# homework2

## xbw

## 2025-09-19

1. a.

```
\texttt{ca\_pa} \gets \texttt{read.csv}(\texttt{"E:/postgraduate/data science/mynotes/data/calif\_penn\_2011.csv"})
```

b.

```
nrow(ca_pa) #[1] 11275
```

```
## [1] 11275
```

```
ncol(ca_pa) #[1] 34
```

```
## [1] 34
```

## c.统计每一列中元素是缺失值(NA)的数量

d.

```
ca_pa_omitna <- na.omit(ca_pa)
```

#### e.omitted 10576 rows

f.(c)和(e)的答案不一致,因为(c)统计的是每一列有多少个元素是na,而(e)删除的是有na元素的行

2. a.

```
install.packages("tidyverse")
```

```
\mbox{\tt \#\#} package 'tidyverse' successfully unpacked and MD5 sums checked \mbox{\tt \#\#}
```

##

## The downloaded binary packages are in

 $\verb| ## C:\Users\\78471\\AppData\\Local\\Temp\\RtmpKa7YJP\\downloaded\_packages$ 

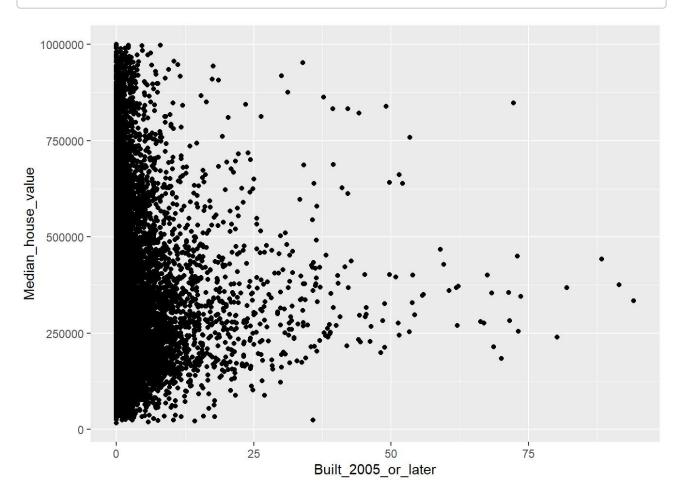
library(tidyverse)

```
## —— Attaching core tidyverse packages -
— tidyverse 2.0.0 —
## √ dplyr
             1.1.4
                        ✓ readr
                                    2.1.5
## / forcats
              1.0.1
                        √ stringr
                                    1.5.2
## ✓ ggplot2 4.0.0
                        ✓ tibble
                                    3. 3. 0
## ✓ lubridate 1.9.4

√ tidyr

                                    1.3.1
## √ purrr
              1.1.0
## —— Conflicts -
---- tidyverse conflicts() ---
## X dplyr::filter() masks stats::filter()
## X dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicts t
o become errors
```

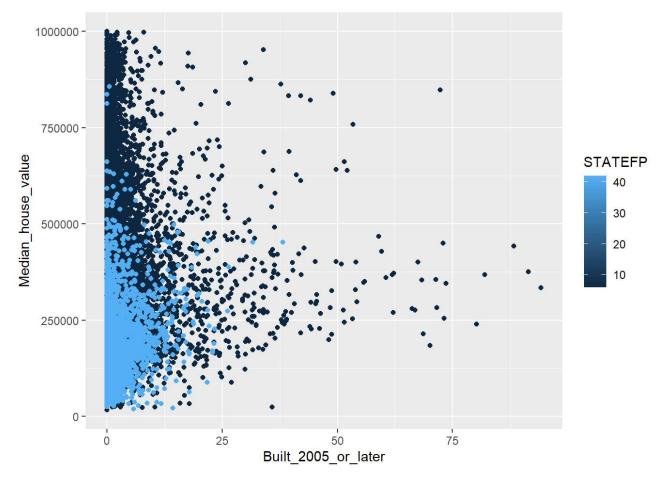
```
# Plot median house prices 与 Built_2005_or_later的关系
ca_pa_omitna |> ggplot()+aes(x=Built_2005_or_later,y=Median_house_value)+geom_point()+ la
bs(x = "Built_2005_or_later", y = "Median_house_value")
```



b.

## #根据所在州对数据进行区分

ca\_pa\_omitna  $\mid$  > ggplot()+aes(x=Built\_2005\_or\_later, y=Median\_house\_value, color=STATEFP)+ge om\_point()+ labs( x = "Built\_2005\_or\_later", y = "Median\_house\_value")



C.

 $ca\_pa\$vacancy\_rate \ \leftarrow \ ca\_pa\$Vacant\_units/ca\_pa\$Total\_units$ 

3. a.

min(ca\_pa\$vacancy\_rate, na.rm=TRUE) #0

## [1] 0

max(ca\_pa\$vacancy\_rate, na. rm=TRUE) #1

## [1] 1

mean(ca\_pa\$vacancy\_rate, na.rm=TRUE) #0.08917878

## [1] 0.08917878

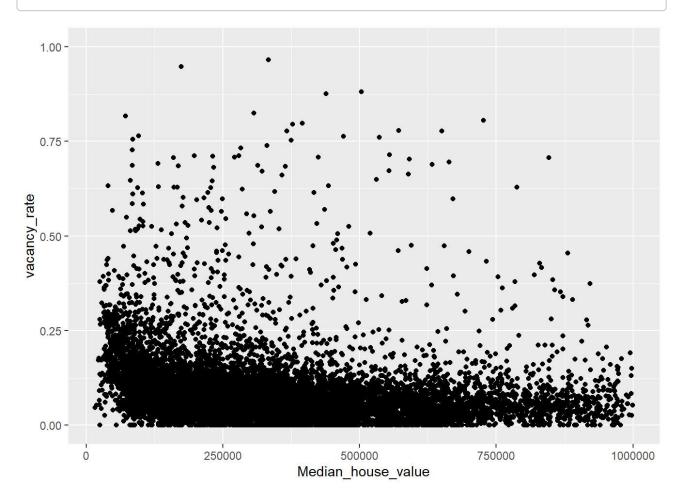
median(ca\_pa\$vacancy\_rate, na.rm=TRUE) #0.06766326

## [1] 0.06766326

b.

ca\_pa |> ggplot()+aes(x=Median\_house\_value, y=vacancy\_rate)+geom\_point()+labs(x="Median\_house\_value", y="vacancy\_rate")

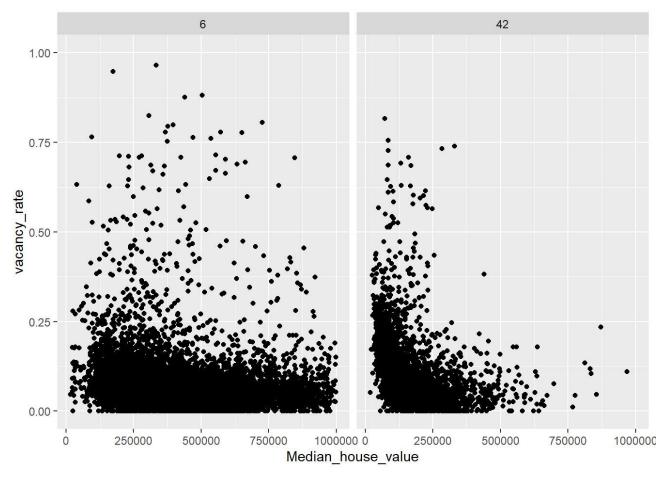
## Warning: Removed 599 rows containing missing values or values outside the scale range ## (`geom\_point()`).



C.

ca\_pa |> ggplot()+aes(x=Median\_house\_value, y=vacancy\_rate)+geom\_point()+labs(x="Median\_house\_value", y="vacancy\_rate")+facet\_wrap( $^{\sim}$ STATEFP)

## Warning: Removed 599 rows containing missing values or values outside the scale range ## (`geom\_point()`).



#宾夕法尼亚州中位房屋价值越高的区域空置率越低,而加利福尼亚州的房屋空置率分布比较均匀

#### 4. a

```
#这段代码要实现将Alameda County, California的median_house_value列成一个vector, 并求中间值acca <- c() #先创建一个空vector, 用于存储行号
for (tract in 1:nrow(ca_pa)) {#从第一行遍历到末尾
if (ca_pa$STATEFP[tract] == 6) {
  if (ca_pa$COUNTYFP[tract] == 1) {
    acca <- c(acca, tract) #将STATEFP==6且COUNTYFP==1的行号添加到acca中
  }
  }
}
accamhv <- c() #创建空vector,
for (tract in acca) { #遍历acca
    accamhv <- c(accamhv, ca_pa[tract,10]) #将每一行的第十列即median_house_value添加到accamhv中
  }
median(accamhv)#求Alameda County, California的median_house_value的中间值
```

## [1] NA

b.

```
median(ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1, 10], na.rm = TRUE)
```

## [1] 473500

C.

```
#Alameda County average percentages of housing built since 2005 is 2.932778 mean(ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1, "Built_2005_or_later"], na.rm = TRUE)
```

```
## [1] 2.932778
```

```
# Santa Clara average percentages of housing built since 2005 is 3.160215
mean(ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85, "Built_2005_or_later"], na.rm = TRUE)
```

```
## [1] 3.160215
```

```
#Allegheny County average percentages of housing built since 2005 is 1.883375 mean(ca_pa[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3, "Built_2005_or_later"], na.rm = TRUE)
```

```
## [1] 1.883375
```

#### d (i) the whole data

```
# the whole data -0.02052684 cor(ca_pa$Median_house_value,ca_pa$Built_2005_or_later,use = "complete.obs")
```

```
## [1] -0.02052684
```

#### (ii). all of California

```
# 0.2339447

cor(ca_pa[ca_pa$STATEFP == 42, c("Median_house_value", "Built_2005_or_later")], use = "comp lete.obs")
```

```
## Median_house_value Built_2005_or_later

## Median_house_value 1.0000000 0.2339447

## Built_2005_or_later 0.2339447 1.0000000
```

#### (iii). all of Pennsylvania

```
# -0.1160322
cor(ca_pa[ca_pa$STATEFP == 6, c("Median_house_value", "Built_2005_or_later")], use = "comple
te.obs")
```

```
## Median_house_value Built_2005_or_later
## Median_house_value 1.0000000 -0.1160322
## Built_2005_or_later -0.1160322 1.00000000
```

## (iv). Alameda County

```
# 0.01432789
cor(ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1,c("Median_house_value","Built_2005_or_
later")], use = "complete.obs")
```

#### (v). Santa Clara County

```
# -0.1726203

cor(ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85,c("Median_house_value","Built_2005_or
_later")],use = "complete.obs")
```

```
## Median_house_value Built_2005_or_later
## Median_house_value 1.0000000 -0.1726203
## Built_2005_or_later -0.1726203 1.0000000
```

## (vi). Allegheny County

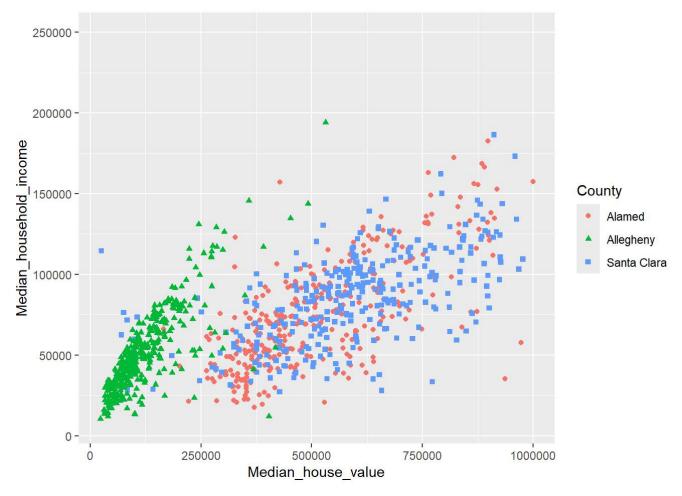
```
# 0.1868602
cor(ca_pa[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3,c("Median_house_value","Built_2005_or
_later")], use = "complete.obs")
```

```
## Median_house_value Built_2005_or_later
## Median_house_value 1.0000000 0.1868602
## Built_2005_or_later 0.1868602 1.0000000
```

e.

```
#先将三个城市的信息提取出来
ca_pa_sub <- ca_pa[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1 |ca_pa$STATEFP == 6 & ca_pa$COUNTYF
P == 85 | ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3,c("STATEFP","COUNTYFP","Median_house_valu
e","Median_household_income")]
#添加城市列名
ca_pa_sub$County <- ifelse(ca_pa_sub$STATEFP == 6 & ca_pa_sub$COUNTYFP == 1, "Alamed", ifelse(c
a_pa_sub$STATEFP == 6 & ca_pa_sub$COUNTYFP == 85, "Santa Clara","Allegheny"))
#画出三个城市的图,用County区分
ca_pa_sub |> ggplot()+aes(x=Median_house_value,y=Median_household_income,color=County,shape = C
ounty)+geom_point() + labs(x="Median_house_value",y="Median_household_income",color="County",shape = "County")
```

```
\#\# Warning: Removed 91 rows containing missing values or values outside the scale range \#\# (`geom_point()`).
```



#### MB.Ch1.11.

#将"female"重复91次, "male"重复92次连接成一个字符串向量, 将字符串向量转化为一个因子gender <- factor(c(rep("female", 91), rep("male", 92)))
table(gender)#统计gender因子中的每个水平的频次

```
## gender
## female male
## 91 92
```

```
gender <- factor(gender, levels=c("male", "female")) #将因子gender的水平顺序从默认顺序改为指定的c("male", "female")顺序

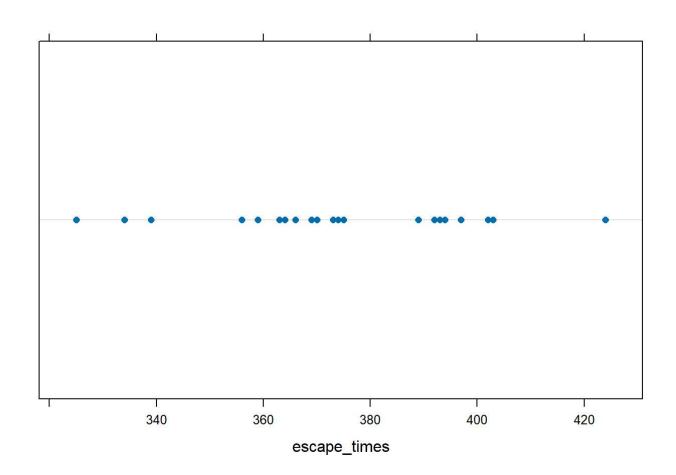
gender <- factor(gender, levels=c("Male", "female"))
#"male"无法匹配"Male", 所以被转为NA
table(gender, exclude=NULL)
```

```
## gender
## Male female <NA>
## 0 91 92
```

#exclude = NULL: 强制显示所有水平,包括NA的计数,上面92个"male"被转换为了NA,所以显示92个NA

#### MB.Ch1.12.

```
proportion_fun <- function(x, cutoff) {</pre>
   #统计x中大于cutoff的数量
   above_cutoff <- sum(x>cutoff, na.rm=TRUE)
   #统计x的总数
   total \leftarrow sum(!is.na(x))
   return (above_cutoff/total)#返回超过cutoff的比例
(a).
 test1 <- 1:100
 proportion_fun(test1, 25) #0.75
 ## [1] 0.75
 proportion_fun(test1,76) #0.24
 ## [1] 0.24
(b).
 install.packages("Devore7")
 ## package 'Devore7' successfully unpacked and MD5 sums checked
 ##
 ## The downloaded binary packages are in
 ## C:\Users\78471\AppData\Local\Temp\RtmpKa7YJP\downloaded_packages
 library (Devore7)
 ## Loading required package: MASS
 ## Attaching package: 'MASS'
 ## The following object is masked from 'package:dplyr':
 ##
 ##
        select
 ## Loading required package: lattice
 data(ex01.36)
 escape times <- ex01.36
 dotplot(~escape_times)
```



proportion\_fun(escape\_times, 420) #0.03846154

## [1] 0.03846154

#### MB.Ch1.18.

```
install.packages("MASS")
```

## Warning: package 'MASS' is in use and will not be installed

```
library(MASS)
#将Rabbit根据Treatment、Dose、Animal的优先级排序
rabbit_sorted <- Rabbit[order(Rabbit$Treatment, Rabbit$Dose, Rabbit$Animal),]
# 使用unstack将Animal展开为列
rabbit_unstack1 <- unstack(rabbit_sorted, BPchange ~ Animal)
# 提取唯一的Treatment和Dose组合
treatment_dose_combos <- unique(rabbit_sorted[, c("Treatment", "Dose")])
# 将treatment_dose_combos 和 rabbit_unstack1 组合成为最终结果
rabbit_unstack <- cbind(treatment_dose_combos, rabbit_unstack1)
print(rabbit_unstack)
```

```
##
     Treatment
                 Dose
                          R1
                                R2
                                      R3
                                            R4
                                                 R5
## 1
       Control
                 6.25
                        0.50
                              1.00
                                    0.75
                                          1.25
                                                1.5
## 2
       Control 12.50
                       4.50
                              1.25
                                    3.00
                                          1.50
                                                1.5
## 3
       Control 25.00 10.00
                             4.00
                                    3.00
                                          6.00
                                                5.0
## 4
       Control 50.00 26.00 12.00 14.00 19.00 16.0
## 5
       Control 100.00 37.00 27.00 22.00 33.00 20.0
## 6
       Control 200.00 32.00 29.00 24.00 33.00 18.0
## 31
            MDL
                  6.25
                        1.25
                              1.40
                                    0.75
                                          2.60
## 32
            MDL
                12.50
                        0.75
                              1.70
                                    2.30
                                          1.20
                                                2.5
## 33
            MDL
                25.00
                        4.00
                              1.00
                                    3.00
                                          2.00
                                                1.5
## 34
                50.00
                        9.00
                              2.00
                                    5.00
            MDL
                                          3.00
## 35
            MDL 100.00 25.00 15.00 26.00 11.00 9.0
## 36
            MDL 200.00 37.00 28.00 25.00 22.00 19.0
```