# Homework 2 - Report

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The data set calif\_penn\_2011.csv contains information about the housing stock of California and Pennsylvania, as of 2011. Information as aggregated into "Census tracts", geographic regions of a few thousand people which are supposed to be fairly homogeneous economically and socially.

- 1. Loading and cleaning
- a. Load the data into a dataframe called ca\_pa.

```
ca_pa <- read.csv("../data/calif_penn_2011.csv", header = TRUE, sep = ",")</pre>
```

b. How many rows and columns does the dataframe have?

```
rows <- dim(ca_pa)[1]
columns <- dim(ca_pa)[2]</pre>
```

c. Run this command, and explain, in words, what this does:

Ans: this command can figure out the number of missing values in every column.

```
colSums(apply(ca_pa,c(1,2),is.na))
```

##	Х	GEO.id2
##	0	0
##	STATEFP	COUNTYFP
##	0	0
##	TRACTCE	POPULATION
##	0	0
##	LATITUDE	LONGITUDE
##	0	0
##	GEO.display.label	Median house value
##	0	599
##	Total units	Vacant units
##	_ 0	_ 0
##	Median rooms	Mean household size owners
##	157	215
##	Mean_household_size_renters	Built_2005_or_later
##	152	98
##	Built_2000_to_2004	Built_1990s
##	98	98
##	Built 1980s	Built_1970s
##	98	98
##	Built 1960s	Built_1950s
##	98	98
##	Built_1940s	Built_1939_or_earlier
##	98	98
##	Bedrooms 0	Bedrooms_1
	Dodioomb_o	Dodfoomb_1

```
##
                              98
                                                             98
##
                     Bedrooms 2
                                                    Bedrooms 3
##
                              98
                                                             98
##
                     Bedrooms_4
                                           Bedrooms_5_or_more
##
##
                          Owners
                                                       Renters
##
                             100
##
       Median_household_income
                                        Mean_household_income
##
```

d. The function na.omit() takes a dataframe and returns a new dataframe, omitting any row containing an NA value. Use it to purge the data set of rows with incomplete data.

```
ca_pa <- na.omit(ca_pa)</pre>
```

e. How many rows did this eliminate?

```
rows - dim(ca_pa)[1]
```

## [1] 670

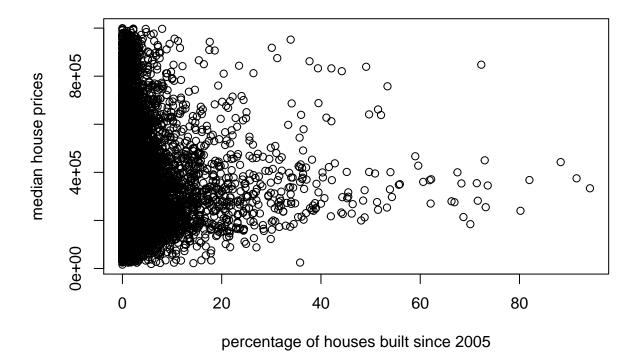
f. Are your answers in (c) and (e) compatible? Explain.

**Ans:** They are compatible. The command in (c) check the number of missing values in every column, and the command in (e) check the number of rows with incomplete data. We can infer that after purging, the number of missing values in every column will be zero. And we can use the command below to check out the truth.

```
sum(colSums(apply(ca_pa,c(1,2),is.na)))
```

## [1] 0

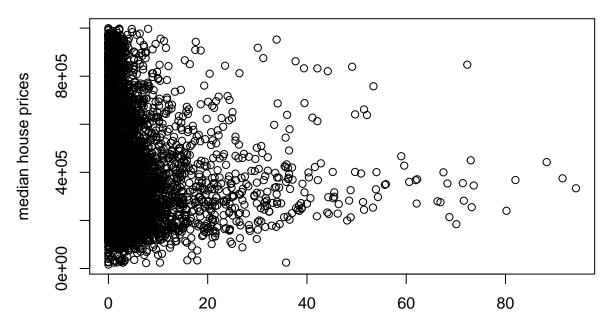
- 2. This Very New House
- a. The variable Built\_2005\_or\_later indicates the percentage of houses in each Census tract built since 2005. Plot median house prices against this variable.



b. Make a new plot, or pair of plots, which breaks this out by state. Note that the state is recorded in the STATEFP variable, with California being state 6 and Pennsylvania state 42.

```
plot(ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6],
    ca_pa$Median_house_value[ca_pa$STATEFP == 6],
    xlab = "percentage of houses built since 2005",
    ylab = "median house prices",
    main = "Houses in California")
```

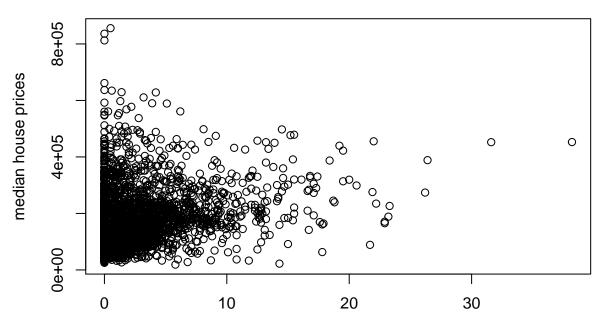
# **Houses in California**



percentage of houses built since 2005

```
plot(ca_pa$Built_2005_or_later[ca_pa$STATEFP == 42],
     ca_pa$Median_house_value[ca_pa$STATEFP == 42],
     xlab = "percentage of houses built since 2005",
     ylab = "median house prices",
     main = "Houses in Pennsylvania")
```

### **Houses in Pennsylvania**



percentage of houses built since 2005

#### 3. Nobody Home

The vacancy rate is the fraction of housing units which are not occupied. The dataframe contains columns giving the total number of housing units for each Census tract, and the number of vacant housing units.

a. Add a new column to the dataframe which contains the vacancy rate. What are the minimum, maximum, mean, and median vacancy rates?

```
Vacancy_rate <- ca_pa$Vacant_units / ca_pa$Total_units
ca_pa <- data.frame(ca_pa, Vacancy_rate)
max(Vacancy_rate)

## [1] 0.965311
min(Vacancy_rate)

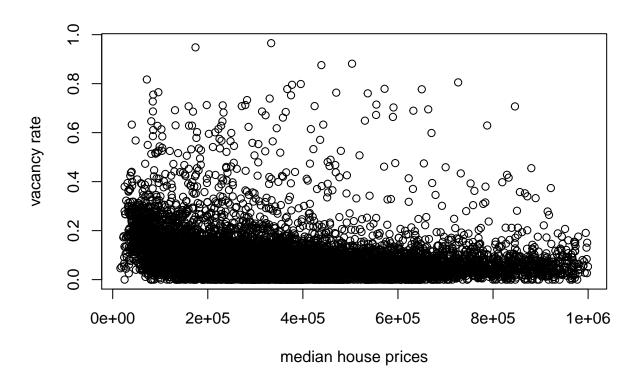
## [1] 0
mean(Vacancy_rate)

## [1] 0.08888789
median(Vacancy_rate)

## [1] 0.06767283
b. Plot the vacancy rate against median house value.</pre>
```

plot(ca\_pa\$Median\_house\_value, ca\_pa\$Vacancy\_rate,

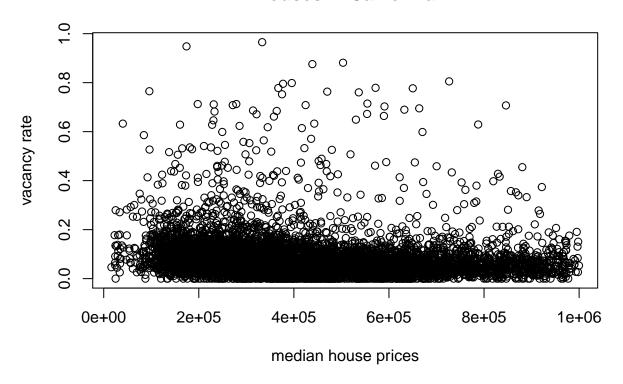
xlab = "median house prices", ylab = "vacancy rate")



c. Plot vacancy rate against median house value separately for California and for Pennsylvania. Is there a difference?

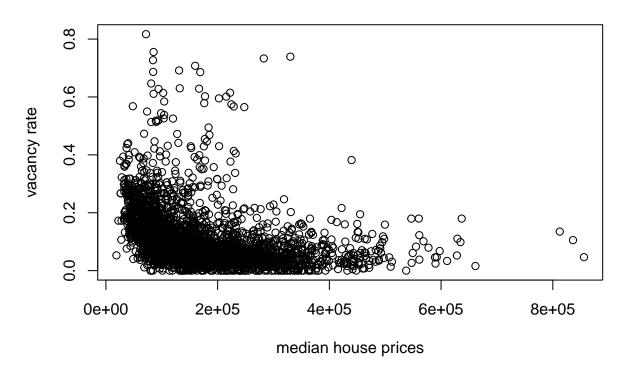
```
plot(ca_pa$Median_house_value[ca_pa$STATEFP == 6],
    ca_pa$Vacancy_rate[ca_pa$STATEFP == 6],
    xlab = "median house prices", ylab = "vacancy rate",
    main = "Houses in California")
```

# **Houses in California**



```
plot(ca_pa$Median_house_value[ca_pa$STATEFP == 42],
    ca_pa$Vacancy_rate[ca_pa$STATEFP == 42],
    xlab = "median house prices", ylab = "vacancy rate",
    main = "Houses in Pennsylvania")
```

### Houses in Pennsylvania



- 4. The column COUNTYFP contains a numerical code for counties within each state. We are interested in Alameda County (county 1 in California), Santa Clara (county 85 in California), and Allegheny County (county 3 in Pennsylvania).
- a. Explain what the block of code at the end of this question is supposed to accomplish, and how it does it.

Ans: This block of code is supposed to pick up the tracts in Alameda County and compute the median of the median house values of those tracts. The code firstly traverse all the tracts in ca\_pa, and if a tract matches condition, it will be stored in a new vector acca. Then the code traverse again to store the median house values of the tracts in acca into accamhv. Finally the code call median function to compute the median of accamhv.

b. Give a single line of R which gives the same final answer as the block of code. Note: there are at least two ways to do this; you just have to find one.

```
median(ca_pa$Median_house_value[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1])
```

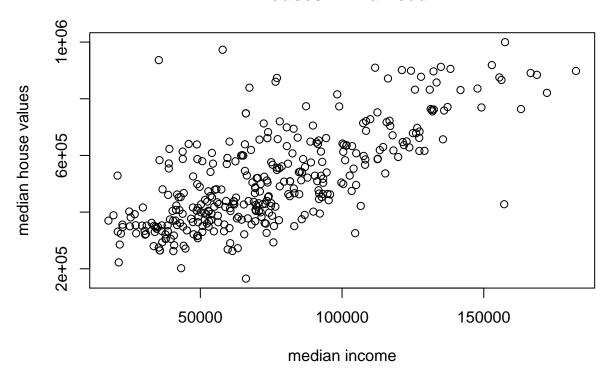
#### ## [1] 474050

c. For Alameda, Santa Clara and Allegheny Counties, what were the average percentages of housing built since 2005?

```
# Alameda
mean(ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1])
## [1] 2.820468
# Santa Clara
mean(ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85])
```

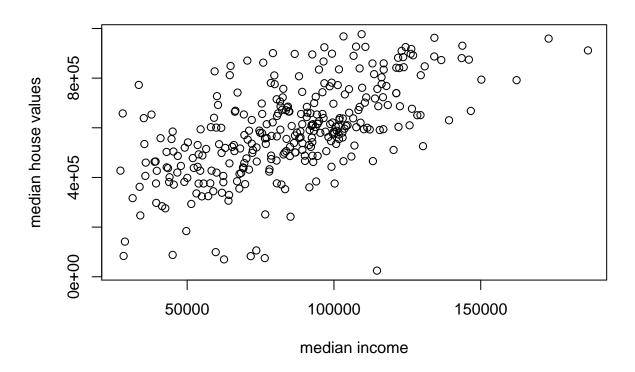
```
## [1] 3.200319
# Allegheny
mean(ca_pa$Built_2005_or_later[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3])
## [1] 1.474219
  d. The cor function calculates the correlation coefficient between two variables. What is the correlation
     between median house value and the percent of housing built since 2005 in (i) the whole data, (ii) all of
     California, (iii) all of Pennsylvania, (iv) Alameda County, (v) Santa Clara County and (vi) Allegheny
     County?
# (i) the whole data
cor(ca pa$Median house value, ca pa$Built 2005 or later)
## [1] -0.01893186
# (ii) all of California
cor(ca_pa$Median_house_value[ca_pa$STATEFP == 6],
    ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6])
## [1] -0.1153604
# (iii) all of Pennsylvania
cor(ca_pa$Median_house_value[ca_pa$STATEFP == 42],
    ca_pa$Built_2005_or_later[ca_pa$STATEFP == 42])
## [1] 0.2681654
# (iv) Alameda
cor(ca pa$Median house value[ca pa$STATEFP == 6 & ca pa$COUNTYFP == 1],
    ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1])
## [1] 0.01303543
# (v) Santa Clara
cor(ca_pa$Median_house_value[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85],
    ca_pa$Built_2005_or_later[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85])
## [1] -0.1726203
# (vi) Allegheny
cor(ca_pa$Median_house_value[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3],
    ca_pa$Built_2005_or_later[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3])
## [1] 0.1939652
  e. Make three plots, showing median house values against median income, for Alameda, Santa Clara,
     and Allegheny Counties. (If you can fit the information into one plot, clearly distinguishing the three
     counties, that's OK too.)
plot(ca_pa$Median_household_income[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1],
     ca_pa$Median_house_value[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 1],
     xlab = "median income", ylab = "median house values",
     main = "Houses in Alameda")
```

# **Houses in Alameda**



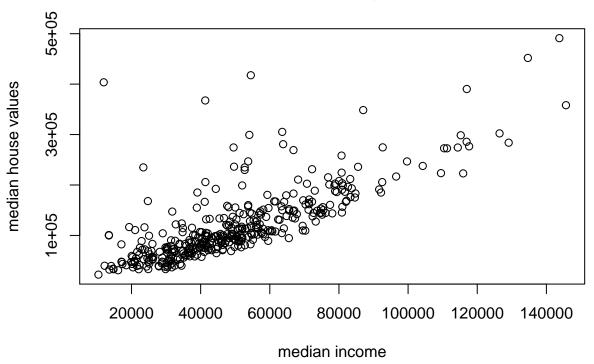
```
# Santa Clara
plot(ca_pa$Median_household_income[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85],
    ca_pa$Median_house_value[ca_pa$STATEFP == 6 & ca_pa$COUNTYFP == 85],
    xlab = "median income", ylab = "median house values",
    main = "Houses in Santa Clara")
```

### **Houses in Santa Clara**



```
# Allegheny
plot(ca_pa$Median_household_income[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3],
    ca_pa$Median_house_value[ca_pa$STATEFP == 42 & ca_pa$COUNTYFP == 3],
    xlab = "median income", ylab = "median house values",
    main = "Houses in Allegheny")
```

# **Houses in Allegheny**



```
acca <- c()
for (tract in 1:nrow(ca_pa)) {
   if (ca_pa$STATEFP[tract] == 6) {
      if (ca_pa$COUNTYFP[tract] == 1) {
        acca <- c(acca, tract)
      }
   }
}
accamhv <- c()
for (tract in acca) {
   accamhv <- c(accamhv, ca_pa[tract,10])
}
median(accamhv)</pre>
```