Yufei_HW3

R Markdown

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When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
#Section A
#Question 1
library(tidyverse)
## - Attaching packages -
                                                        tidyvers
e 1.2.1 —
## ✓ ggplot2 3.2.1
                      ✓ purrr 0.3.3

✓ dplyr 0.8.3

## ✓ tibble 2.1.3
                      ✓ stringr 1.4.0
## ✓ tidyr 1.0.0
           1.3.1
                      ✓ forcats 0.4.0
## ✔ readr
## -- Conflicts -
                                                   - tidyverse conf
licts() —
## * dplyr::filter() masks stats::filter()
```

* dplyr::lag() masks stats::lag()

```
library(dplyr)
#create the area code column
set.seed(1050)
Area code1 <-sample(LETTERS, 1000, replace=TRUE)</pre>
Area code2 <-sample(LETTERS, 1000, replace=TRUE)</pre>
#create the company column
set.seed(1050)
Company1<- sample("Alpha", 1000, replace = TRUE)</pre>
Company2<- sample("Beta", 1000, replace = TRUE)</pre>
#create the employee height column
set.seed(1005)
height1 <- sample(rnorm(1000, mean=160, sd=5), replace = TRUE)
height2 <- sample(rnorm(1000, mean=170, sd=5), replace=TRUE)</pre>
#create the data frame
df1 <- tibble("Area Code"=Area code1, "Company"=Company1, "Employee H
eight"=height1)
df2 <- tibble("Area Code"=Area code2, "Company"=Company2, "Employee H
eight"=height2)
df3 <-full join(df1, df2)
```

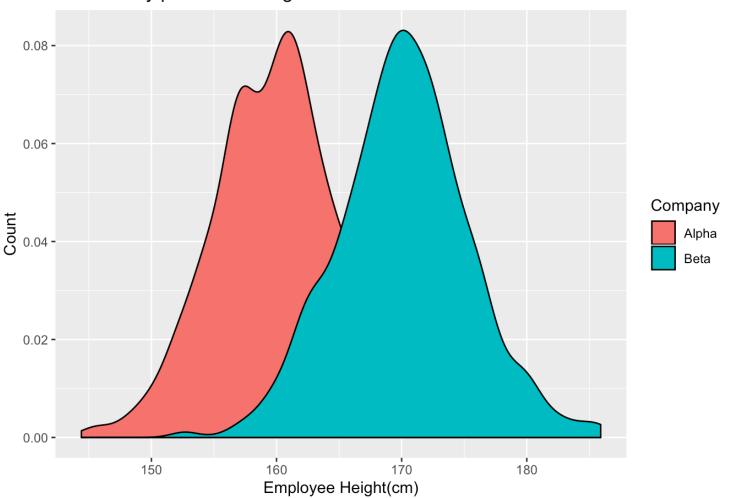
```
## Joining, by = c("Area_Code", "Company", "Employee_Height")
```

df3

```
## # A tibble: 2,000 x 3
      Area Code Company Employee Height
##
##
      <chr>
                <chr>
                                    <dbl>
                Alpha
                                     160.
##
    1 Z
    2 W
                Alpha
##
                                     158.
##
   3 H
                Alpha
                                     159.
##
   4 S
                Alpha
                                     156.
##
   5 P
                Alpha
                                     163.
                Alpha
## 6 Z
                                     160.
                Alpha
##
                                     162.
   7 M
                                     159.
##
   8 D
                Alpha
## 9 Z
                Alpha
                                     158.
## 10 B
                Alpha
                                     156.
## # ... with 1,990 more rows
```

```
plot1 <- ggplot(data=df3)+
   geom_density(mapping=aes(x=df3$Employee_Height,fill=Company))+
scale_fill_discrete() +
scale_y_continuous() +
xlab("Employee Height(cm)")+
ylab("Count")+
   ggtitle("The density plot on the height")</pre>
```

The density plot on the height



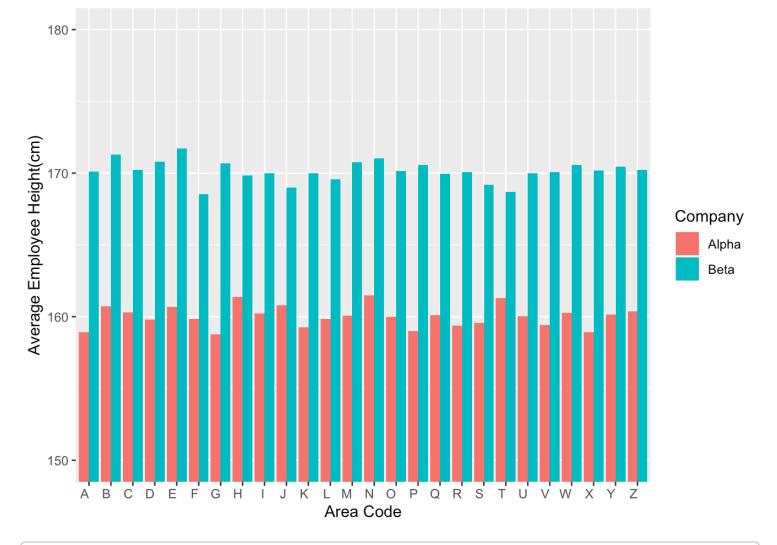
```
# Question 2
#part 1
#create the new datafram
# caculate the average height for each company
target1 <-group_by(df1,Company)
target2 <- summarize(target1, AvgHeight=mean(df1$Employee_Height,na.
rm=TRUE))
target2</pre>
```

```
df4 <- full_join(target2, target4, by = c("Company", "AvgHeight"))
df4</pre>
```

```
#part 2
new <- df3 %>% group_by(Company,Area_Code) %>%
  summarise(AvgHeight=mean(Employee_Height,na.rm=TRUE))
new
```

```
## # A tibble: 52 x 3
## # Groups:
               Company [2]
      Company Area Code AvgHeight
##
##
      <chr>
              <chr>
                             <dbl>
##
   1 Alpha
                              159.
              Α
##
    2 Alpha
                              161.
              В
##
    3 Alpha
              C
                              160.
##
   4 Alpha
                              160.
              D
   5 Alpha
                              161.
##
              E
   6 Alpha
##
              F
                              160.
   7 Alpha
##
              G
                              159.
##
   8 Alpha
              Η
                              161.
##
   9 Alpha
                              160.
              Ι
## 10 Alpha
              J
                              161.
## # ... with 42 more rows
```

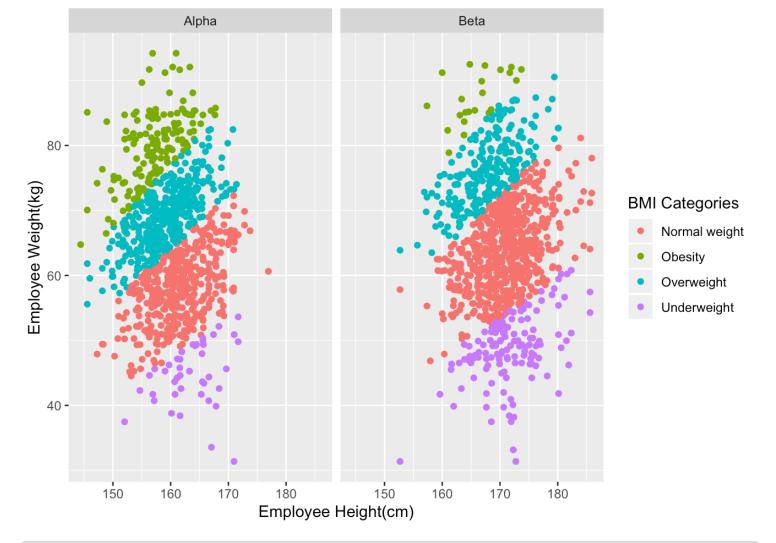
```
#plot the target chart
plot2 <- ggplot(data=new)+
   geom_bar(mapping=aes(x=Area_Code,y=AvgHeight,fill=Company),stat="i
dentity",position ="dodge")+
   scale_x_discrete() +
scale_fill_discrete() +
scale_y_continuous() +
theme(axis.text.x=element_text(angle = 360, hjust = 1))+
coord_cartesian(ylim=c(150,180))+
xlab("Area Code")+
ylab("Average Employee Height(cm)")
plot2</pre>
```



```
#Question 3
#create the random employee weight and add it to the previous datafr
ame
set.seed(1000)
df5<- mutate(df3, "Employee_Weight(kg)"=sample(rnorm(2000,mean=65,sd
=10), replace = TRUE))
df6<- mutate(df5,"BMI"=`Employee_Weight(kg)`/((Employee_Height/100)^
2))
df7<- mutate(df6,BMI_Categories= ifelse(BMI <=18.5,"Underweight",ife
lse(BMI>18.5 & BMI <= 25,"Normal weight",ifelse(BMI >25 & BMI<=30 ,"
Overweight", ifelse(BMI >30, "Obesity","")))))
df7
```

```
## # A tibble: 2,000 x 6
      Area Code Company Employee Height `Employee Weight... BMI BMI
##
Categories
##
      <chr>
                <chr>
                                   <dbl>
                                                      <dbl> <dbl> <chr
>
##
    1 Z
                Alpha
                                    160.
                                                       66.8
                                                             26.1 Over
weight
##
    2 W
                Alpha
                                    158.
                                                       64.9
                                                             25.9 Over
weight
##
    3 H
                Alpha
                                    159.
                                                       66.2
                                                             26.4 Over
weight
## 4 S
                Alpha
                                    156.
                                                       81.8
                                                             33.7 Obes
ity
## 5 P
                Alpha
                                    163.
                                                             32.1 Obes
                                                       85.0
ity
## 6 Z
                Alpha
                                    160.
                                                       50.5
                                                             19.7 Norm
al weight
## 7 M
                Alpha
                                    162.
                                                       79.2
                                                             30.1 Obes
ity
## 8 D
                Alpha
                                    159.
                                                             19.6 Norm
                                                       49.3
al weight
## 9 Z
                Alpha
                                    158.
                                                       61.8
                                                             24.9 Norm
al weight
## 10 B
                                    156.
                                                       71.1
                                                             29.2 Over
                Alpha
weight
## # ... with 1,990 more rows
#create the target plot
df7 %>% ggplot(mapping=aes(df7$Employee Height,df7$`Employee Weight(
```

```
#create the target plot
df7 %>% ggplot(mapping=aes(df7$Employee_Height,df7$`Employee_Weight(
kg)`))+
   geom_point(mapping=aes(color=BMI_Categories))+
   facet_grid(.~Company)+
   xlab("Employee Height(cm)")+
   ylab("Employee Weight(kg)")+
   labs(color="BMI Categories")
```



```
#Section B
#Problem 1
library(tidyverse)
library(plyr)
```

```
## -----
```

```
## You have loaded plyr after dplyr - this is likely to cause proble
ms.
## If you need functions from both plyr and dplyr, please load plyr
first, then dplyr:
## library(plyr); library(dplyr)
```

```
## -----
```

```
##
## Attaching package: 'plyr'
## The following objects are masked from 'package:dplyr':
##
##
       arrange, count, desc, failwith, id, mutate, rename, summarise
##
       summarize
## The following object is masked from 'package:purrr':
##
##
       compact
library(dplyr)
library(haven)
#upload the dataset
dataset1 <- read xpt('https://wwwn.cdc.gov/Nchs/Nhanes/2015-2016/DE</pre>
MO I.XPT')
#filter the ratio without missing values
new111 <- dataset1[!is.na(dataset1$INDFMPIR),]</pre>
#remove the ratio's decimals
new111$INDFMPIR <- trunc(new111$INDFMPIR)</pre>
#count1
dataset2 <- group by(new111,RIDRETH1,INDFMPIR)</pre>
```

dataset3 <- dataset2 %>% dplyr::count(RIDRETH1, name="count1")

dataset3 %>% dplyr::filter(!is.na(INDFMPIR))

```
## # A tibble: 30 x 3
## # Groups:
                RIDRETH1, INDFMPIR [30]
      RIDRETH1 INDFMPIR count1
##
         <dbl>
                   <dbl>
##
                             666
##
    1
              1
                        0
##
    2
              1
                        1
                             512
##
    3
              1
                        2
                             265
                        3
##
    4
              1
                             98
   5
              1
                        4
##
                              55
                        5
##
    6
              1
                             69
##
   7
              2
                        0
                             343
##
    8
              2
                        1
                             351
##
              2
                        2
    9
                             156
              2
## 10
                        3
                             117
## # ... with 20 more rows
```

caculate the proportion of each ethnic families among all families

dataset4 <- plyr::ddply(dataset3,.(INDFMPIR),transform,prop1=count1/</pre>

at each annual family income value

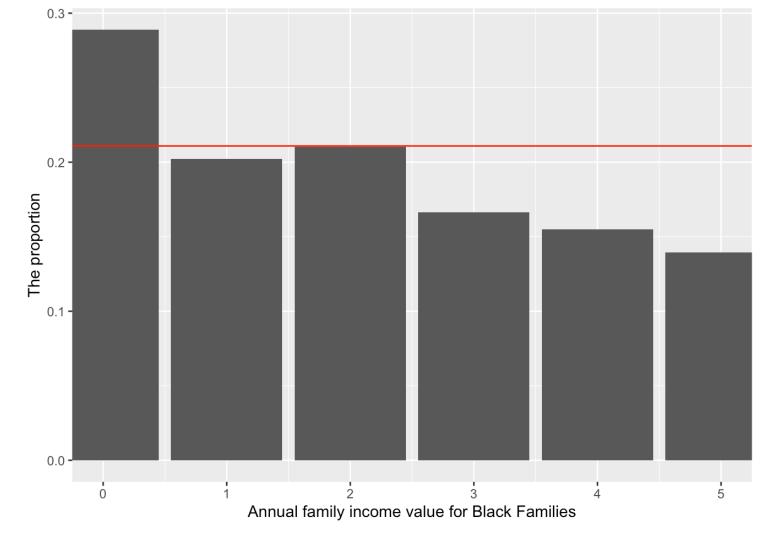
```
sum(count1))
dataset4 %>% dplyr::filter(!is.na(INDFMPIR)) %>% group by( RIDRETH1)
## # A tibble: 30 x 4
## # Groups: RIDRETH1 [5]
##
      RIDRETH1 INDFMPIR count1 prop1
##
         <dbl>
                   <dbl> <int> <dbl>
##
                       0
                             666 0.288
    1
              1
##
    2
              2
                             343 0.148
                       0
##
    3
              3
                       0
                             367 0.159
##
    4
              4
                       0
                             669 0.289
##
                             270 0.117
    5
              5
                       0
                             512 0.207
##
    6
              1
                       1
##
   7
              2
                       1
                             351 0.142
##
    8
              3
                       1
                             783 0.316
##
                       1
                             501 0.202
    9
## 10
              5
                       1
                             330 0.133
## # ... with 20 more rows
```

```
#caculate the proportion of each ethnic families among all families
dataset5 <- group by(new111,RIDRETH1)</pre>
dataset6 <- dataset5 %>% dplyr::count(RIDRETH1, name="count2")
library(data.table)
##
## Attaching package: 'data.table'
## The following objects are masked from 'package:dplyr':
##
##
      between, first, last
## The following object is masked from 'package:purrr':
##
##
      transpose
setDT(dataset6)[,prop2:=count2/sum(count2)]
dataset6
## RIDRETH1 count2 prop2
            1 1665 0.1866801
## 1:
## 2:
           2 1132 0.1269201
## 3:
            3 2877 0.3225698
       4 1881 0.2108981
## 4:
            5 1364 0.1529319
## 5:
#get the target data frame
dataset7 <- full join(dataset4,dataset6)</pre>
## Joining, by = "RIDRETH1"
dataset7 %>% dplyr::filter(!is.na(INDFMPIR)) %>% select(RIDRETH1,IND
```

FMPIR,prop2,prop1)

```
##
      RIDRETH1 INDFMPIR
                              prop2
                                         prop1
## 1
              1
                       0 0.1866801 0.28768898
## 2
              2
                       0 0.1269201 0.14816415
## 3
              3
                       0 0.3225698 0.15853132
              4
                       0 0.2108981 0.28898488
## 4
## 5
              5
                       0 0.1529319 0.11663067
## 6
              1
                       1 0.1866801 0.20670166
              2
                       1 0.1269201 0.14170367
## 7
              3
## 8
                       1 0.3225698 0.31610820
## 9
              4
                       1 0.2108981 0.20226080
## 10
              5
                       1 0.1529319 0.13322568
## 11
              1
                       2 0.1866801 0.18088737
## 12
              2
                       2 0.1269201 0.10648464
## 13
              3
                       2 0.3225698 0.35631399
## 14
              4
                       2 0.2108981 0.21023891
## 15
              5
                       2 0.1529319 0.14607509
## 16
              1
                       3 0.1866801 0.11475410
## 17
              2
                       3 0.1269201 0.13700234
## 18
              3
                       3 0.3225698 0.41803279
## 19
              4
                       3 0.2108981 0.16627635
## 20
              5
                       3 0.1529319 0.16393443
## 21
              1
                       4 0.1866801 0.09353741
## 22
              2
                       4 0.1269201 0.12074830
## 23
              3
                       4 0.3225698 0.44557823
## 24
                       4 0.2108981 0.15476190
              4
## 25
              5
                       4 0.1529319 0.18537415
## 26
              1
                       5 0.1866801 0.05655738
## 27
              2
                       5 0.1269201 0.07704918
## 28
                       5 0.3225698 0.48032787
              3
                       5 0.2108981 0.13934426
## 29
              4
## 30
              5
                       5 0.1529319 0.24672131
```

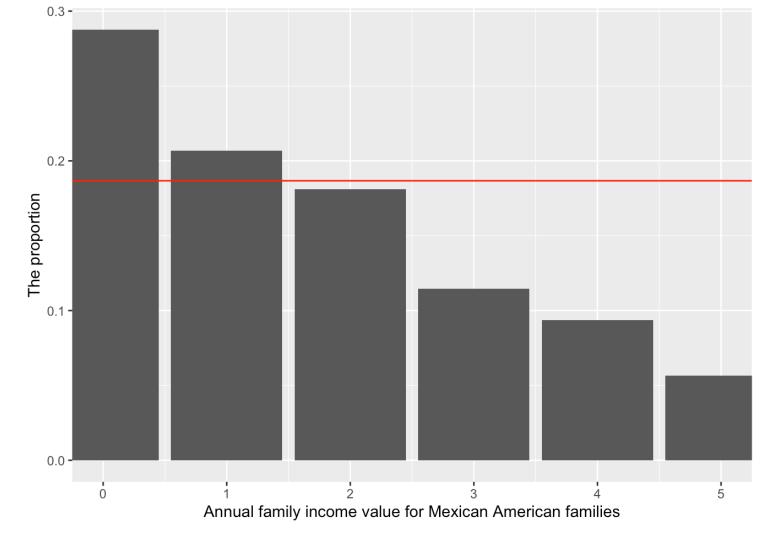
```
#plot1 for black families (4)
dataset7 %>% dplyr::filter(RIDRETH1==4) %>%
    ggplot()+
    geom_bar(mapping=aes(INDFMPIR,prop1),stat="identity")+
    coord_cartesian(xlim=c(0:5))+
    geom_hline(yintercept=0.2108981,color="red")+
    xlab("Annual family income value for Black Families")+
    ylab("The proportion")
```



Answer

Most balck families are under the poverty. The proportions of the annual family income values (0) are above the proportion of famility over all familities, and the proportion of "0" annual family income value is the highest number in the plot.

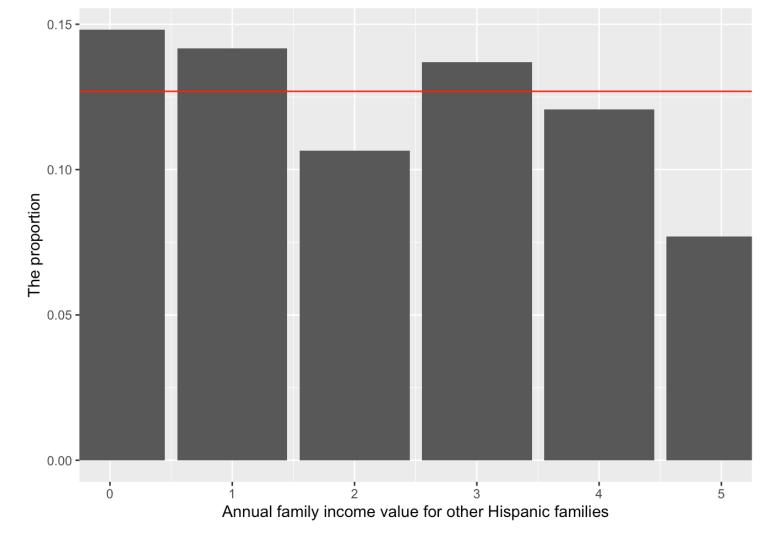
```
#plot2 for Mexican American families (1)
dataset7 %>% dplyr::filter(RIDRETH1==1) %>%
   ggplot()+
   geom_bar(mapping=aes(INDFMPIR,prop1),stat="identity")+
   coord_cartesian(xlim=c(0:5))+
   geom_hline(yintercept=0.1866801,color="red")+
   xlab("Annual family income value for Mexican American families")+
   ylab("The proportion")
```



Answer

Most Mexican American families are under the poverty. The proportions of the annual family income values (0 and 1) are above the proportion of famility over all familities, and the proportion of "0" annual family income value is the highest number in the plot.

```
#plot3 for other Hispanic families (2)
dataset7 %>% dplyr::filter(RIDRETH1==2) %>%
    ggplot()+
    geom_bar(mapping=aes(INDFMPIR,prop1),stat="identity")+
    coord_cartesian(xlim=c(0:5))+
    geom_hline(yintercept=0.1269201,color="red")+
    xlab("Annual family income value for other Hispanic families")+
    ylab("The proportion")
```



Answer

Most other Hispanic families are over poverty. The proportions of the annual family income values (0,1,3) are above the proportion of famility over all familities, and the proportion of "0" annual family income value is the highest number in the plot.
