SS2864B Assignment 5

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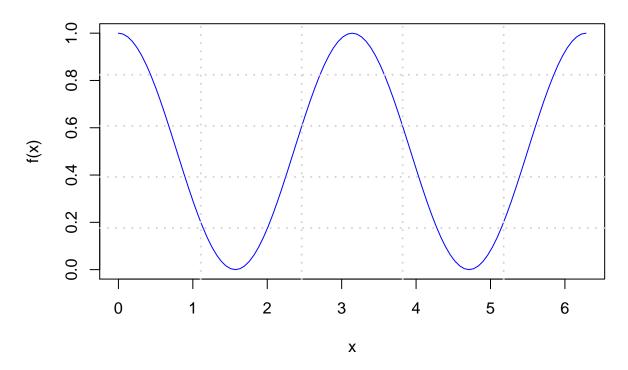
Question 1

```
# define the given function
f1 = function(x) {
    return ((cos(x))^2)
}

# define a CI function
ci.t.95 = function(x) {
    # compute lower and upper bounds
    lower = mean(x) - 1.96*sd(x)/sqrt(length(x))
    upper = mean(x) - 1.96*sd(x)/sqrt(length(x))
# construct a list object
    result = list(Lower.Bound = lower, Mean = mean(x), Upper.Bound = upper)
    return (result)
}
```

(a)

A plot of $[\cos(x)]^2$



(b)

```
# set seed for reproducibility
set.seed(123)
# define a random uniform variable on [0, 2pi]
U = runif(n = 1e6, min = 0, max = 2*pi)
# perform MC integration
I1 = mean(f1(U))*(2*pi)
# compute a 95% ci
ci = ci.t.95(f1(U)*(2*pi))
# display ci
ci
## $Lower.Bound
## [1] 3.13889
##
## $Mean
## [1] 3.143241
## $Upper.Bound
## [1] 3.13889
(c)
```

We use some identities to simplify the square of cosine before integrating it:

$$\cos^2(x) = 1 - \sin^2(x)$$
 (Pythagorean Identity)

$$\sin^2(x) = \cos^2(x) - \cos(2x) \text{ (Cosine Double Angle Identity)}$$

$$\therefore 2\cos^2(x) = 1 - \cos(2x) \text{ (Combine Two Identities)}$$

$$I = \int_0^{2\pi} f(x)dx = \int_0^{2\pi} \cos^2(x)dx = \frac{1}{2} \int_0^{2\pi} 1 - \cos(2x)dx$$

$$I = \frac{1}{2} \left(x - \frac{1}{2}\sin(2x) \right) \Big|_0^{2\pi}$$

$$\therefore I = \frac{1}{2} (2\pi) = \pi$$

Based on the output in part b, indeed, the CI covers the true value if we assume the true value = 3.14.

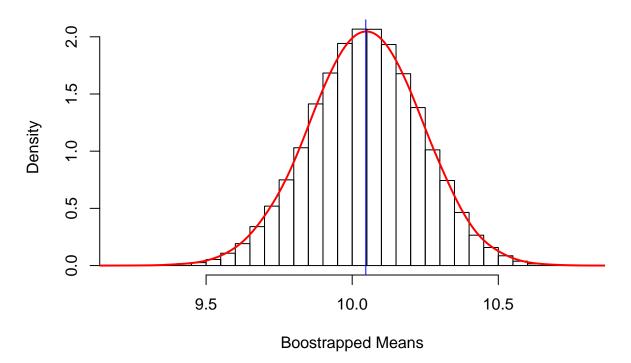
Question 2

(a)

(c)

```
set.seed(222)
# Generate 100 x_i such that X \sim N(10, 4)
x = rnorm(n = 100, mean = 10, sd = 2)
# Generate summary statistics of the x_i's
min(x)
## [1] 4.451066
max(x)
## [1] 14.41816
mean(x)
## [1] 10.04396
sd(x)
## [1] 1.916051
(b)
# Generate 50000 different samples and calculate their means
out = replicate(50000, mean(sample(x, replace=TRUE)))
# Generate summary statistics of the means
min(out)
## [1] 9.243029
max(out)
## [1] 10.78305
mean(out)
## [1] 10.04469
sd(out)
## [1] 0.1905054
```

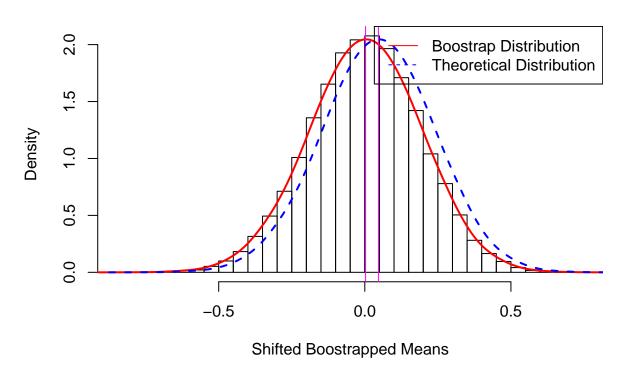
Sampling Distribution of Sample Mean



This output looks like a translated normal distribution centered at the true sample mean.

(d)

Sampling Distribution of Sample Mean (Shifted)



It looks like the boostrap distribution is identical to theoretical distribution except that their centres are shifted (translated). This shift may be called an 'error'. Note that the distributions are equivariant.

```
We will omit the code from the output since it is identical to the last parts. ** Round 2 **
```

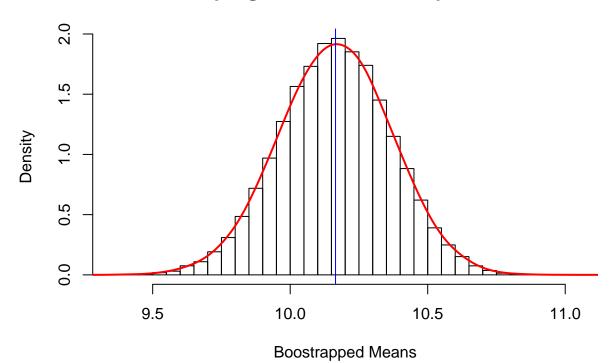
- ## [1] "summary statistics of the sample: "
- ## [1] "min:"

(e)

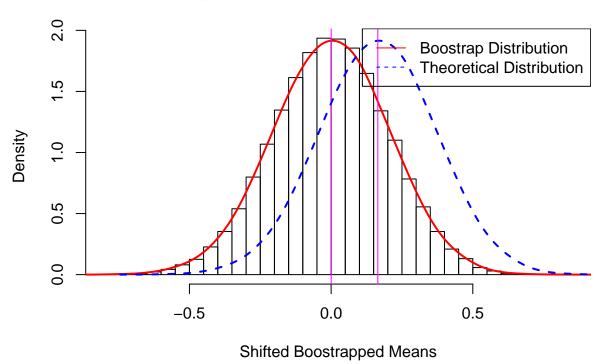
- ## [1] 4.162518
- ## [1] "max:"
- ## [1] 14.9448
- ## [1] "mean:"
- ## [1] 10.16458
- ## [1] "sd:"
- ## [1] 2.039159
- ## [1] "summary statistics of the means (boostrapped data): "
- ## [1] "min:"
- ## [1] 9.381324

- ## [1] "max:"
 ## [1] 11.00874
 ## [1] "mean:"
 ## [1] 10.16348
- ## [1] "sd:"
- ## [1] 0.2026687

Sampling Distribution of Sample Mean



Sampling Distribution of Sample Mean (Shifted)



Note in general the results are similar, except that this time, the 'error' is larger.

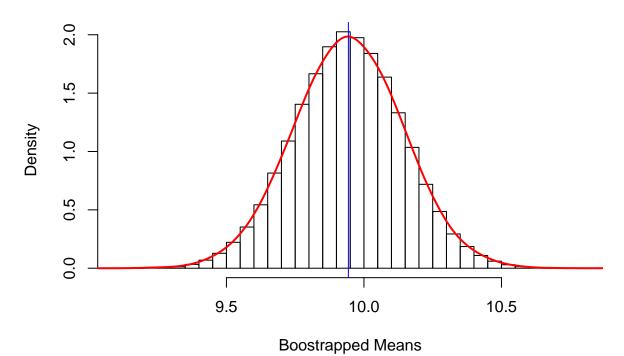
```
** Round 3 **
```

- ## [1] "summary statistics of the sample: "
- ## [1] "min:"
- ## [1] 5.251747
- ## [1] "max:"
- ## [1] 14.22127
- ## [1] "mean:"
- ## [1] 9.94272
- ## [1] "sd:"
- ## [1] 1.991693
- ## [1] "summary statistics of the means (boostrapped data): "
- ## [1] "min:"
- ## [1] 9.14236
- ## [1] "max:"
- ## [1] 10.75656
- ## [1] "mean:"
- ## [1] 9.943207

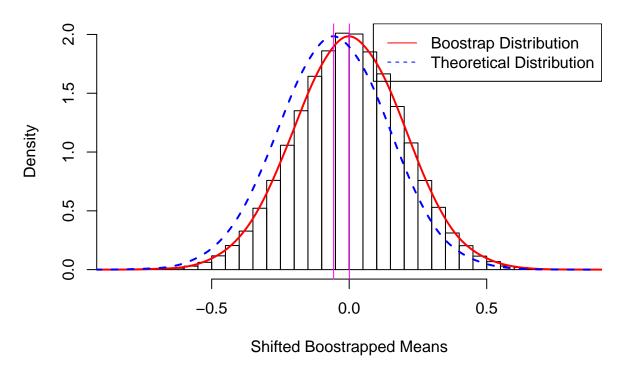
[1] "sd:"

[1] 0.1979312

Sampling Distribution of Sample Mean



Sampling Distribution of Sample Mean (Shifted)



Again, we get similar results, and this time the 'error' has shrunk.

Question 3

(a)

```
my.obj = function(theta, x) {
    # We allow theta to be a vector.
    # If dim(theta) > 1, we return a vector of dim(theta)
    # and we re-use the data in x for all theta given
    # Else, we return a scalar
    if (length(theta) > 1) {
        x.mat = matrix(rep(x, times = length(theta)), nrow = length(x))
        theta.mat = matrix(rep(theta, each = length(x)), nrow = length(x))
        result = colSums(abs(x.mat - theta.mat))
    }
    else {
        result = sum(abs(x - theta))
    }
    return(result)
}
```

(b)

```
# define a function that minimizes my.obj with respect to theta
my.minimizer1 = function(x, interval = c(min(x), max(x))) {
```

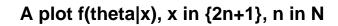
```
o = optimize(my.obj, interval = c(min(x), max(x)), x)
  return(o$minimum)
}
# test the function on the given data
data = c(3, 7, 9, 12, 15, 18, 21)
my.minimizer1(data)
## [1] 12
median(data)
## [1] 12
Our function works perfectly!
(c)
# define a function that minimizes my.obj with respect to theta
my.minimizer2 = function(x, start = mean(x)) {
 o = nlminb(start = start, objective = my.obj, x = x)
  return(o$par)
}
\# test the function on the given data
my.minimizer2(data)
## [1] 12
median(data)
## [1] 12
This function works just as good too.
(d)
# create test data
data = c(1, 3, 7, 9, 12, 15, 18, 21)
# test the functions from b and c
tb1 = my.minimizer1(data, interval = c(min(data) - 2, max(data) + 2))
tb2 = my.minimizer1(data, interval = c(min(data) - 4, max(data) + 4))
tb3 = my.minimizer1(data, interval = c(min(data) - 6, max(data) + 6))
tc1 = my.minimizer2(data, start = min(data))
tc2 = my.minimizer2(data, start = max(data))
tc3 = my.minimizer2(data, start = mean(data)/2)
tb1
## [1] 10.2539
tb2
## [1] 10.2539
tb3
## [1] 10.2539
```

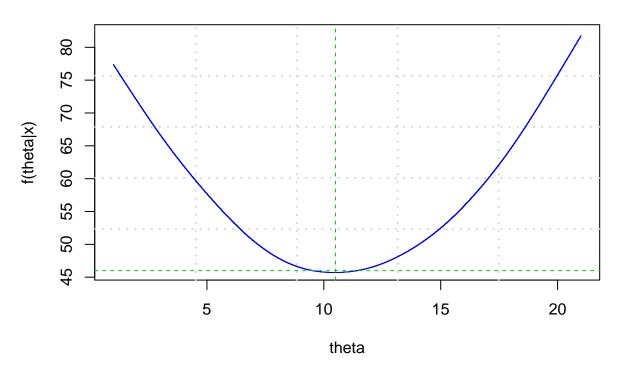
```
## [1] 10.33333
tc2
## [1] 9
tc3
## [1] 9.375
median(data)
```

[1] 10.5

It seems that both functions fail to converge to the minimizer.

```
library('MASS')
# evaluate theta on interval [1, 21]
theta = seq(min(data), max(data), length.out = 1e5)
# evaluate f on theta and the given data
f = my.obj(theta, data)
# set up a grid and plotting area
plot(x = theta, y = f, type = 'n', main = 'A plot f(theta|x), x in {2n+1}, n in N',
    xlab = 'theta', ylab = 'f(theta|x)', col = 4)
grid(5, lwd = 2)
# create a smooth plot of the data
sm = smooth.spline(theta, f, spar=1)
# add the smooth plot
lines(sm, col = 4)
abline(v = median(data), col = 3, lty = 2)
abline(h = my.obj(median(data), data), col = 3, lty = 2)
```





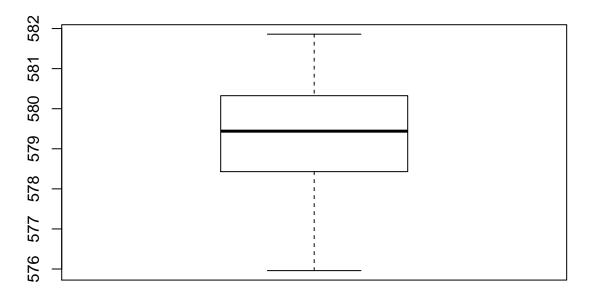
We find that indeed, the value of θ that minimizes $f(\theta|x)$ is the median of x. In this case, it is 10.5.

Question 4

We will omit the code from showing on the output due to its length.

```
mean(huron)
## [1] 579.3091
sd(huron)
## [1] 1.335657
min(huron)
## [1] 575.96
max(huron)
## [1] 581.86
boxplot(huron, main = "Huron Lake Mean Water Level Over 1860-1986")
```

Huron Lake Mean Water Level Over 1860–1986



ts.plot(huron, main = "Huron Lake Mean Water Level Over 1860-1986", ylab = "Mean Water Level")

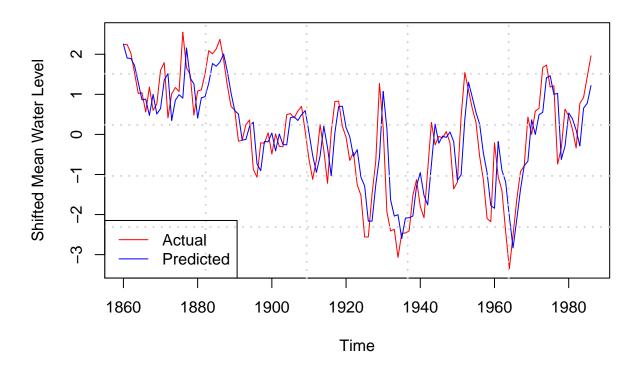
Huron Lake Mean Water Level Over 1860–1986

```
582
       581
Mean Water Level
       580
       579
       578
       577
       9/9
                             1880
               1860
                                            1900
                                                           1920
                                                                         1940
                                                                                        1960
                                                                                                      1980
                                                            Time
```

```
\#\#\# (b)
# define the given likelihood function
log.likelihood=function(par, x){
  n=length(x)
  v=x[1]^2
  for (i in 2:n)
    v=v+(x[i]-par[1]*x[i-1])^2
  return(v/par[2]+n*log(par[2]))
}
# define huron2
huron2 = huron - mean(huron)
# do optimization
my.par = nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2)
## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
## NA/NaN function evaluation
## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
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## NA/NaN function evaluation
```

```
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## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
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## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
## NA/NaN function evaluation
## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
## NA/NaN function evaluation
## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
## NA/NaN function evaluation
## Warning in log(par[2]): NaNs produced
## Warning in nlminb(start = c(-0.1, 540), objective = log.likelihood, x = huron2):
## NA/NaN function evaluation
# extract the parameters
par = my.par$par
# display the parameters
## [1] 0.8459369 0.5250192
(c)
# predict the time series data using the first entry in huron2
pred.huron2 = huron2
for (i in 2:length(huron2)) {
 pred.huron2[i] = par[1]* huron2[i-1]
# plot the predicted and actual time series
ts.plot(huron2, pred.huron2, col = c('red', 'blue'),
        main = "Huron Lake Shifted Mean Water Level Over 1860-1986",
        ylab = "Shifted Mean Water Level")
# add a legend and a grid
grid(5, lwd = 2)
legend("bottomleft", col = c('red', 'blue'), lty = 1, legend = c("Actual", "Predicted"))
```

Huron Lake Shifted Mean Water Level Over 1860–1986



Our model captures the fluctuations of the time series data; however, it does not match the original data and we can see it is off by a constant.