MA 354: Data Analysis – Fall 2021 – Due 10/8 at 5p Homework 2:

Complete the following opportunities to use what we've talked about in class. These questions will be graded for correctness, communication and succinctness. Ensure you show your work and explain your logic in a legible and refined submission.

The starting jobs will be applied in alphabetical order (last name) for question two.

- 1. Solver: provide a solution, if possible, and reasoning for the solution. Due to group 10/5 or earlier.
- 2. Code Checker: provides a first check of the solver's worked solutions and ensures they are correct with a solid interpretation.
- 3. **Checker** checks the solution for completeness, proposes and implements changes if agreed upon by the group. Provides a short paragraph summarizing the discussion of proposals and their reason for acceptance or non-acceptance.
- 4. **Double Checker** checks the solution for completeness, communication and polish. The Double Checker ensures that the solution is correct and highly polished for submission.

For subsequent questions student roles will move down one position. The rolls change as follows.

- 1. Solver \Longrightarrow Code Checker
- 2. Code Checker \Longrightarrow Checker
- 3. Checker \Longrightarrow Double Checker
- 4. Double Checker \Longrightarrow Solver

While students have assigned jobs for each question I encourage students to help each other complete the homework in collaboration.

- 1. Continue with the discrete distribution you selected for Question
 - (a) Provide the mean, standard deviation, skewness, and kurtosis of the PMF. Ensure to interpret each.

Bernoulli Distribution:

PMF:

$$f(k;p) = p^{k} (1-p)^{1-k}$$
 for $k \in \{0,1\}$

Mean:

$$E(X) = p$$

Standard Deviation:

$$\sigma = \sqrt{p\left(1 - p\right)}$$

Skewness:

$$\xi_X = \frac{(1-p)-p}{\sqrt{p(1-p)}}$$

Kurtosis:

$$\kappa_Y = 3 + \frac{1 - 6p(1 - p)}{p(1 - p)}$$

(b) Generate a random sample of size n = 10, 25, 100, and 1000 for your two sets of parameter(s). Calculate the sample mean, standard deviation, skewness, and kurtosis. Interpret the results.

```
library(e1071)
                                 #number of observations
even.10 <- rbinom(n=10,
              size=1,
                                 #number of trials (size=1 for a Bernoulli distribution)
                                 #probability of success
              prob=.5)
mean(even.10)
                                  #mean
## [1] 0.7
sd(even.10)
                                  #standard deviation
## [1] 0.4830459
skewness(even.10)
                                  #skewness
## [1] -0.7452708
kurtosis(even.10)
                                  #kurtosis
## [1] -1.572857
```

```
even.25 \leftarrow rbinom(n=25,
                                  #number of observations
                                  #number of trials (size=1 for a Bernoulli distribution)
              size=1,
              prob=.5)
                                  #probability of success
mean(even.25)
                                  #mean
## [1] 0.56
sd(even.25)
                                  #standard deviation
## [1] 0.5066228
skewness(even.25)
                                  #skewness
## [1] -0.2273881
kurtosis(even.25)
                                 #kurtosis
## [1] -2.02454
even.100 <- rbinom(n=100,
                                  #number of observations
              size=1,
                                  #number of trials (size=1 for a Bernoulli distribution)
              prob=.5)
                                  #probability of success
mean(even.100)
                                  #mean
## [1] 0.45
sd(even.100)
                                 #standard deviation
## [1] 0.5
skewness(even.100)
                                 #skewness
## [1] 0.198
kurtosis(even.100)
                                 #kurtosis
## [1] -1.9803
even.1000 <- rbinom(n=1000,
                                 #number of observations
                                  #number of trials (size=1 for a Bernoulli distribution)
              size=1,
              prob=.5)
                                  #probability of success
mean(even.1000)
                                  #mean
## [1] 0.467
sd(even.1000)
                                  #standard deviation
## [1] 0.4991595
skewness(even.1000)
                                  #skewness
## [1] 0.1320901
kurtosis(even.1000)
                                  #kurtosis
## [1] -1.984534
```

```
uneven.10 <- rbinom(n=10,
                                  #number of observations
                                  #number of trials (size=1 for a Bernoulli distribution)
                  size=1,
                  prob=.7)
                                  #probability of success
mean(uneven.10)
                                  #mean
## [1] 0.6
sd(uneven.10)
                                 #standard deviation
## [1] 0.5163978
skewness(uneven.10)
                                 #skewness
## [1] -0.3485685
kurtosis(uneven.10)
                                 #kurtosis
## [1] -2.055
uneven.25 <- rbinom(n=25,
                                 #number of observations
                   size=1,
                                 #number of trials (size=1 for a Bernoulli distribution)
                   prob=.7)
                                 #probability of success
mean(uneven.25)
                                  #mean
## [1] 0.76
sd(uneven.25)
                                 #standard deviation
## [1] 0.4358899
skewness (uneven.25)
                                 #skewness
## [1] -1.145243
kurtosis(uneven.25)
                                 #kurtosis
## [1] -0.7121684
uneven.100 \leftarrow rbinom(n=100,
                                 #number of observations
                                 #number of trials (size=1 for a Bernoulli distribution)
                   size=1,
                   prob=.7)
                                 #probability of success
mean(uneven.100)
                                  #mean
## [1] 0.72
sd(uneven.100)
                                  #standard deviation
## [1] 0.4512609
skewness(uneven.100)
                                 #skewness
## [1] -0.9652953
kurtosis(uneven.100)
                                  #kurtosis
## [1] -1.078693
```

```
uneven.1000 <- rbinom(n=1000,
                                  #number of observations
                    size=1,
                                  #number of trials (size=1 for a Bernoulli distribution)
                    prob=.7)
                                  #probability of success
mean(uneven.1000)
                                  #mean
## [1] 0.69
                                  #standard deviation
sd(uneven.1000)
## [1] 0.4627247
skewness(uneven.1000)
                                  #skewness
## [1] -0.8204015
kurtosis (uneven. 1000)
                                  #kurtosis
## [1] -1.328267
```

(c) Generate a random sample of size n = 10 for your two sets of parameter(s). Calculate the method of moments estimator(s) and maximum likelihood estimator(s). In a 1×2 grid, plot a histogram (with bin size 1) of each set of data with (1) the method of moments estimated distribution, (2) the maximum likelihood estimated distribution, and superimpose the true distribution in both.

```
library(ggplot2)
second.moment <- function(data, x.bar){ #function to find the second moment
  temp = 0
  for(i in data){
    temp = temp + (i - x.bar)^2
  return (temp/length(data))
MOM.bernoulli <- function(data){</pre>
                                  #function to find the MOM Estimator for p in a Berno
  mu.hat <- mean(data)</pre>
  var.hat <- second.moment(data,mu.hat)</pre>
  p.hat <- (mu.hat-var.hat)/mu.hat</pre>
  return(p.hat)
# Negative of the Likelihood function
bernoulliLogLik.neg <- function(par, data){
  -sum(dbinom(x=data, prob=par, size=1, log=TRUE))} #Sum is negative, as we are looking for a
MLE.bernoulli <- function(data){</pre>
  MLE <- optim(par = .5,</pre>
                                        # best guess for the parameter
               fn = bernoulliLogLik.neg, # function to minimize
               data = data, # data (an argument to fn)
               method = "Brent",
                                     # required for univariate optimization
               lower = 0,
                                        # lowest possible lambda
               upper = 1)
                                      # reasonable upper bound
  # Note that our mle is hat(lambda) = xbar
  return(MLE$par)
```

```
bernoulli.plot <- function(data){</pre>
  dfActual <- data.frame(matrix(nrow = 2, ncol = 2))</pre>
  colnames(dfActual) <- c("Prop","Value")</pre>
  rownames(dfActual) <- c("0","1")</pre>
  ActualTab <- as.data.frame(data) %>% group_by(data) %>%
    summarize(count = n()) # Calculate the counts
  dfActual[1,1] <- prop.table(ActualTab$count)[1]</pre>
  dfActual[2,1] <- prop.table(ActualTab$count)[2]</pre>
  dfActual[1,2] \leftarrow 0
  dfActual[2,2] <- 1
  dfEstimate <- data.frame(matrix(nrow = 2, ncol = 2))</pre>
  colnames(dfEstimate) <- c("Prop", "Value")</pre>
  rownames(dfEstimate) <- c("0","1")</pre>
  EstimateTab <- as.data.frame(data) %>% group_by(data) %>%
    summarize(count = n()) # Calculate the counts
  dfEstimate[1,1] <- 1-MOM.bernoulli(data)</pre>
  dfEstimate[2,1] <- MOM.bernoulli(data)</pre>
  dfEstimate[1,2] \leftarrow 0
  dfEstimate[2,2] <- 1
  p1 <- ggplot(mapping = aes(y=Prop, x=Value)) +
    geom_col(data = dfActual, position = "dodge", fill = "lightblue") +
    geom_col(data = dfEstimate, aes(group = Value),
              fill = "black", width = 0.01, position = position_dodge(width = 0.9))
  dfActual <- data.frame(matrix(nrow = 2, ncol = 2))</pre>
  colnames(dfActual) <- c("Prop", "Value")</pre>
  rownames(dfActual) <- c("0","1")</pre>
  ActualTab <- as.data.frame(data) %>% group_by(data) %>%
    summarize(count = n()) # Calculate the counts
  dfActual[1,1] <- prop.table(ActualTab$count)[1]</pre>
  dfActual[2,1] <- prop.table(ActualTab$count)[2]</pre>
  dfActual[1,2] <- 0
  dfActual[2,2] <- 1
  dfEstimate <- data.frame(matrix(nrow = 2, ncol = 2))</pre>
  colnames(dfEstimate) <- c("Prop", "Value")</pre>
  rownames(dfEstimate) <- c("0","1")</pre>
  EstimateTab <- as.data.frame(data) %>% group_by(data) %>%
```

```
summarize(count = n()) # Calculate the counts
 dfEstimate[1,1] <- 1-MLE.bernoulli(data)</pre>
 dfEstimate[2,1] <- MLE.bernoulli(data)</pre>
 dfEstimate[1,2] <- 0
 dfEstimate[2,2] <- 1
 p2 <- ggplot(mapping = aes(y=Prop, x=Value)) +
   geom_col(data = dfActual, position = "dodge", fill = "lightblue") +
   geom_col(data = dfEstimate, aes(group = Value),
             fill = "black", width = 0.01, position = position_dodge(width = 0.9))
 return(p1+p2)
even.10 <- rbinom(n=10,
                               #number of observations
                    size=1,
                                   #number of trials (size=1 for a Bernoulli distribution)
                    prob=.5)
                                   #probability of success
MOM.bernoulli(even.10)
## [1] 0.4
MLE.bernoulli(even.10)
## [1] 0.4
bernoulli.plot(even.10)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
uneven.10 <- rbinom(n=10,
                                 #number of observations
                                  #number of trials (size=1 for a Bernoulli distribution)
                    size=1,
                    prob=.7)
                                   #probability of success
MOM.bernoulli(uneven.10)
## [1] 0.6
MLE.bernoulli(uneven.10)
## [1] 0.6
bernoulli.plot(uneven.10)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
```

(d) Generate a random sample of size n=25 for your two sets of parameter(s). Calculate the method of moments estimator(s) and maximum likelihood estimator(s). In a 1×2 grid, plot a histogram (with bin size 1) of each set of data with (1) the method of moments estimated distribution, (2) the maximum likelihood estimated distribution, and superimpose the true distribution in both.

```
even.25 \leftarrow rbinom(n=25,
                                #number of observations
                  size=1,
                                  #number of trials (size=1 for a Bernoulli distribution)
                                  #probability of success
                  prob=.5)
MOM.bernoulli(even.25)
## [1] 0.56
MLE.bernoulli(even.25)
## [1] 0.56
bernoulli.plot(even.25)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
                                 #number of observations
uneven.25 <- rbinom(n=25,
                                   #number of trials (size=1 for a Bernoulli distribution)
                    size=1.
                    prob=.7)
                                    #probability of success
MOM.bernoulli(uneven.25)
## [1] 0.6
MLE.bernoulli(uneven.25)
## [1] 0.6
bernoulli.plot(uneven.25)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
```

(e) Generate a random sample of size n = 100 for your two sets of parameter(s). Calculate the method of moments estimator(s) and maximum likelihood estimator(s). In a 1×2 grid, plot a histogram (with bin size 1) of each set of data with (1) the method of moments estimated distribution, (2) the maximum likelihood estimated distribution, and superimpose the true distribution in both.

```
even.100 <- rbinom(n=100,
                                 #number of observations
                  size=1,
                                 #number of trials (size=1 for a Bernoulli distribution)
                                 #probability of success
                  prob=.5)
MOM.bernoulli(even.100)
## [1] 0.56
MLE.bernoulli(even.100)
## [1] 0.56
bernoulli.plot(even.100)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
uneven.100 <- rbinom(n=100,
                                   #number of observations
                    size=1,
                                   #number of trials (size=1 for a Bernoulli distribution)
                                   #probability of success
                    prob=.7)
MOM.bernoulli(uneven.100)
```

```
## [1] 0.68
MLE.bernoulli(uneven.100)
## [1] 0.68
bernoulli.plot(uneven.100)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
```

(f) Generate a random sample of size n=100 for your two sets of parameter(s). Calculate the method of moments estimator(s) and maximum likelihood estimator(s). In a 1×2 grid, plot a histogram (with bin size 1) of each set of data with (1) the method of moments estimated distribution, (2) the maximum likelihood estimated distribution, and superimpose the true distribution in both.

```
even.1000 <- rbinom(n=1000,
                                   #number of observations
                                 #number of trials (size=1 for a Bernoulli distribution)
                  size=1,
                  prob=.5)
                                 #probability of success
MOM.bernoulli(even.1000)
## [1] 0.5
MLE.bernoulli(even.1000)
## [1] 0.5
bernoulli.plot(even.1000)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
uneven.1000 <- rbinom(n=1000,
                                     #number of observations
                    size=1,
                                   #number of trials (size=1 for a Bernoulli distribution)
                    prob=.7)
                                   #probability of success
MOM.bernoulli(uneven.1000)
## [1] 0.68
MLE.bernoulli(uneven.1000)
## [1] 0.68
bernoulli.plot(uneven.1000)
## Error in as.data.frame(data) %>% group_by(data) %>% summarize(count = n()): could
not find function "%>%"
```

(g) Comment on the results of parts (c)-(f).