Data 556 HW 2

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```
#loading the package that includes rv distributions and ggplot
library(stats)
library(ggplot2)
```

2a.Use simulations to numerically estimate mean & variance of A.

```
#setting a seed so its reproducible
set.seed(99)

nsims=10000
records = rep(0,nsims)

for(i in 1:nsims){
    radius <- runif(1, min = 0, max = 1)
        records[i] <- ((radius^2)*pi)
}

#expectation of the area (the area is the records)
e_a = sum(records)/nsims #1.048
var_a = var(records) #.8768</pre>
```

3a. Use simulations to gain understadning about the distribution of R. Numerically estimate the expected value of R and 1/R.

```
#set seed
set.seed(100)

nsims=10000
records_x = rep(0,nsims)
records_y = rep(0,nsims)
records_ratio = rep(0,nsims)
one_r = rep(0,nsims)

for(i in 1:nsims){
    x <- runif(1, min = 0, max = 1)
    y = 1-x</pre>
```

```
records_x[i] <- x
records_y[i] <- y
records_ratio[i] <- (x/y)
one_r[i] <- (y/x)
}

e_r = sum(records_ratio)/10000 #9.0554
e_r_one = sum(one_r)/10000 #8.438</pre>
```

4a. Let U_1, \ldots, U_n be iid Unif(0,1) and x=max(iids). Use R to numerically estimate E(x).

```
set.seed(100)

nsims = 10000
xrecords = rep(0,nsims)

for(i in 1:nsims){
    x <- runif(1, min = 0, max = 1)
    xrecords[i] <- x
}</pre>
e_r = sum(xrecords)/10000 # .50029
```

5b. Use R to numerically estimate P(X < Y) for X - N(0,1), Y - N(1,5) w X and Y being independent.

```
set.seed(100)

nsims = 10000
records_x_y = rep(0,nsims)

for(i in 1:nsims){
    x <- rnorm(1, mean = 0, sd = 1)
    y <- rnorm(1, mean = 1, sd = 5)
    records_x_y[i] <- (x<y)
}

exy = sum(records_x_y)/nsims # .5759</pre>
```

6b. Use R to calculate the Monte Carlo estimates of mean + SD of x-y.

```
set.seed(100)

nsims=10000
records_x = rep(0,nsims)
records_y = rep(0,nsims)
records_ratio = rep(0, nsims)

for(i in 1:nsims){
    x <- rnorm(1, mean = 69.1, sd = 2.9)
    y <- rnorm(1, mean = 63.7, sd = 2.7)
    records_x[i] <- x
    records_y[i] <- y
    records_ratio[i] <- (x - y)
}

e_r = sum(records_ratio)/10000 # 5.396
sd <- sd(records_ratio) #3.942</pre>
```

6c. What is the probability that a randomly sampled man is taller than a randomly sampled woman?

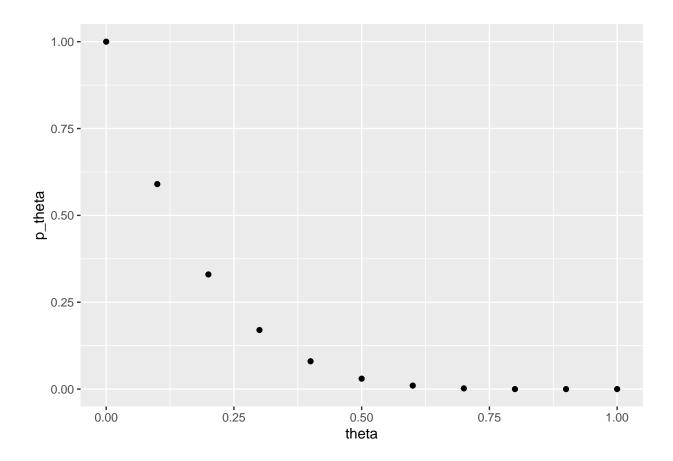
```
pnorm(63.7, mean = 5.4, .2, lower.tail = FALSE)
## [1] 0
```

7b. For y = 0 make a plot for P(theta given y) for each theta within $\{0.0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.0\}$. Make a plot with horizontal axis the 11 values of theta and verticle the P(theta given y).

```
#making a df
theta <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
p_theta <- c(1, .59, .33, .17, .08, .03, .01, .002, 0, 0, 0)

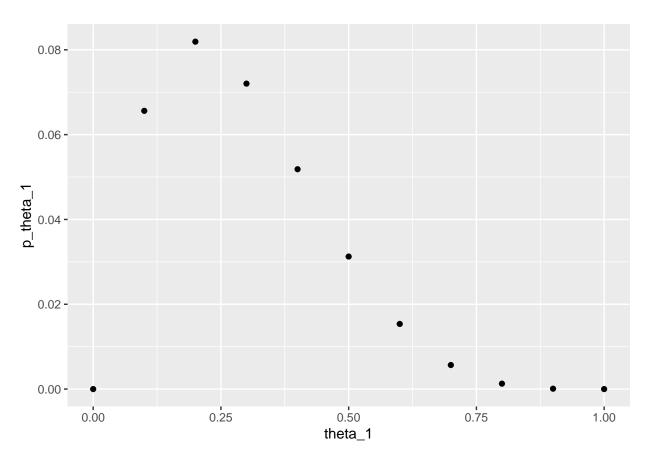
df_theta <- data.frame(theta, p_theta)

ggplot(df_theta, aes(x= theta, y = p_theta)) + geom_point()</pre>
```

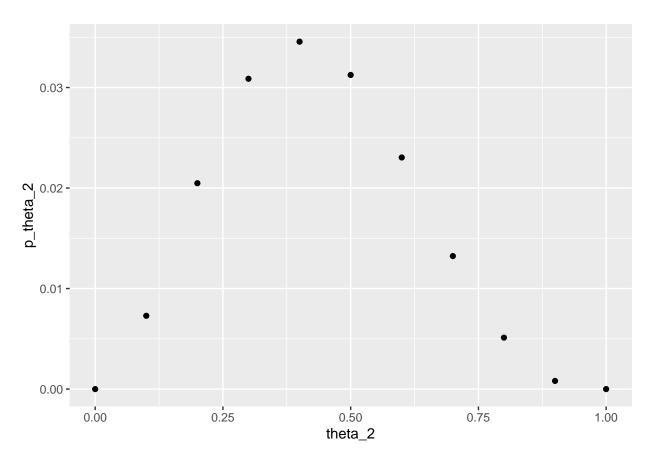


7c. Repeat (b) for each y within $\{1,2,3,4,5\}$.

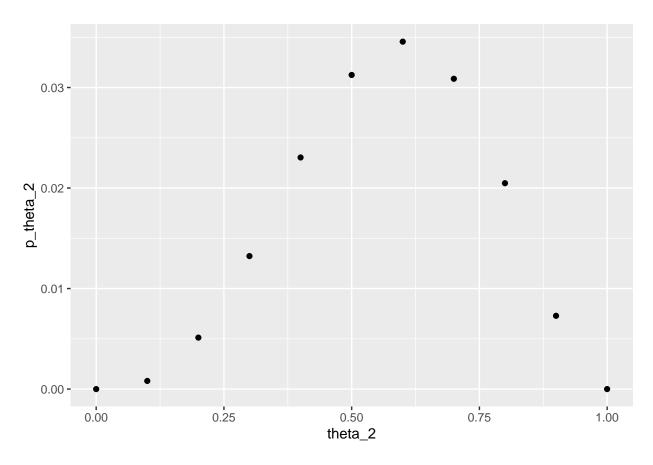
```
#making a df for 1 and graph
theta_1 <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
p_theta_1 <- c((0^1)*(1^4), (.1^1)*(.9^4), (.2^1)*(.8^4), (.3^1)*(.7^4), (.4^1)*(.6^4), (.5^1)*(.5^4),
df_theta_1 <- data.frame(theta, p_theta)
ggplot(df_theta_1, aes(x= theta_1, y = p_theta_1)) + geom_point()</pre>
```



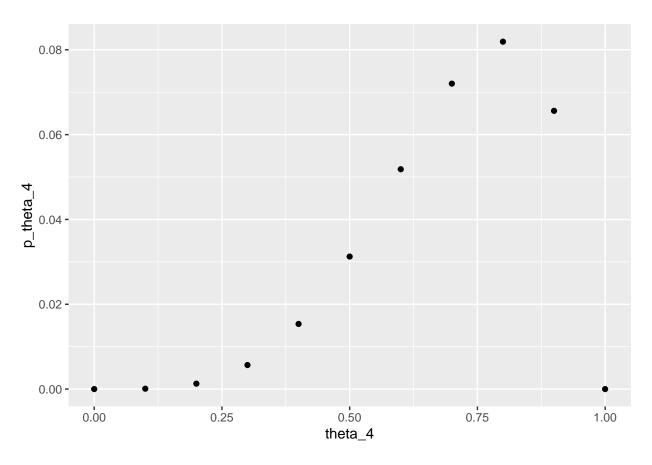
```
#df and graph for 2
theta_2 <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
p_theta_2 <- c((0^2)*(1^3), (.1^2)*(.9^3), (.2^2)*(.8^3), (.3^2)*(.7^3), (.4^2)*(.6^3), (.5^2)*(.5^3),
df_theta_2 <- data.frame(theta, p_theta)
ggplot(df_theta_2, aes(x= theta_2, y = p_theta_2)) + geom_point()</pre>
```



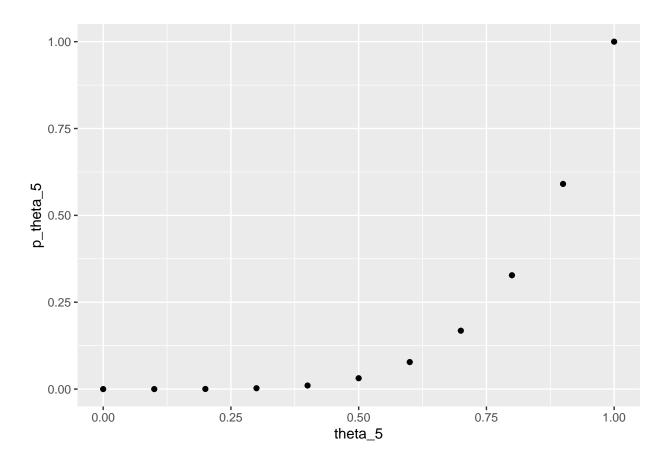
```
#df and graph for 3
theta_2 <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
p_theta_2 <- c((0^3)*(1^2), (.1^3)*(.9^2), (.2^3)*(.8^2), (.3^3)*(.7^2), (.4^3)*(.6^2), (.5^3)*(.5^2),
df_theta_2 <- data.frame(theta, p_theta)
ggplot(df_theta_2, aes(x= theta_2, y = p_theta_2)) + geom_point()
```



```
#df and graph for 4 theta_4 <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1) p_theta_4 <- c((0^4)*(1^1), (.1^4)*(.9^1), (.2^4)*(.8^1), (.3^4)*(.7^1), (.4^4)*(.6^1), (.5^4)*(.5^1), df_theta_4 <- data.frame(theta, p_theta) ggplot(df_theta_4, aes(x= theta_4, y = p_theta_4)) + geom_point()
```



```
#df and graph for 5
theta_5 <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
p_theta_5 <- c((0^5)*(1^0), (.1^5)*(.9^0), (.2^5)*(.8^0), (.3^5)*(.7^0), (.4^5)*(.6^0), (.5^5)*(.5^0),
df_theta_5 <- data.frame(theta, p_theta)
ggplot(df_theta_5, aes(x= theta_5, y = p_theta_5)) + geom_point()</pre>
```



8b. Implement your sampling algoritm in R and use your code to produce a Monte Carlo estimate of P(X lives within (2,3)) where X is a r.v. with a logistic distribution.

```
#setting a seed so its reproducible
set.seed(99)
nsims=10000
records = rep(0, nsims)
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```

```
for(i in 1:nsims){
  x <- runif(1, min = 0, max = 1)</pre>
```

```
xmake_change <- ((1/x) + 1)
xlogistic <- -log(xmake_change, base = exp(1))
records[i] <- (2 < xlogistic & xlogistic < 3)
}
# The probability is zero because non of ours fall between 2 and three?</pre>
```