean<popMean<-(LowBound-ageMean)+ageMean]
```

#Question 1E - Bootstrapping

#hhincM is a vector in which the household income bootstrapped medians are stored

```
hhincM<-vector("numeric", length = 4000)
```

#bootstrapping

```
for(i in 1:4000){
  samp =sort(sample(hhinc, size=30, replace = TRUE))
  #find the median value in the bootstrap
  hhincM[i]=median(samp)
}
```

#sorting the vector

```
hhincM = sort(hhincM)
```

#Question 2 code & Logic

#reading in required data

```
quest2 = read.table("F:\\sta3030 - Project 1\\A1.txt",header= TRUE)
```

```
city1=quest2$City1
```

```
summary(city1)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  444.0   482.0   552.0   555.5   613.5   712.0
```

```
city2 =quest2$City2
```

```
summary(city2)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  464.0   529.5   557.0   575.9   632.5   759.0
```

#Question 2A)

*#A combined data set is created by combining the data from the two cities
#Bootstraps of the combined are created and the means of the bootstraps are stored in the bootstrap mean vectors(city1B and city2B). The mean differences are also then computed and stored in vector cityDiff*

```
combined = c(city1,city2)
```

```
city1B <-vector(mode="numeric", length = 5000)
```

```
city2B <-vector(mode="numeric", length = 5000)
```

```
cityDiff <-vector(mode="numeric", length = 5000)
```

```
for(i in 1:5000){
  samp = sample(combined, size = 20, replace = TRUE)
  city1B[i] = mean(samp[1:10])
  city2B[i] = mean(samp[11:20])
  cityDiff[i] = city1B[i]-city2B[i]
}
```

```
sortCityDiff =sort(cityDiff)
```

#Question 2B)

#95% confint - find index of (5000.025) & (5000*.975)*

*#LowBound<bmean(1-2) - smean(1-2)<uppBound **note that is smean(1-2)=0*

#LowBound<bmean(1-2) <uppBound

#LowBound<smean(1-2)- u(1-2)<uppBound

#-uppBound<u(1-2)-smean(1-2)<-LowBound

#therefore: -uppBound+smean(1-2)<u(1-2)<-LowBound+smean(1-2)

#Question 2C)

#the following code block was used to create a vector (cityFs) containing the bootstrap f-stats of city one and two

```
cityFs <-vector(mode="numeric", length = 5000)
```

```
for(i in 1:5000){
```

```

samp = sample(combined, size = 20, replace = TRUE)
city1v= c(samp[1:10])
varCity1v = var(city1v) #calculate variance of bootstrap of city1
city2v= c(samp[11:20])
varCity2v=var(city2v) #calculate variance of bootstrap of city2
cityFs[i] = varCity1v/varCity2v #calculate Fstat and retain that
}

```

#Question 3

#reading the data

```

quest3 = read.table("F:\\sta3030 - Project 1\\aov5.txt",header= TRUE)
quest3

```

##	Form_1	Form_2	Form_3	Form_4
## 1	23	88	116	103
## 2	59	114	123	122
## 3	68	81	64	105
## 4	122	41	136	73
## 5	74	108	99	87
## 6	90	92	156	81
## 7	70	52	175	120
## 8	87	54	93	169
## 9	155	103	77	130
## 10	120	50	88	56
## 11	124	135	91	101
## 12	103	76	118	143
## 13	54	143	86	106
## 14	90	124	164	129
## 15	124	151	101	104
## 16	80	96	74	169
## 17	69	76	124	69
## 18	123	128	137	76
## 19	76	60	69	55
## 20	71	127	136	138
## 21	94	109	127	122
## 22	167	122	135	139
## 23	69	88	97	138
## 24	105	109	103	132
## 25	98	90	86	99
## 26	73	56	121	64
## 27	79	105	98	89
## 28	61	64	59	128
## 29	121	127	91	127
## 30	56	104	61	161

#treatment groups

```

form1 = quest3$Form_1
form2 = quest3$Form_2
form3 = quest3$Form_3

```

```

form4 = quest3$Form_4

#number pof treatment groups
k = 4
#number of participants per group
ni = 30
#total no. participants
N = 30*4
#overall mean
Y.. = (sum(form4,form3, form2, form1))/N
#Y..

B = 4000

#calculate values for the provided sample
sSSE=sSST=0
#treatment groups placed in list
forms = list(form1, form2, form3, form4)
#mean of each treatment group
Yi. = vector( mode = "numeric", length = 4)
for(i in 1:k){
  Yi.[k]=mean(forms[[k]])
  #sum((forms[[k]]- Yi.[k])^2) does the (Yij-Yi)^2 for all values in that loc
  ked list
  #vector minus a vector basically
  sSSE=sSSE+sum((forms[[k]]- Yi.[k])^2)
  sSST =sSST+ni*((Yi.[k]-Y..)^2)
}
Forig = (sSST/(k-1))/(sSSE/(N-1))

#bootstraps
combined = unlist(forms) #combined list of all our sample data
Ratios = vector("numeric") #empty vector to store bootstrap F-stats
sampleToPrint = list("numeric", length = 3)
for (i in 1:B){
  bSSE=bSST=0
  Yi.=vector("numeric")
  samp = sample(combined, size = 120, replace = TRUE)
  bstrp = list("numeric", length = k)
  bstrp[[1]]= samp[1:30]
  bstrp[[2]]= samp[31:60]
  bstrp[[3]]= samp[61:90]
  bstrp[[4]]= samp[91:120]
  if(i<4){
    sampleToPrint[[i]]=samp
  }
  Y.. = mean(unlist(bstrp))
  for(j in 1:k)
  {Yi.[j]=mean(bstrp[[j]])}
}

```

```

bSSE=bSSE+sum((bstrp[[j]]-Yi.[j])^2)
bSST=bSST+ni*(Yi.[j]-Y..)^2
}
Ratios[i]=(bSST/(k-1))/(bSSE/(N-k))
}

```

Question 3B)

#3 bootstrap samples generated -
print(sampleToPrint)

```

## [[1]]
## [1] 175 96 70 167 122 76 109 124 59 61 120 90 23 54 99 76 90
101
## [19] 169 64 104 127 104 64 139 90 80 59 59 56 118 169 156 77 76
132
## [37] 124 103 103 127 137 122 69 60 97 96 99 87 122 97 118 64 122
52
## [55] 54 105 124 122 81 77 56 81 87 105 56 175 90 88 23 138 128
155
## [73] 87 156 139 96 90 59 128 55 64 50 169 103 68 64 103 128 81
50
## [91] 91 81 121 80 121 69 76 64 137 161 98 137 122 122 76 120 137
99
## [109] 54 101 59 56 104 104 56 128 161 121 104 90
##
## $length
## [1] 120 69 109 161 123 127 70 88 69 135 97 135 124 135 161 90 91
169
## [19] 127 41 123 87 87 129 76 69 79 124 64 56 120 155 130 120 76
88
## [37] 108 105 155 109 88 87 87 73 136 127 103 127 169 76 99 56 124
103
## [55] 169 127 167 169 143 137 120 89 76 169 23 114 106 127 101 94 86
101
## [73] 71 76 123 151 59 60 129 64 77 122 136 103 128 64 64 143 64
130
## [91] 106 81 79 156 86 88 88 73 137 41 127 90 70 41 73 143 118
135
## [109] 143 88 137 76 64 92 138 132 52 86 130 69
##
## [[3]]
## [1] 98 76 69 127 135 68 108 61 92 88 91 143 81 55 81 135 90
90
## [19] 156 114 161 64 124 127 97 114 127 136 55 124 52 60 74 76 64
69
## [37] 77 61 88 105 122 76 151 155 122 99 61 73 138 81 103 109 103
89
## [55] 108 90 76 105 155 122 87 123 169 90 105 123 76 104 127 103 124
74

```

```
## [73] 56 108 90 80 60 54 167 76 143 109 80 114 99 54 104 56 105
81
## [91] 87 175 56 132 93 143 127 143 104 161 99 90 128 155 64 103 52
74
## [109] 41 121 86 73 122 50 129 71 122 128 69 124
```