# MINI PROJECT 3 GROUP NO. 28

## Members JEEVAN DSOUZA SAMARTH SAIRAM

### Contributions Rendered

Both the team members rendered contributions equally with respect to analysis and design of the solution for the problem statement and programming for the same.

```
1. a) #using Monte Carlo method
```

# we first have to set the population parameter that is theta

# Next we need to generate the samples from the population and need to calculate cap theta

# Then we can get the MSE which is the squared difference of the population parameter theta and calculated cap theta.

#### 1. b)

#Trying to calculate theta1 which is from MLE and Theta 2 that is method of moments estimator

#First we need to calculate both mle and moments from the same sample and then we need calculate MSE for both of them and we see in

```
MSE_moments_Esimate = function(sample_size,theta_value) {

#we use Monte carlo simulation thus we use runif function

monte_carlo_value = runif(sample_size,min=0,max=theta_value)

#MLE calucation is obtained by taking the derivative of the log function and equating it to zero, but in here it is maximum of sample

Estimated_MLE = max(monte_carlo_value)

#Method of moments is 2 times the mean of generated value
```

#Next we return both the generated values
return (c(Estimated\_MLE,Estimated\_Moments))
}

Estimated\_Moments = 2 \* mean(monte carlo value)

```
#we need to calculate if for 1000 replications
MSE moments Calculte = function(sample size,theta value){
#We need to calculate MLE for both the estimators, so we call the
function to estimate cap theta and and apply the formula to calculate
both MLE and moments
Cap Theta =
replicate(1000,MSE moments Esimate(sample size,theta value))
#we use the definition of MSE to calculate the mean square difference
Mean Squared Error = (theta value - Cap Theta) ^ 2
#To get different values for both of the estimators
Moments Calculated = Mean Squared Error[c(TRUE,FALSE)]
MLE Calculated = Mean Squared Error[c(FALSE,TRUE)]
Moments Calculated = mean(Moments Calculated)
MLE Calculated = mean(MLE Calculated)
return(c(MLE Calculated, Moments Calculated))
#For sample n=1 and theta =1
MSE Calulate 1 1 = MSE moments Calculte(1,1)
MSE Calulate 1 1
1.c)
#To calculate for all other combinations
```

#we will have 2 values one for MLE and the other for moments we need to plot them we will have 2 variations fixed value theta and

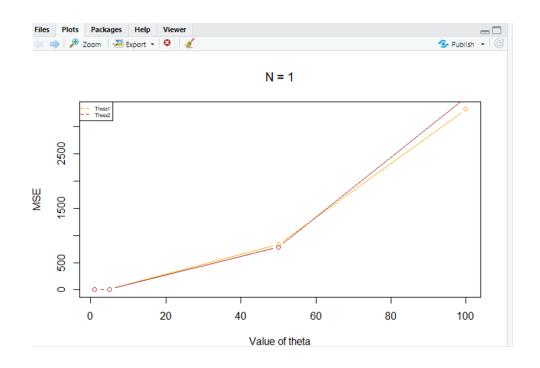
#constant n and vice versa

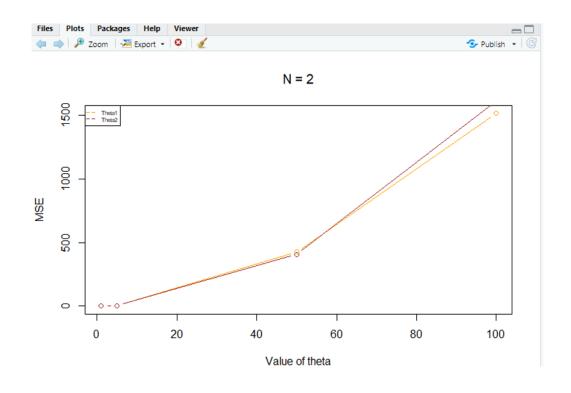
#First we add values of theta and n to a column then calculate the values by looping through 2 times and estimating those values

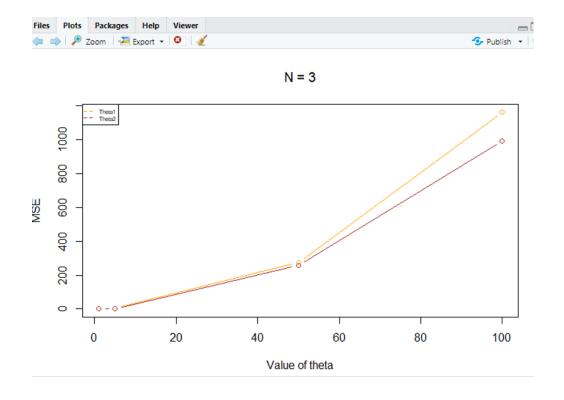
#This is for n

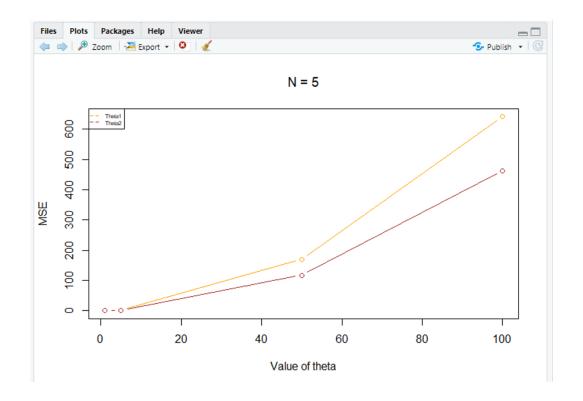
```
vals of n = c(1,2,3,5,10,30)
vals of theta = c(1,5,50,100)
counter = 0
#initializing both theta and
MSE cal theta1 = c(0,0,0,0)
MSE Calc theta2 = c(0,0,0,0)
for(itr1 in vals of n)
 counter=1
 for(itr2 in vals of theta)
 {
  calc = MSE moments Calculte(itr1,itr2)
  MSE cal theta1[counter] = calc[1]
  MSE Calc theta2[counter] = calc[2]
  counter = counter + 1
```

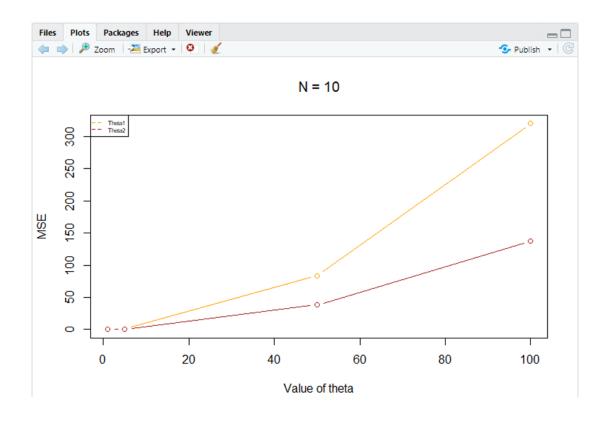
```
plot(vals of theta, MSE cal theta1, ylab = 'MSE', main=bquote(paste("N = ", .
(itr1)), xlab = 'Value of theta',
    type = 'b',col='orange')
 lines(vals of theta,MSE Calc theta2,col='brown',type = 'b')
 legend("topleft",legend=c("Theta1","Theta2"),col=c('orange','brown'),cex =
0.5, lty=c(2,2), merge = TRUE)
#This is for theta values
vals of n = c(1,2,3,5,10,30)
vals of theta = c(1,5,50,100)
counter = 0
#initializing both theta and
MSE cal theta1 = c(0,0,0,0,0,0)
MSE Calc theta2 = c(0,0,0,0,0,0)
for(itr1 in vals of theta)
 counter=1
 for(itr2 in vals of n)
 {
  calc = MSE moments Calculte(itr2,itr1) #Interchanging the values in here
  MSE cal theta1[counter] = calc[1]
```





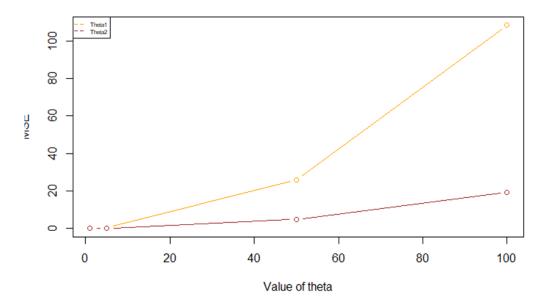


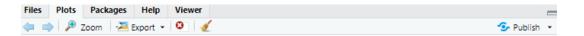




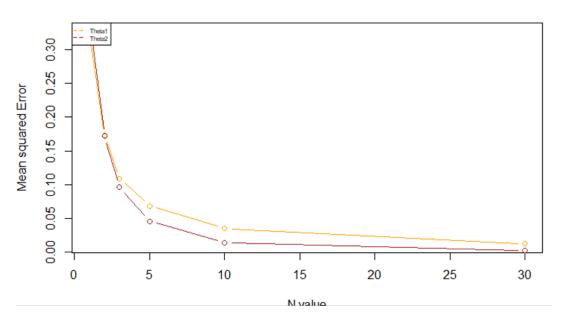


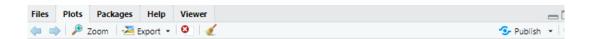
N = 30



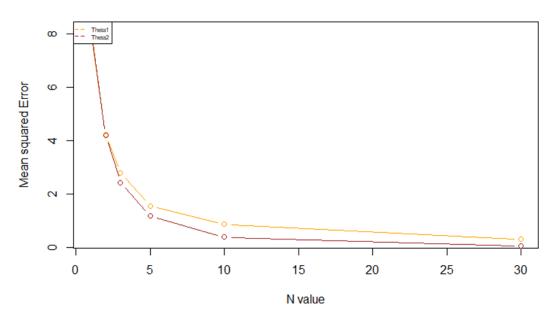


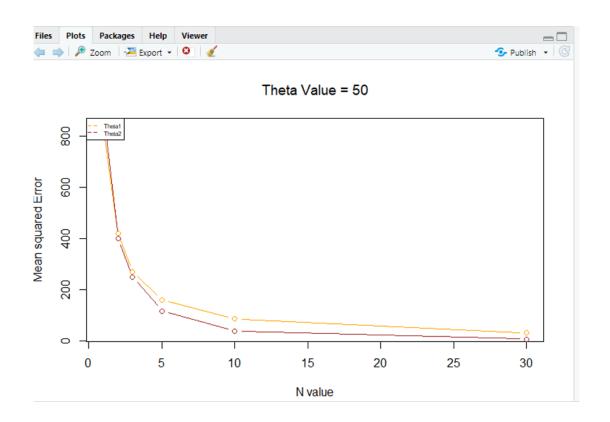
#### Theta Value = 1

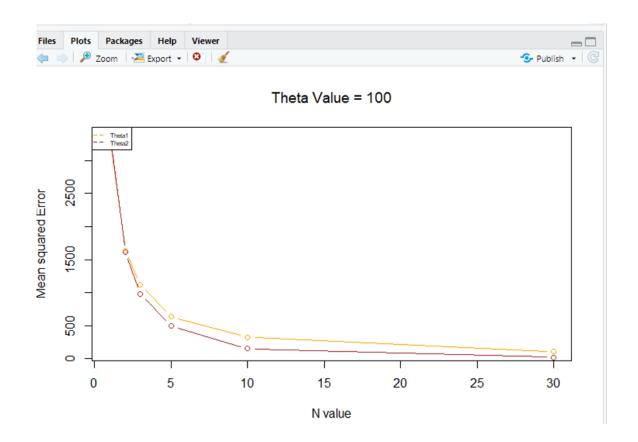




Theta Value = 5







## 1.d)

It can be inferred that as the sample value increases mean squared error goes on decreasing also MLE is better than Moments estimatoras it shows lower MSE, thus we infer MLE is better for bigger values n #in the second case Theta values might differ but the resulting graphs are similar, thus inferring that estimator wouldn't depend on theta value #Thus MLE is better than method of moments as sample size increases

Soln: We denote MLE with L(0) and in here, we have  $L(0) = \prod_{i=1}^{n} \left( \frac{0}{x_i^{n+1}} \right)$ 

Since, this is a product, taking logarithm on both sides it becomes a summation & is strictly increasing. In probability theory, we take natural logarithm.

$$log(L(\theta)) = log_{e}\left(\prod_{i=1}^{n}\left(\frac{\theta}{x_{i}^{\theta+i}}\right)\right).$$

On time is  $0^n$  & we can take it out of summation,  $\log(L(0)) = \log(0^n \times \prod_{i=1}^n \left(\frac{1}{x_i^{n+1}}\right))$ 

using (1) we have,

$$log_{\mathcal{L}}(L(0)) = nlog_{\mathcal{O}} + \sum_{i=1}^{n} log_{\mathcal{X}_{i}}^{-(\mathcal{O}+1)}$$

$$log_{e}(L(\theta)) = n log_{\theta} - (\theta+1) \sum_{i=1}^{n} log_{e}x_{i}$$

$$= n log_{\theta} - \theta \sum_{i=1}^{n} log_{e}x_{i} - \sum_{i=1}^{n} log_{e}x_{i}$$

On partially differentiating the above function,

$$\hat{\theta}_{MLE} = \frac{5}{15.461}$$

```
#Add the values to a vector

max_log_func <- c(21.72,14.65,50.42,28.78,11.23)

#writing a function to get the likelihood parameter

log_likelihood <- function(par,data_pt) {

res = length(data_pt)*log(par)-(par+1)*sum(log(data_pt))

return(-res)
}

#using optim function and giving hessian as true

max_likely <-optim(par = 0.5,data_pt=max_log_func,
fn=log_likelihood,method="L-BFGS-B",hessian = TRUE,lower = 0.01)

max_likely
```

```
. #using optim function and giving hessian as true
> max_likely <-optim(par = 0.5,data_pt=max_log_func, fn=log_likelihood,method="L-BFGS-B",hessian = TRUE,lower = 0.01)
> max_likely
    $par
[1] 0.3233885
    $value
    [1] 26.10585
    $counts
    function gradient
    $convergence
    [1] 0
    [1] "CONVERGENCE: REL_REDUCTION_OF_F <= FACTR*EPSMCH"
    $hessian
    [1,] 47.81116
   #result
   #Yes the answers match
   2.d
   #We need to get the hessian matrix to calculate this estimate
   err<- sqrt( diag(solve(max likely$hessian)))
   err
    alp val < -0.05
   # confidence interval is mean +/- z-score multiplied by error, we use
   gnorm function to find it
   Confidence Int = max likelypar + c(-1, 1)* qnorm(1-(alp val/2))* err
   Confidence Int
> #We need to get the hessian matrix to calculate this estimate
> err<- sqrt( diag(solve(max_likely$hessian)))</pre>
[1] 0.1446223
> alp_val <- 0.05
> # confidence interval is mean +/- z-score multiplied by error, we use gnorm function to find it
> Confidence_Int = max_likely$par + c(-1, 1)* qnorm(1-(alp_val/2)) * err
> Confidence_Int
[1] 0.0399339 0.6068430
```

> max\_log\_tunc <- c(21,/2,14.65,50.42,28./8,11.23)

> #writing a function to get the likelihood parameter
> log\_likelihood <- function(par,data\_pt){
+ res = length(data\_pt)\*log(par)-(par+1)\*sum(log(data\_pt))</pre>

#infer-> The true estimate lies within the given confidence interval i,e out of 100 trials 95 percent of the trials will be there

#in the given interval