

Name: Niranjana Poudel

Homework Assignment 04 (Before class starts)

70 Points — Due before week 3

General Instructions

For this fourth homework assignment, you have to create your own RMarkdown (.Rmd) file, based on files from class and from Homework 1, copy the question numbers and the answer options into your .Rmd file, and knit that file into a pdf file. **Alternatively** (and much easier!!!), use this .Rnw file as a template, just fill in the answers into the provided spaces, and knit into a pdf file.

Only the final resulting pdf file (from .Rmd or .Rnw) has to be submitted via Canvas. As previously stated, I would like to encourage potential and current MS and PhD students to work with .Rnw and L^AT_EX instead of .Rmd.

You need to learn how to write R code that is easily readable for others. There exists *Google's R Style Guide* that summarizes rules for good R style. These rules are accessible at <https://google.github.io/styleguide/Rguide.xml>. In particular, make sure that you always have a space after a comma and that you consistently use the same type of assignment operator, ideally <-. Look at the examples on this web page and follow the style whenever you write your own R code from now on.

Do not forget to replace my name and include your name instead! We will print the homeworks, so a homework with no name/my name on it can't be graded!

In all question parts, show your R code and the results!

(i) (20 Points) **Family Data Revisited:**

In the following exercises, try to write your code to be as general as possible so that it would still work if the family had 27 members in it or if the variables were in a different order in the data frame.

Show your R code and the final results produced from within R for all question parts!

- (a) (3 Points) Copy the family data set for this homework from Canvas into your local folder for this homework. Then load the `hw04_familyDF.rda` data set into R. Show the objects that have been loaded. Is the first object that is listed a data frame? Search for help if you don't recall how to check whether something is a data frame.

Answer:

```
> load(file = "hw04_familyDF.rda")
> print(load(file = "hw04_familyDF.rda"))

[1] "family"

> head(family) # To show object is loaded
```

	firstName	gender	age	height	weight	bmi	overWt
1	Tom	m	77	70	175	25.16239	TRUE
2	May	f	33	64	125	21.50106	FALSE
3	Joe	m	79	73	185	24.45884	FALSE
4	Bob	m	47	67	156	24.48414	FALSE
5	Sue	f	27	64	105	18.06089	FALSE
6	Liz	f	33	68	190	28.94981	TRUE

```
> options(width = 80) # For pdf display
> is.data.frame(family)

[1] TRUE
```

- (b) (4 Points) The NHANES survey used different cut-off values for men and women when classifying them as overweight. Suppose that a man is classified as obese if his bmi exceeds 26 and a woman is classified as obese if her bmi exceeds 25. Write a logical expression to create a logical vector, called `OW.NHANES`, that is `TRUE` if a member of the family is obese and `FALSE` otherwise. Display its content.

Answer:

```
> OW.NHANES <- as.logical((family$gender == 'm' & family$bmi > 26)
+                           | (family$gender == 'f' & family$bmi > 25))
> OW.NHANES

[1] FALSE FALSE FALSE FALSE FALSE  TRUE  TRUE FALSE  TRUE  TRUE  TRUE FALSE
[13] FALSE FALSE
```

- (c) (4 Points) Here is an alternative way to create the same vector that introduces some useful functions and ideas. We first create a numeric vector called `OW.limit` that is 26 for each male in the family and 25 for each female in the family. To do this, we create a vector of length 2, called `OW.val`, where the first element is 26 and second element is 25. Then we create the `OW.limit` vector by subsetting `OW.val` by position, where the positions are the numeric values in the gender variable (i.e., use `as.numeric` to coerce the factor vector to a numeric vector). Notice that we can “subset” a vector of length 2 by a much longer vector:

```
> OW.val <- 26:25
> OW.limit <- OW.val[as.numeric(family$gender)]
> OW.limit
```

Finally, use `OW.limit` and the `bmi` vector in `family` to create the desired logical vector, and call it `OW.NHANES2`. Display its content. Compare with your results from part (b) via the `any` function. Did you get the intended result? If not, check your R code again!

Answer:

```
> OW.val <- 26:25
> OW.limit <- OW.val[as.numeric(family$gender)]
> OW.limit

[1] 26 25 26 26 25 25 26 25 26 26 25 26 26 25

> OW.NHANES2 <- as.logical(family$bmi > OW.limit)
> OW.NHANES2

[1] FALSE FALSE FALSE FALSE FALSE  TRUE  TRUE FALSE  TRUE  TRUE  TRUE FALSE
[13] FALSE FALSE
```

- (d) (4 Points) Use the vector `OW.limit` and each person’s height to find the weight that they would have if their `bmi` was right at the limit (26 for men and 25 for women). Call this weight `OW.weight` and display its content. To do this, start with the formula

$$\text{bmi} = (\text{weight} / 2.2) / (2.54 / 100 * \text{height})^2$$

and re-express it in terms of weight (i.e., `weight = ...`).

Answer:

```
> OW.weight <- OW.limit * (2.54 / 100 * family$height) ^ 2 * 2.2
> OW.weight

[1] 180.8254 145.3416 196.6569 165.6582 145.3416 164.0771 170.6402 149.9191
[9] 170.6402 186.0288 159.2868 160.7501 160.7501 136.3997
```

- (e) (5 Points) Create the following plot of actual weight (on the vertical axis) against the weight at which they would be overweight (on the horizontal axis). If you get an error when you run this code, check whether you are using the correct variable names in your code earlier on.

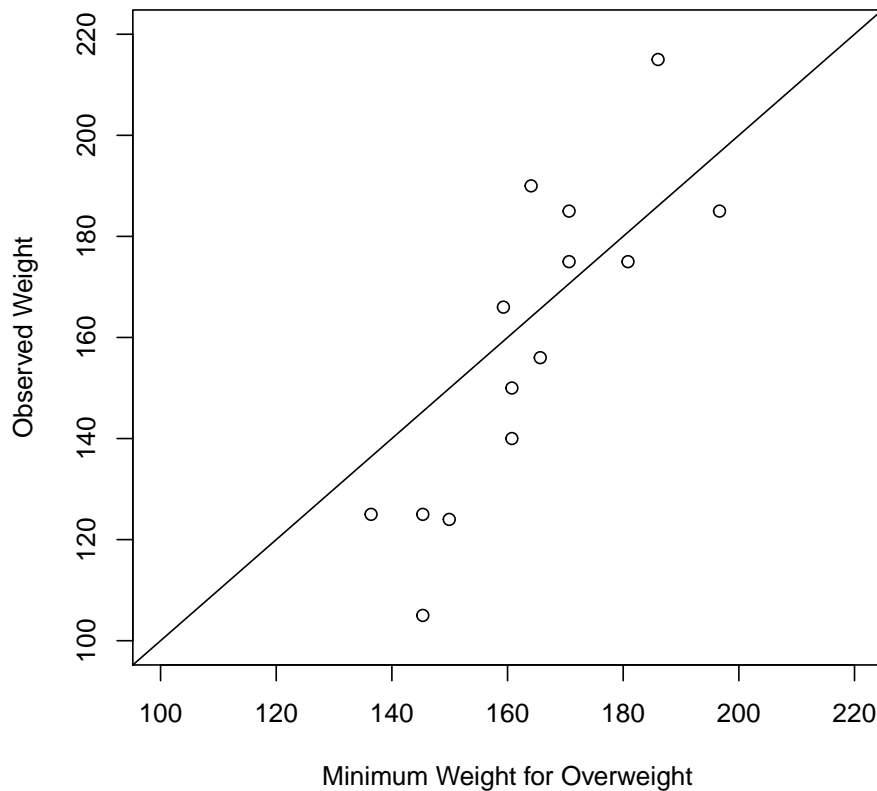
```
> plot(OW.weight, family$weight,  
+      xlab = "Minimum Weight for Overweight",  
+      xlim = c(100, 220), # !!!  
+      ylab = "Observed Weight",  
+      ylim = c(100, 220)) # !!!  
> abline(a = 0, b = 1)
```

`abline` adds a straight line (here with y-intercept $a = 0$ and slope $b = 1$) to the plot. Note that this is not the regression line! Thus, points that fall exactly on the line belong to individuals where the observed weight exactly qualifies to be overweight. Points above the line represent individuals who are overweight, and points below the line represent individuals who are not overweight.

We can easily count in the plot how many points are above the line and how many points are below the line, but we want that R does this counting for us! So, write two R expressions that do this counting for us and display their results.

Answer:

```
> plot(OW.weight, family$weight,
+      xlab = "Minimum Weight for Overweight",
+      xlim = c(100, 220), # !!!
+      ylab = "Observed Weight",
+      ylim = c(100, 220)) # !!!
> abline(a = 0, b = 1)
```



```
> # Number of points above the line
> sum(OW.weight < family$weight)
[1] 5
> # Number of points below the line
> sum(OW.weight > family$weight)
[1] 9
```

(ii) (34 Points) **San Francisco Housing Data:**

In this question, you have to work with actual housing data from the San Francisco area.

Show your R code and the final results produced from within R for all question parts!

- (a) (4 Points) Copy the San Francisco housing data set (`hw04_SFhousing.rda`) for this homework from Canvas into your local folder for this homework. Then load this data set into R. Show the objects that have been loaded. Are `cities` and `housing` both data frames? Let R answer this question! Search for help if you don't recall how to check whether something is a data frame.

Answer:

```
> load(file = "hw04_SFhousing.rda")
> print(load(file = "hw04_SFhousing.rda"))

[1] "cities" "housing"

> head(cities) # To show objects are loaded
```

	longitude	latitude	county	medianPrice	medianSize
Alameda	-122.2485	37.75993	Alameda County	580000	1489.0
Alamo	-122.0205	37.85522	Contra Costa County	1250000	2723.5
Albany	-122.2940	37.89107	Alameda County	520250	1170.0
Almaden	NA	NA	Santa Clara County	835000	2139.0
American Canyon	-122.2580	38.16664	Napa County	419000	1344.0
Angwin	-122.4499	38.57451	Napa County	662000	1822.0

```

numHouses medianBR
Alameda      2339      3
Alamo         760      4
Albany        640      2
Almaden       1705      4
American Canyon 463      3
Angwin         79      3

> head(family) # To show objects are loaded
```

	firstName	gender	age	height	weight	bmi	overWt
1	Tom	m	77	70	175	25.16239	TRUE
2	May	f	33	64	125	21.50106	FALSE
3	Joe	m	79	73	185	24.45884	FALSE
4	Bob	m	47	67	156	24.48414	FALSE
5	Sue	f	27	64	105	18.06089	FALSE
6	Liz	f	33	68	190	28.94981	TRUE

```
> is.data.frame(cities)

[1] TRUE

> is.data.frame(housing)

[1] TRUE
```

- (b) (2 Points) What are the names of the vectors in `housing`?

Answer:

```
> names(housing)

[1] "county" "city" "zip" "street" "price" "br" "lsqft"
[8] "bsqft" "year" "date" "long" "lat" "quality" "match"
[15] "wk"
```

(c) (2 Points) How many observations are in housing?

Answer:

```
> nrow(housing)
```

```
[1] 281506
```

- (d) (6 Points) Explore the housing data using the summary function. Describe in words at least three problems that you see with the data.

Answer:

```
> summary(housing)
```

```

      county      city      zip
Santa Clara County :70424  Oakland      : 14730  94565 : 4595
Alameda County     :60410  Santa Rosa  : 9917  94509 : 4302
Contra Costa County:59381  Fremont    : 9414  95123 : 4023
Solano County       :23404  San Francisco: 8137  95687 : 3652
San Mateo County    :22558  Evergreen   : 7947  94533 : 3472
Sonoma County       :21676  Antioch     : 7726  (Other):261457
(Other)             :23653  (Other)     :223635  NA's   : 5

      street      price      br      lsqft
Length:281506   Min.    : 22000   Min.    :1.000   Min.    : 19
Class :character 1st Qu.: 400000   1st Qu.:2.000   1st Qu.: 4000
Mode  :character Median : 530000   Median :3.000   Median : 5760
      Mean : 602000   Mean :3.024   Mean : 65939
      3rd Qu.: 700000   3rd Qu.:4.000   3rd Qu.: 7701
      Max. :20000000   Max. :8.000   Max. :418611600
                        NA's :21687

      bsqft      year      date
Min.    : 122   Min.    : 0   Min.    :2003-04-27 01:00:00
1st Qu.: 1121   1st Qu.:1954   1st Qu.:2004-02-08 01:00:00
Median : 1430   Median :1971   Median :2004-10-24 01:00:00
Mean : 1624   Mean :1966   Mean :2004-11-01 17:06:12
3rd Qu.: 1882   3rd Qu.:1985   3rd Qu.:2005-07-24 01:00:00
Max. :1868120   Max. :3894   Max. :2006-06-04 01:00:00
NA's :426   NA's :9202

      long      lat
Min.    :-123.6   Min.    :36.98
1st Qu.: -122.3   1st Qu.:37.50
Median : -122.1   Median :37.77
Mean : -122.1   Mean :37.78
3rd Qu.: -121.9   3rd Qu.:38.00
Max.    :-121.5   Max.    :38.85
NA's :23316   NA's :23316

      quality      match
QUALITY_ADDRESS_RANGE_INTERPOLATION :170719   Exact      :197044
gpsvisualizer      : 31084   Relaxed     : 30570
QUALITY_CITY_CENTROID      : 20473   Relaxed; Soundex: 23338
QUALITY_EXACT_PARCEL_CENTROID      : 17208   Soundex     : 2573
QUALITY_ZIP_CODE_TABULATION_AREA_CENTROID: 14980   1           : 2244
(Other)             : 3726   (Other)     : 2421
NA's                : 23316   NA's        : 23316

      wk
Min.    :2003-04-21
1st Qu.:2004-02-01
Median :2004-10-18
Mean :2004-10-26
3rd Qu.:2005-07-18
Max.    :2006-05-29

```


Problems:

- i. Problem 1. High number of NA's present in some columns.
 - ii. Problem 2. The year column has years greater than 2020.
 - iii. Problem 3. The column lsqft seems to have a very large area (Max area)
- (e) (4 Points) We will work with houses in Albany, Berkeley, Piedmont, and Emeryville only. Subset the data frame so that we have only houses in these cities, and keep only the variables city, zip, price, br, bsqft, and year. Call this new data frame BerkArea. This data frame should have 4059 observations and 6 variables (check it!).

Answer:

```
> BerkArea <- subset(housing, housing$city %in%
+                   c('Albany', 'Berkeley', 'Piedmont', 'Emeryville'),
+                   select = c('city', 'zip', 'price', 'br', 'bsqft',
+                               'year'))
> dim(BerkArea)
[1] 4059    6
```

- (f) (4 Points) We are interested in studying the relationship between price and size of house, but first we will further subset the data frame to remove the unusually large values. Use the quantile function to determine the 99th percentile of price and bsqft and eliminate all of those houses that are above either of these 99th percentiles. Call this new data frame BerkArea, as well. It should have 3999 observations (check it!). Write your code so that it is very general and does not depend on the actual numeric value for these quantiles.

Answer:

```
> BerkArea <- subset(BerkArea, (
+   (BerkArea$price <= quantile(BerkArea$price, 0.99))
+   & (BerkArea$bsqft <= quantile(BerkArea$bsqft, 0.99,
+   na.rm = T) | is.na(BerkArea$bsqft))))
> nrow(BerkArea)
[1] 3999
```

- (g) (2 Points) Create a new vector that is called `pricepsqft` by dividing the sale price by the square footage of the house. Add this new variable to the `BerkArea` housing data frame.

Answer:

```
> BerkArea$pricepsqft <- BerkArea$price / BerkArea$bsqft
```

- (h) (4 Points) Create a vector called `br5` that contains the number of bedrooms in the house, except when this number is greater than 5, it is set to 5. That is, if a house has 5 or more bedrooms then `br5` will be 5. Otherwise it will be the number of bedrooms in the house. Note that there is no need for any “if”-statements or loops to create this vector — just basic R expressions discussed so far will be sufficient! Recall how `TRUE` and `FALSE` are represented numerically or how to reassign a different value to a subset!

Answer:

```
> br5 <- BerkArea$br # Makes it easy for to code as below
> br5[br5 >= 5] <- 5 # Felt more easy
```

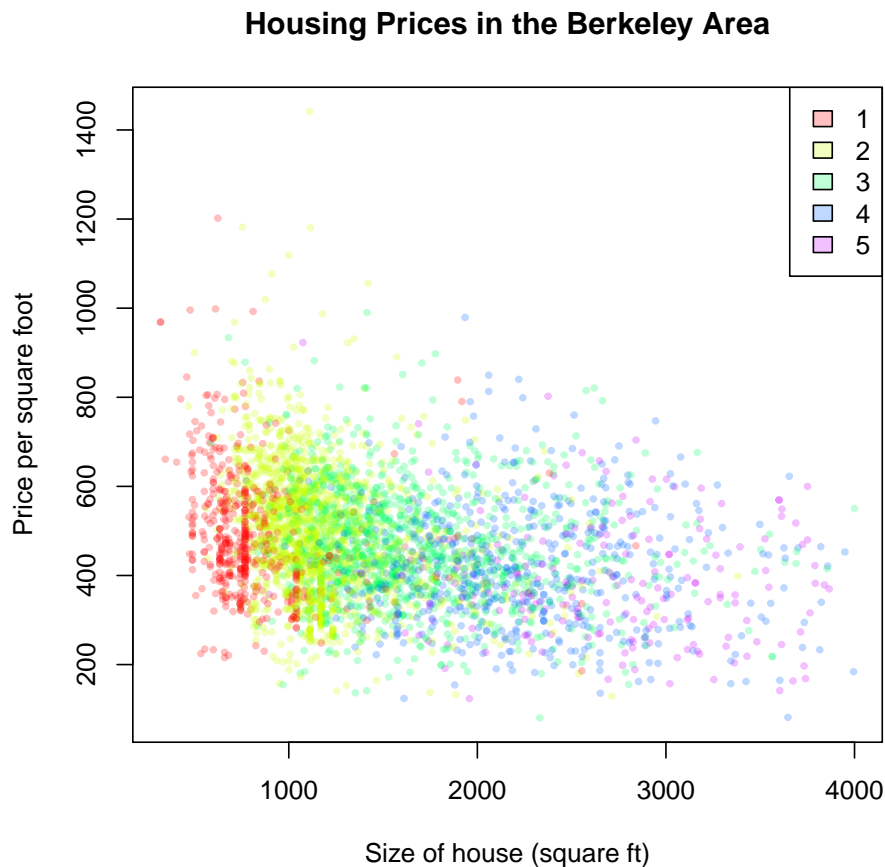
- (i) (6 Points) Recreate the following plot on your side. Then answer the question below. If you get an error when you run this code, check whether you are using the correct variable names in your code earlier on.

```
> rCols <- rainbow(5, alpha = 0.25)
> brCols <- rCols[br5]
> plot(pricepsqft ~ bsqft, data = BerkArea,
+      main = "Housing Prices in the Berkeley Area",
+      xlab = "Size of house (square ft)",
+      ylab = "Price per square foot",
+      col = brCols, pch = 19, cex = 0.5)
> legend(legend = 1:5, fill = rCols, "topright")
```

What interesting feature do you see that you didn't know before making this plot? Numerically quantify (use only 3 decimal digits!) and interpret this feature!

Answer:

```
> rCols <- rainbow(5, alpha = 0.25)
> brCols <- rCols[br5]
> plot(pricepsqft ~ bsqft, data = BerkArea,
+      main = "Housing Prices in the Berkeley Area",
+      xlab = "Size of house (square ft)",
+      ylab = "Price per square foot",
+      col = brCols, pch = 19, cex = 0.5)
> legend(legend = 1:5, fill = rCols, "topright")
```



Before making this plot I was unaware of the housing prices based on the area of the houses (number of bedroom). Obviously as expected with large area there are more number of bedroom, i thought the houses with large area might be more expensive (Price per square feet) but it looks opposite as it seems like there is slight negative correlation (dipping downward trend). But obviously we have not accounted for other factors like the location (which seems to me as a major factor). **Frankly speaking i am not still sure in what context to explain numerically. (may be correlation or just the average price values per bedroom or per area).** I will just calculate the

correlation between pricesqft and bsqft.

```
> # I have used correlation (among many more possibility)
> cor(BerkArea$pricesqft, BerkArea$bsqft, use = "complete.obs")
[1] -0.2929954
```

As expected negative correlation of 0.293 between the price per square feet and area of the house.

(iii) (16 Points) **Survival of Passengers on the Titanic:**

Work with the `Titanic` data set, a 4-dimensional array related to the survival of passengers and crew on board of the Titanic ocean liner. For further details, refer to the help page via `?Titanic`. Technically, the Titanic data set is a table, but we can access it similar to a multi-dimensional array.

Show your R code and the final results produced from within R for all question parts!

- (a) (4 Points) Write an R expression that extracts the numbers of females in all three classes (but not crew) who survived the sinking of the Titanic. Provide data for children and adults. The result should look as follows:

	Age	
Class	Child	Adult
1st	1	140
2nd	13	80
3rd	14	76

Answer:

```
> # The Titanic data is pre-loaded in the packages(datasets)
> Titanic[, "Female", , "Yes"][c("1st", "2nd", "3rd"), ]
```

```
      Age
Class Child Adult
1st      1   140
2nd     13    80
3rd     14    76
```

- (b) (4 Points) Write an R expression that extracts the numbers of male crew members (adults only) who survived or did not survive the sinking of the Titanic. The result should be a vector of length 2.

Answer:

```
> Titanic["Crew", "Male", "Adult", ]
```

```
  No Yes
670 192
```

- (c) (4 Points) Write an R expression that extracts the following matrix from the Titanic data set:

```
      Sex
Class Female Male
Crew      20  192
1st      140   57
2nd       80   14
3rd       76   75
```

Describe what this matrix represents, i.e., which subgroup(s) from the Titanic passengers and crew.

Answer:

```
> row <- c("Crew", "1st", "2nd", "3rd")
> col <- c("Female", "Male")
> Titanic[, , "Adult", "Yes"][row, col]
```

```
      Sex
Class Female Male
Crew      20  192
1st      140   57
2nd       80   14
3rd       76   75
```

This matrix represents the number of adults (male and female) who survived in all class categories.

- (d) (4 Points) Write an R expression that extracts the following vector from the Titanic data set:

```
[1] 35 17 387 89
```

Describe what this vector represents, i.e., which subgroup(s) from the Titanic passengers and crew. Hint: I first extracted a matrix and then transformed this into a vector using `as.vector`.

Answer:

```
> as.vector(Titanic["3rd", , , "No"])
```

```
[1] 35 17 387 89
```

This vector represents the number of people who did not survive (Both child and adult) for male and female separately, the first two values are male and female for child group who did not survive and the later two for adults in similar category.

The end !!!!