

A1.R

yuantien

2022-01-21

```
#A1 Yuan Tien
```

```
library(tidyverse)
```

```
## — Attaching packages — tidyverse 1.3.1 —
```

```
## ✓ ggplot2 3.3.5      ✓ purrr 0.3.4
## ✓ tibble 3.1.6       ✓ dplyr 1.0.7
## ✓ tidyr 1.1.4        ✓ stringr 1.4.0
## ✓ readr 2.1.1        ✓ forcats 0.5.1
```

```
## — Conflicts — tidyverse_conflicts() —
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
```

```
getwd()
```

```
## [1] "/Users/yuantien/Desktop/R/613/Data"
```

```
setwd("/Users/yuantien/Desktop/R/613/Data")
getwd()
```

```
## [1] "/Users/yuantien/Desktop/R/613/Data"
```

```
dathh2007 <- read.csv("dathh2007.csv")
dathh07 <- dathh2007
rm(dathh2007)
```

```
#1.1
class(dathh07$idmen)
```

```
## [1] "numeric"
```

```
a <- unique(dathh07$idmen) #find unique value
length(a) #10498
```

```
## [1] 10498
```

```
getwd()
```

```
## [1] "/Users/yuantien/Desktop/R/613/Data"
```

```
#1.2
```

```
dathh05 <- read.csv("dathh2005.csv")
```

```
length(dathh05$mstatus[dathh05$mstatus == "Couple, with Kids"]) #3374
```

```
## [1] 3374
```

```
table(dathh05$mstatus) #redo this with a more convenient way
```

```
##  
## Couple, No kids Couple, with Kids Other Single  
## 2656 3374 275 2663  
## Single Parent  
## 785
```

```
#1.3
```

```
datind08 <- read.csv("datind2008.csv")
```

```
b <- unique(datind08$idind)
```

```
length(b) #it shows 10825, but this individual level data has 25510 obs.
```

```
## [1] 10825
```

```
#1.4
```

```
datind16 <- read.csv("datind2016.csv")
```

```
a <- datind16 %>%
```

```
  filter(age>= 25 & age<=35) %>%
```

```
  nrow()
```

```
a #2765
```

```
## [1] 2765
```

```
#1.5
```

```
datind09 <- read.csv("datind2009.csv")
```

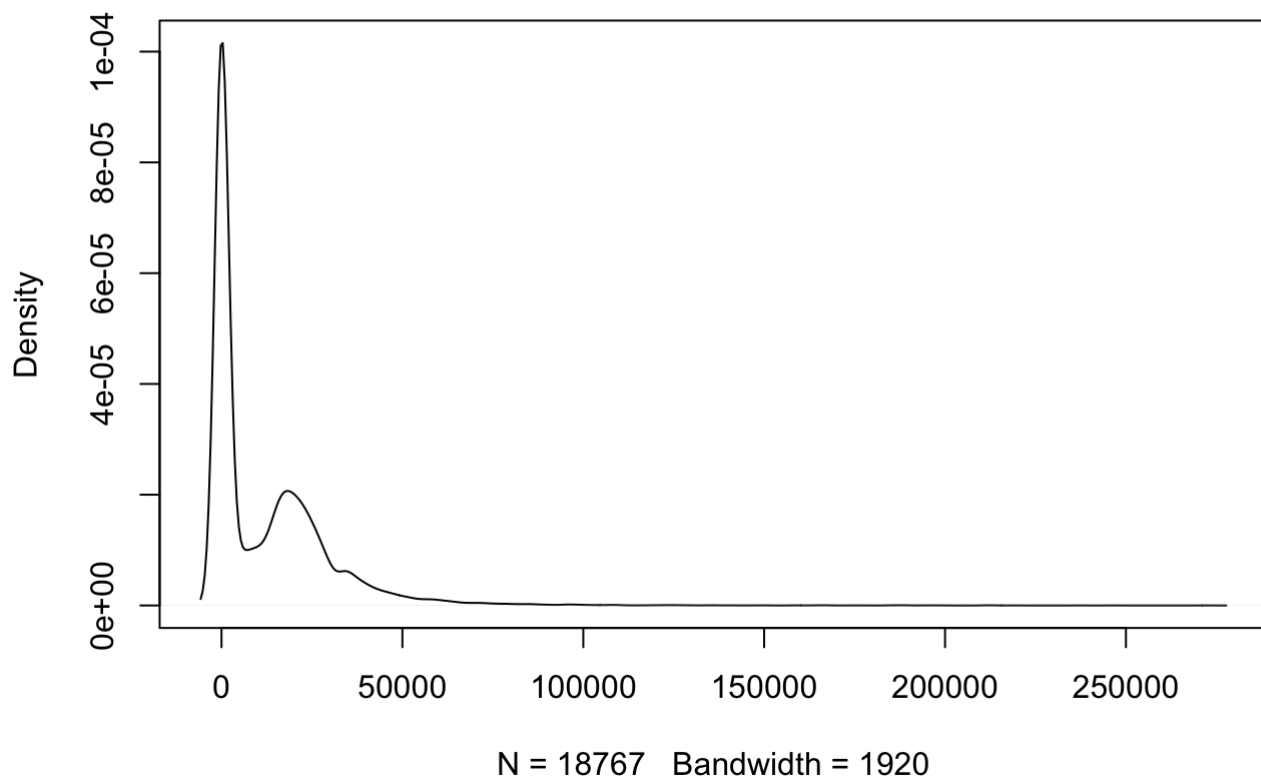
```
CrossTable <- table(datind09$gender, datind09$profession)
```

```
CrossTable
```

```
##  
## 0 11 12 13 21 22 23 31 33 34 35 37 38 42 43 44 45  
## Female 11 30 8 29 63 65 8 68 85 184 50 179 78 258 437 1 153  
## Male 19 57 19 78 213 114 48 98 107 142 59 260 368 110 117 2 95  
##  
## 46 47 48 52 53 54 55 56 62 63 64 65 67 68 69  
## Female 410 82 22 782 27 584 353 696 64 35 29 19 147 120 40  
## Male 340 429 215 169 182 98 101 74 443 520 246 159 237 177 82
```

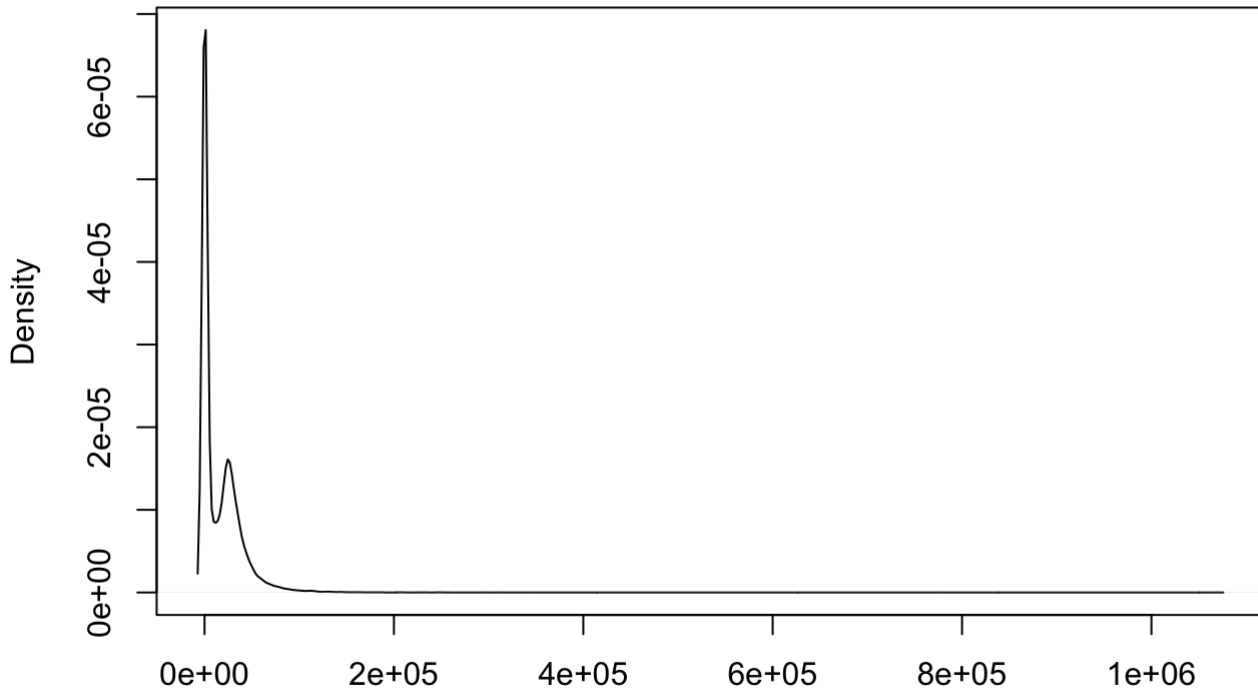
```
#1.6
datind05 <- read.csv("datind2005.csv")
datind19 <- read.csv("datind2019.csv")
plot(density(datind05$wage, na.rm = TRUE)) #plot the distribution
```

density.default(x = datind05\$wage, na.rm = TRUE)



```
plot(density(datind19$wage, na.rm = TRUE))
```

density.default(x = datind19\$wage, na.rm = TRUE)



N = 21421 Bandwidth = 2416

```
inter_decile <- function(x) {  
  quantileX = quantile(x, prob = c(0.1, 0.9))  
  ratio = quantileX[2]/quantileX[1] #because 2nd element represent the 90% and the 1st element represent the 10%  
  return(ratio)  
}  
  
gini <- function(y) {  
  n = length(y)  
  a = 1/(n-1)  
  b = (n+1)  
  c = - 2*((sum((n+1-1:n)*y)))  
  d = sum(y)  
  return(a*(b-c/d))  
} #this is sample gini coefficient. Reference: http://www3.nccu.edu.tw/~jthuang/Gini.pdf page 2  
  
dist_report <- function(x) {  
  return(c(mean = mean(x), sd = sd(x), ratio = inter_decile(x), gini = gini(x)))  
}  
datind05_rm <- na.omit(datind05$wage) #clear out rows with NA in wage  
datind19_rm <- na.omit(datind19$wage)  
datind05_rm <- datind05_rm[datind05_rm != 0] #clear out wage = 0  
datind19_rm <- datind19_rm[datind19_rm != 0]  
dist_report(datind05_rm) #mean = 22443.029, sd = 18076.708, ratio = 8.896, gini = 2.0  
01
```

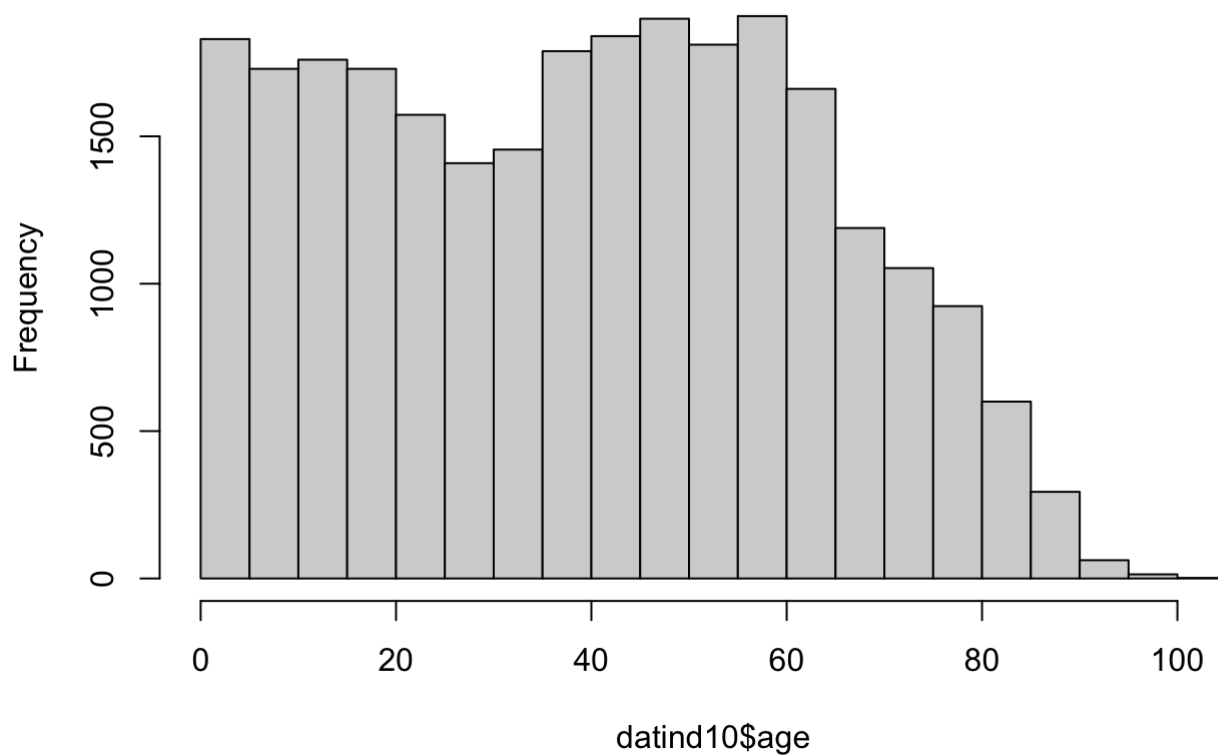
##	mean	sd	ratio.90%	gini
##	22443.029118	18076.708882	8.896525	2.001934

```
dist_report(datind19_rm) #mean = 27578.839, sd = 25107.187, ratio = 13.862, gini = 2.041
```

##	mean	sd	ratio.90%	gini
##	27578.839302	25107.187196	13.862300	2.041109

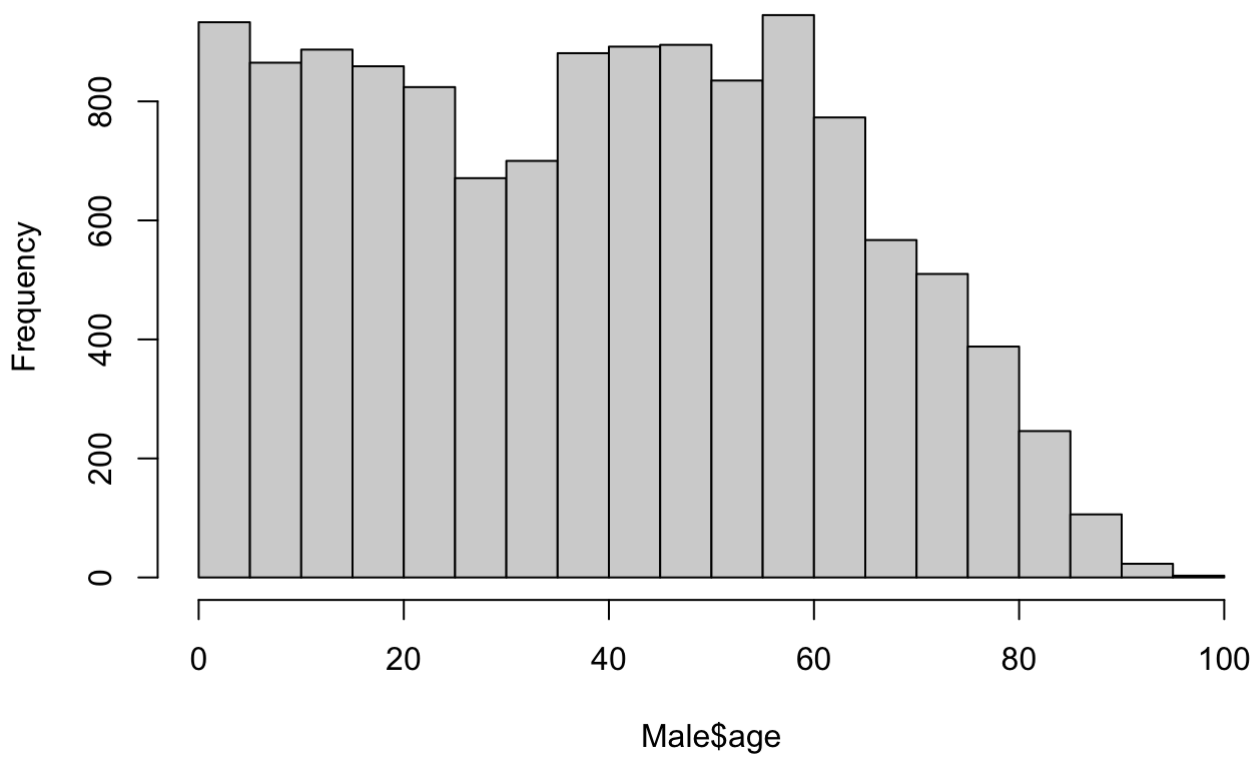
```
#1.7
datind10 <- read.csv("datind2010.csv")
hist(datind10$age)
```

Histogram of datind10\$age



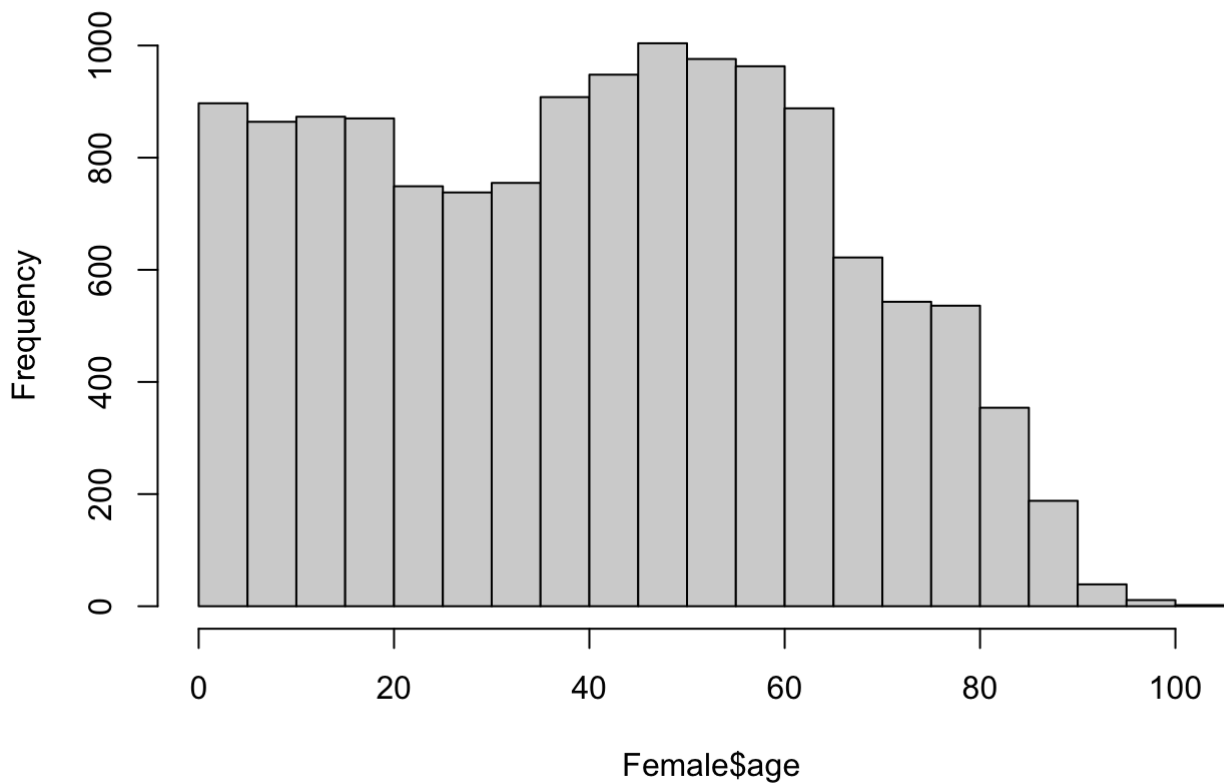
```
Male <- datind10[datind10$gender == "Male",] #remember to put in Comma to select the
rows we like
Female <- datind10[datind10$gender == "Female",]
hist(Male$age)
```

Histogram of Male\$age



```
hist(Female$age)
```

Histogram of Female\$age



```

# the most represented male group in the samples is around 60 years old, while the most represented female group is around 50
#

#1.8
datind11 <- read.csv("datind2011.csv")
dathh11 <- read.csv("dathh2011.csv")
m11 <- datind11 %>%
  inner_join(dathh11, by = "idmen")
#merge household dataset which contains location to ind data
#I use inner join to delete individuals who do not appear in the household dataset
nrow(m11[m11$location == "Paris",]) #3531

```

```
## [1] 3531
```

#Exercise 2

#2.1

```

dind = list.files(pattern="datind")
for (i in 1:length(dind)) assign(dind[i], read.csv(dind[i])) #find data in my file and read multiple files

```

```

Mind <- do.call("rbind", list(datind2004.csv, datind2005.csv, datind2006.csv, datind2007.csv,
                             datind2008.csv, datind2009.csv,
                             datind2010.csv, datind2011.csv, datind2012.csv, datind2013.csv,
                             datind2014.csv, datind2015.csv,
                             datind2016.csv, datind2017.csv, datind2018.csv, datind2019.csv))

```

#2.2

```
dhh = list.files(pattern="dathh")
```

```
for (i in 1:length(dhh)) assign(dhh[i], read.csv(dhh[i]))
```

```

Mhh <- do.call("rbind", list(dathh2004.csv, dathh2005.csv, dathh2006.csv, dathh2007.csv,
                             dathh2008.csv, dathh2009.csv,
                             dathh2010.csv, dathh2011.csv, dathh2012.csv, dathh2013.csv,
                             dathh2014.csv, dathh2015.csv,
                             dathh2016.csv, dathh2017.csv, dathh2018.csv, dathh2019.csv))

```

#2.3

```
colnames(Mind)
```

```
## [1] "X"          "idind"      "idmen"      "year"       "empstat"
## [6] "respondent" "profession" "gender"     "age"        "wage"
```

```
colnames(Mhh)
```

```
## [1] "X"          "idmen"      "year"       "datent"     "myear"      "mstatus"    "move"
## [8] "location"
```

```
y = c(colnames(Mind), colnames(Mhh))
y[duplicated(y) == TRUE] #X, idmen, year
```

```
## [1] "X"      "idmen" "year"
```

```
#find duplicated column names --> find variables that appear in both datasets)
```

```
#2.4
```

```
M <- inner_join(Mhh, Mind, by = c("idmen", "year"))
#I use inner_join because I believe those household ids that appear in both datasets
more reliable data
```

```
#2.5
```

```
M1 <- M #create M1 in case of unexpected accident
members_more_4 = function(x) {
  M2 = M1 %>%
    filter(year == x)
  z = table(M2$idmen)
  y = as.data.frame(z)
  nrow( y[ y$Freq >= 4, ] )
}
# I do this by year
# I create a frequency table by household -> turn to a dataframe -> calculate frequency
year = 2004:2019
more_4_by_year = sapply(year, members_more_4)
sum(more_4_by_year) #37108
```

```
## [1] 37108
```

```
#2.6
```

```
more_1_unemp = function(x) {
  M2 = M1 %>%
    filter(year == x)
  z = table(M2$idmen, M2$empstat)
  y = as.data.frame(z)
  h = y %>%
    filter(Var2 == "Unemployed")
  nrow( h[ h$Freq >=1, ] )
}
more_unemp_year = sapply(year, more_1_unemp)
sum(more_unemp_year) #17241
```

```
## [1] 17241
```

```
#2.7
```

```
unique(M1$profession) #check professions, I am not sure if "X1" "X2" "HO" are professions. Below, I assume they are
```



```
## [1] "67" "56" "" "38" "45" "34" "42" "46" "37" "54" "11" "63" "55" "48" "52"
## [16] "68" "23" "53" "31" "21" "22" "62" "43" "47" "33" "69" "65" "64" "12" "35"
## [31] "13" "44" "00" "X1" "X2" "HO" NA "0" "50" "36" "66" "61"
```

```
twoprof = function(x) {
  M2 = M1 %>%
    filter(year == x)
  z = table(M2$idmen, M2$profession)
  y = as.data.frame.matrix(z)
  nrow(y[y[,2:ncol(y)] >= 2,])
}
#By this, I return the rows that from column2 to column_n_professions where the frequency >= 2
#why from column 2? because column 1 appears to be the freq of NA, the first profession 00 starts with 2nd column

two_prof_year <- sapply(year, twoprof)
sum(two_prof_year) #7509
```

```
## [1] 7509
```

```
#2.8
M1 %>%
  filter(mstatus == "Couple, with Kids") %>%
  nrow() #209382
```

```
## [1] 209382
```

```
#2.9
M1 %>%
  filter(location == "Paris") %>%
  nrow() #51904
```

```
## [1] 51904
```

```
#2.10
most_mem <- function(x) {
  M2 = M1 %>%
    filter(year == x)
  z = table(M2$idmen)
  y = as.data.frame(z)
  y[which.max(y$Freq),] #which.max will find me the index of maximum value
}
most_mem_year <- sapply(year, most_mem)
most_mem_year #the most in 2007 row 9903, and 2010 row 10991. Both have 14 members
```

```
##      [,1]      [,2]      [,3]      [,4]
## Var1 1208045118450100 1607839058220100 1607839058220100 2207811124040100
## Freq 10              11              10              14
##      [,5]      [,6]      [,7]      [,8]
## Var1 1700707001000100 1700707001000100 2510263102990100 1905191114960100
## Freq 10              11              14              10
##      [,9]      [,10]     [,11]     [,12]
## Var1 1905191114960100 2202243098040100 2106457101960100 3000896115750100
## Freq 10              10              9              12
##      [,13]     [,14]     [,15]     [,16]
## Var1 3000896115750100 3000896115750100 3000896115750100 2806477001000100
## Freq 12              12              11              9
```

```
most_mem(2007) #idem: 2207811124040100
```

```
##              Var1 Freq
## 9903 2207811124040100 14
```

```
most_mem(2010) #idem: 2510263102990100
```

```
##              Var1 Freq
## 10991 2510263102990100 14
```

```
#2.11
M2 <- M1 %>%
  filter(year == 2010)
length(unique(M2$idmen)) #11048 households in 2010
```

```
## [1] 11048
```

```
M2 <- M1 %>%
  filter(year == 2011)
length(unique(M2$idmen)) #11360 households in 2011
```

```
## [1] 11360
```

#Exercise 3

#3.1

```
Mhh2 <- Mhh #I only work on the household dataset
```

```
z = table(Mhh2$idmen, Mhh2$year)
```

```
y = as.data.frame.matrix(z)
```

```
# the df is now sorted by idmen and showcase the year of entry (the earliest column with 1) and the year of exit (the last column with 1)
```

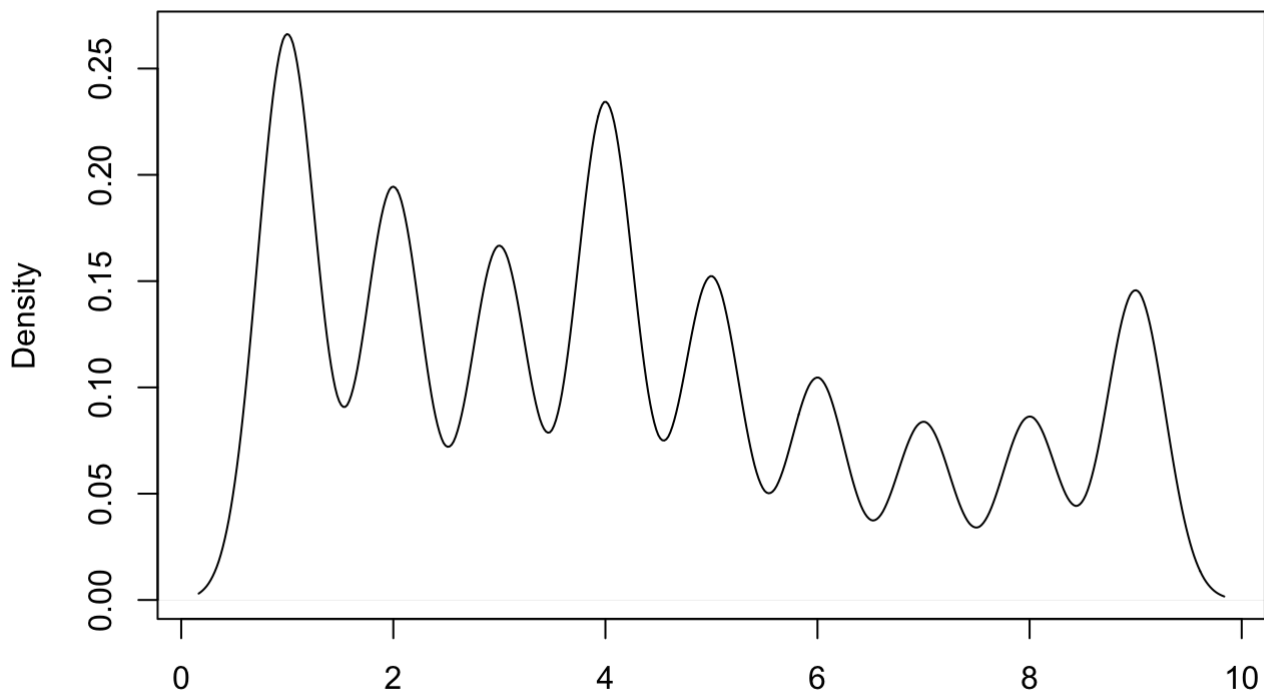
```
y$year_spent <- rowSums(y)
```

```
#the columns are years, and the value is either 1 (participate) or 0 (not participate).
```

```
#So, by summing all the columns I get the years spent in survey
```

```
plot(density(y$year_spent)) #plot the distribution of time spent in the survey
```

density.default(x = y\$year_spent)



N = 41084 Bandwidth = 0.2784

#3.2

```
Mhh2$move_in = Mhh2$year - Mhh2$datent == 0 # create variable "move_in" return 0 -> respondents moved in the same year as the survey
```

```
head(Mhh2$move_in, 10) # first 10 rows, all false
```

```
## [1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

```

#the second part of question requires us to plot the share of "individuals"

h = as.data.frame.matrix( table(M1$idmen, M1$year) ) #create households members by ye
ar and household

dwelling <- function(x) {
  u <- h %>%
    select(members = as.character(x)) #I put as.character to make it work. Without it
R cannot execute my function.
  u$idmen <- rownames(u)
  u$idmen <- as.numeric(u$idmen)

  p <- Mhh2 %>%
    filter(year == x)

  K <- inner_join(p, u, by = "idmen") #here I create members count in 2004, and put i
t in hh dataset
  #duplicate household data by members
  n.times <- K$members
  N <- K[ rep( seq_len( nrow(K) ), n.times), ]
  z <- table(N$year, N$move_in)
  y <- as.data.frame.matrix(z)

  y$ratio <- y$"TRUE"/(y$"TRUE"+y$"FALSE")
  return(y)
}

dwelling(2004) #check if this work, and it works

```

```

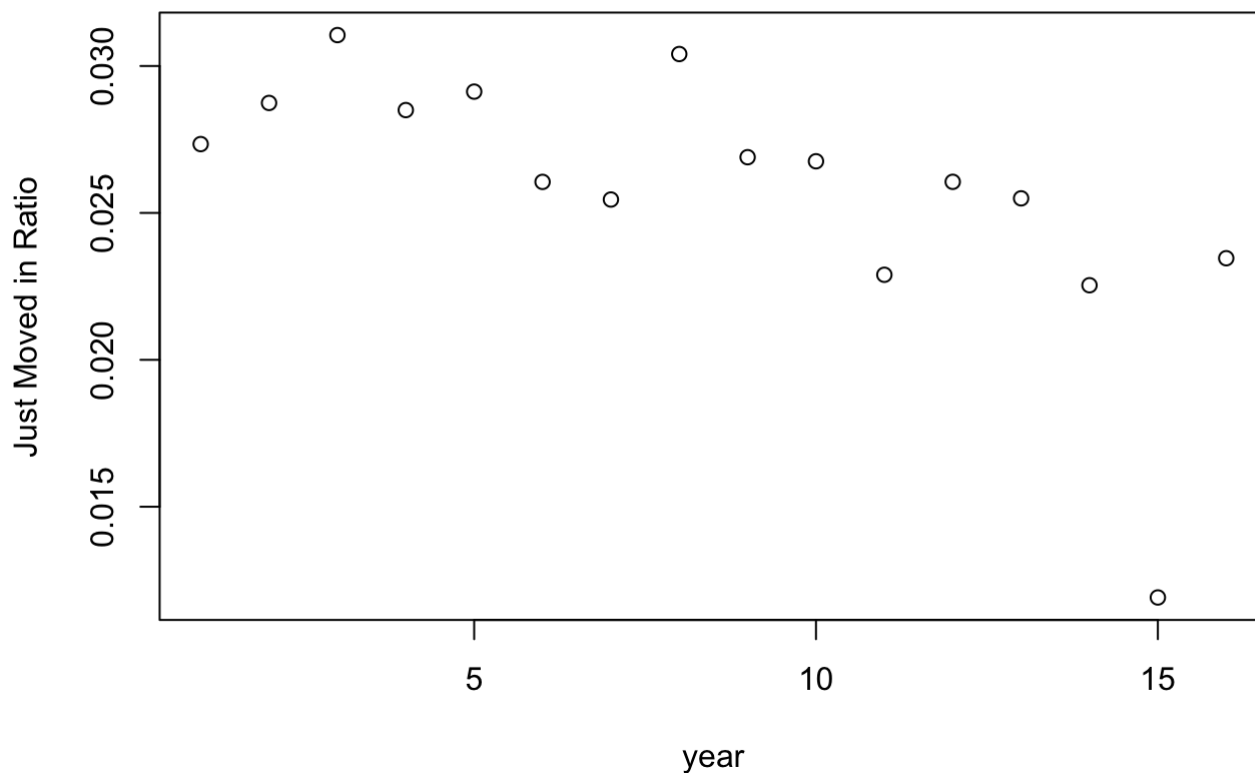
##      FALSE TRUE      ratio
## 2004 21522  605 0.02734216

```

```

dwelling_year <- sapply(year, dwelling)
dwelling_year <- t(dwelling_year)
dwelling_year <- as.data.frame(dwelling_year)
dwelling_year$ratio <- as.numeric(dwelling_year$ratio)
plot(dwelling_year$ratio, xlab = "year", ylab = "Just Moved in Ratio")

```



```
#Below is the code for plotting the "share of households" instead of individuals
#z <- table(Mhh2$year, Mhh2$move_in)
#y <- as.data.frame.matrix(z)
#y

#y$ratio <- y$"TRUE"/y$"FALSE"

#plot(y$ratio, xlab = "year", ylab = "just moved in ratio")
#axis(1, 4:19)

#3.3
#household migrated at the year of survey

Mhhold <- Mhh2 %>%
  filter(year <= 2014)
Mhhold$mig_survey = Mhhold$year - Mhhold$myear == 0

Mhhnew <- Mhh2 %>%
  filter(year >= 2015)

Mhhnew$mig_survey = Mhhnew$move == 2
#By this method I exert that when the household reports migration in year x, I assume
the whole family migrate that year

Mhh3 <- rbind(Mhhold, Mhhnew)
head(Mhh3$mig_survey, 10) #all false for the first 10 rows
```

```
## [1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

```

migrate <- function (x){
  u <- h %>%
    select(members = as.character(x))
  u$idmen <- rownames(u)
  u$idmen <- as.numeric(u$idmen)
  p <- Mhh3 %>% #Mhh3 contains migration data I created
    filter(year == x)

  K <- inner_join(p, u, by = "idmen")

  n.times <- K$members
  N <- K[rep(seq_len(nrow(K)), n.times),] #by this time I have individuals' household
data
  z <- table(N$year, N$mig_survey)
  y <- as.data.frame.matrix(z)

  y$ratio <- y$"TRUE"/(y$"FALSE"+y$"TRUE")
  return(y)
}
migrate(2004) #check if my function works

```

```

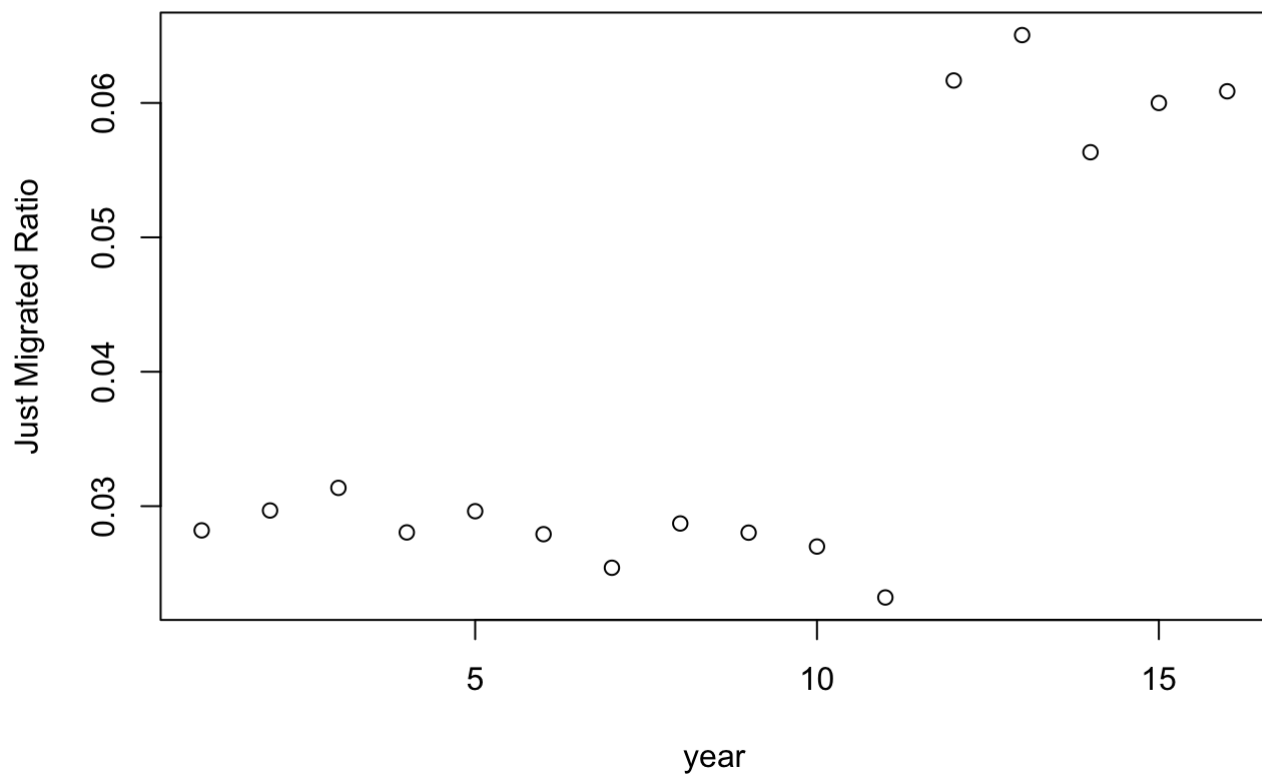
##      FALSE TRUE      ratio
## 2004 21091  612 0.02819887

```

```

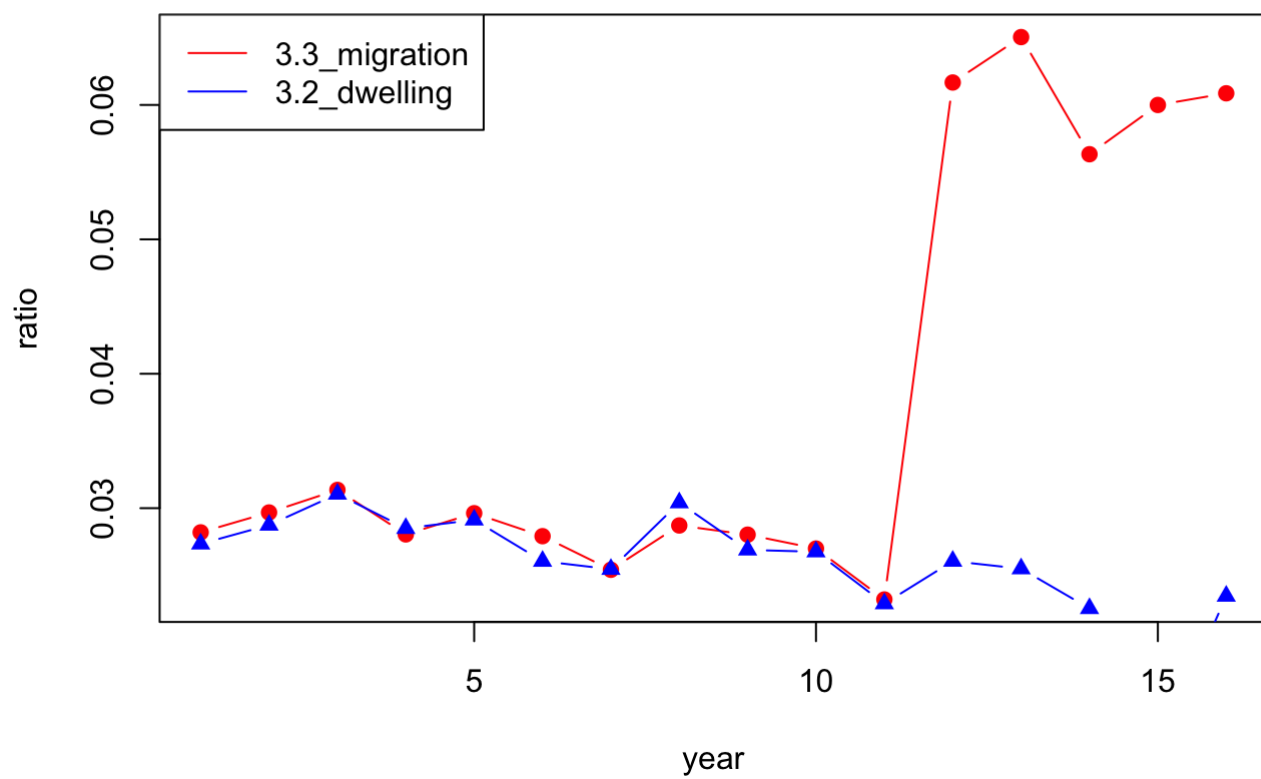
migrate_year <- sapply(year, migrate)
migrate_year <- as.data.frame(t(migrate_year))
migrate_year$ratio <- as.numeric(migrate_year$ratio)
plot(migrate_year$ratio, xlab = "year", ylab = "Just Migrated Ratio")

```



#3.4

```
plot(migrate_year$ratio, type = "b", pch = 19, col = "red", xlab = "year", ylab = "ratio")
lines(dwelling_year$ratio, type = "b", pch = 17, col = "blue") #lines() is used to add a line on a plot
legend("topleft", c("3.3_migration", "3.2_dwelling"), lty = c(1,1), col = c("red", "blue"))
```



I prefer method in 3.2 because there is a unnatural spike for the method in 3.3 during 2015.

One reason for this is the change of data collection method.

The newly introduced "move" data captures who have moved since the last survey

while "datent" captures whether they move the same year as they filling the survey

Suppose John responded in 2013 and 2018, and he moved in 2017. In 2018, we would say he did not move for the 3.2 method

However, we would code he have moved for the variable "move". If re-entries are plenty, the spike for method in 3.3 can be caused by this situation.

Consequently, if we want to know whether the respondent has just migrated, method in 3.2 would be better.

#3.5

```
x <- Mhh3%>%
  select(idmen, move_in)
Mig <- M1 %>%
  left_join(x, by = "idmen") %>%
  filter(move_in == TRUE)

Mig2 <- Mig[ order(Mig$idmen, Mig$idind, Mig$empstat), ] #sort by idmen, idind, and empstat
Mig2 <- Mig2 %>%
  select(idmen, idind, year, empstat, profession)

try <- function (x) {
  Mig2 %>%
    filter(idind == x) %>%
    mutate(change = length(unique(empstat)) >= 2) # because 2 represents change in condition (1 is no change)
}

tryls <- sapply(unique(Mig2$idind), try)
tryMatrix <- as.data.frame.matrix(t(tryls))
tryMatrix$change <- as.vector(tryMatrix$change)

for (i in 1: nrow(tryMatrix)) {
  tryMatrix$change2[i] = sample(unlist(tryMatrix$change[i]), 1)
}
#I use sample because "change" is either all true or all false
#right now, I have compiled "whether an individual changes their profession" in variable change 2

full <- function (x) {
  Mig2 %>%
    filter(idind == x) %>%
    mutate(change = length(unique(empstat)) >= 2 | length(unique(profession)) >= 2)
}

fullls <- sapply(unique(Mig2$idind), full)
fullMx <- as.data.frame.matrix(t(fullls))
fullMx$change <- as.vector(fullMx$change)

for (i in 1: nrow(fullMx)) {
  fullMx$change2[i] = sample(unlist(fullMx$change[i]), 1) #I use sample because "chan
```

```

ge" is either all true or all false
}
# I have complied "whether an individual changes profession or changes employment sta
tus in change2
# Note, in this way if one couple where the husband fills in their household record u
nder one id,
# and the couple's professions are different, I may have coded it as change in profes
sion.
# Overall, I may have overestimated the number.

for (i in 1: nrow(fullMx)) {
  fullMx$idmen[i] = sample(unlist(fullMx$idmen[i]), 1) #turn the list idmen to a vect
or
}

answer <- fullMx %>%
  filter(change2 == TRUE)
length(unique(answer$idmen)) #find the unique households that experience this situati
on

```

```
## [1] 3090
```

#Exercise 4

```

entry_exit <- function (x) {
  a <- M1 %>%
    filter(idind == x) %>%
    arrange(year)%>%
    mutate(entry = year[1])
  a$exit = year[nrow(a)]
  a <- a%>%
    select(idind, entry, exit)
  return(a)
} #this function creates a data.frame for every individual.
#Since the data.frame is already sorted by year, the year of the first row must be en
try, and the last must be exit
#By this I rule out re-entry and multiple exits.

entry_exit(M1$idind[678]) #just testing my function

```

```

##           idind entry exit
## 1 1.120274e+18  2004 2005
## 2 1.120274e+18  2004 2005

```

```

a <- lapply(unique(M1$idind), entry_exit)
b <- bind_rows(a)
y <- b[!duplicated(b),] #clean out duplicated individual observation
z <- table(y$exit) #this lists exits across years

d <- as.data.frame(z)

active <- function(x) {
  length( which ( M1$year == x))
}
d$active <- sapply(year, active)

d$attrition <- d$Freq/d$active
#Report your final result as a table in proportions.
d

```

```

##      Var1 Freq active attrition
## 1  2004 3631  22144 0.16397218
## 2  2005 4758  24241 0.19627903
## 3  2006 2888  24940 0.11579791
## 4  2007 4530  25907 0.17485622
## 5  2008 1960  25510 0.07683261
## 6  2009 3136  25611 0.12244739
## 7  2010 1138  26528 0.04289807
## 8  2011 3528  27071 0.13032396
## 9  2012 1826  28534 0.06399383
## 10 2013 1822  26353 0.06913824
## 11 2014  380  26787 0.01418599
## 12 2015 2176  26644 0.08166942
## 13 2016  310  26647 0.01163358
## 14 2017  908  25402 0.03574522
## 15 2018  882  24698 0.03571139
## 16 2019 1594  26484 0.06018728

```