

Problem Set 1

Yifan Li

Line to GitHub

The link to my GitHub repository is <https://github.com/FYlee39/Stats-506/tree/main/PS1>.

Problem 1

a

Using `read.table` to read a file written in txt format. For the separation, using `'.'`. Then according to the description file, 'wine.names', there are 14 attributes in the data file with a class number listed in the first column. So adding `col.names` in the code `read.table`. Such that, one can produce a `data.frame` object with appropriate columns names.

```
wines_data <- read.table("wine.data",
                        sep = ",",
                        col.names=c('class_number',
                                    'Alcohol',
                                    'Malic_acid',
                                    'Ash',
                                    'Alcalinity_of_ash',
                                    'Magnesium',
                                    'Total_phenols',
                                    'Flavanoids',
                                    'Nonflavanoid_phenols',
                                    'Proanthocyanins',
                                    'Color_intensity',
                                    'Hue',
                                    'OD280_OD315_of_diluted_wines',
                                    'Proline'))
```

b

First, using `wines_data['class_number']==i` for `i` in `[1, 2, 3]` to create a new `data.frame` that has `True` only if the class numbers match with `i`. After that, using a `sum` function to compute the number of `True`, which is the number of the wine class.

```
num_class_one <- sum(wines_data['class_number'] == 1)
num_class_two <- sum(wines_data['class_number'] == 2)
num_class_three <- sum(wines_data['class_number'] == 3)
```

The results are:

```
num_class_one
```

```
[1] 59
```

```
num_class_two
```

```
[1] 71
```

```
num_class_three
```

```
[1] 48
```

So, the number of wines within each class is correct as reported in the file “wine.names”.

c

1.

The correlation between alcohol content and color intensity can be derived from a function `cor`. The alcohol content has variable name `Alcohol`, the color intensity has variable name `Color_intensity`. So the input of the function will be:

```
cor(wines_data['Alcohol'], wines_data['Color_intensity'])
```

```
           Color_intensity
Alcohol      0.5463642
```

2.

For each class, first the whole data from that class will be extracted, then the correlation between alcohol content and color intensity will be calculated.

For class one:

```
class_one <- wines_data[wines_data['class_number'] == 1, ]
class_one_cor <- cor(class_one['Alcohol'], class_one['Color_intensity'])
class_one_cor
```

```
          Color_intensity
Alcohol      0.4082913
```

For class two:

```
class_two <- wines_data[wines_data['class_number'] == 2, ]
class_two_cor <- cor(class_two['Alcohol'], class_two['Color_intensity'])
class_two_cor
```

```
          Color_intensity
Alcohol      0.2697891
```

For class three:

```
class_three <- wines_data[wines_data['class_number'] == 3, ]
class_three_cor <- cor(class_three['Alcohol'], class_three['Color_intensity'])
class_three_cor
```

```
          Color_intensity
Alcohol      0.3503777
```

Through comparison, one will find that class one has the highest correlation which is 0.4082913, while class two has the lowest correlation which is 0.2697891.

3.

To find the wine with highest color intensity, using `which.max` function, with attributes `wines_data$Color_intensity`. This will yield the index of the wine with highest color intensity. Then using this index to find the wine, after that extract its alcohol content.

```
index <- which.max(wines_data$Color_intensity)
target_wine <- wines_data[index, ]
target_wine$Alcohol
```

```
[1] 14.34
```

Finally extract the alcohol content from the target wine, which is 14.34.

4.

First, find the number of wines that have a higher content of proanthocyanins than ash. Then divide it by the sum of three classes of wines, which will give us the percentage of wines had a higher content of proanthocyanins compare to ash, which is 8.426966%.

```
num <- sum(wines_data$'Proanthocyanins' > wines_data$'Ash')
percentage <- num * 100 / (num_class_one + num_class_two + num_class_three)
percentage
```

```
[1] 8.426966
```

d

```
average_table <- data.frame(id = 1: 4,
                             class_number = c('overall', '1', '2', '3'),

                             Mean_Alcohol = c(mean(wines_data$Alcohol),
                                                mean(class_one$Alcohol),
                                                mean(class_two$Alcohol),
                                                mean(class_three$Alcohol)),

                             Mean_Malic_acid = c(mean(wines_data$Malic_acid),
                                                  mean(class_one$Malic_acid),
                                                  mean(class_two$Malic_acid),
                                                  mean(class_three$Malic_acid)),

                             Mean_Ash = c(mean(wines_data$Ash),
                                             mean(class_one$Ash),
                                             mean(class_two$Ash),
```

```

        mean(class_three$Ash)),

Mean_Alcalinity_of_ash = c(
    mean(wines_data$Alcalinity_of_ash),
    mean(class_one$Alcalinity_of_ash),
    mean(class_two$Alcalinity_of_ash),
    mean(class_three$Alcalinity_of_ash)),

Mean_Magnesium = c(mean(wines_data$Magnesium),
                    mean(class_one$Magnesium),
                    mean(class_two$Magnesium),
                    mean(class_three$Magnesium)),

Mean_Total_phenols = c(mean(wines_data$Total_phenols),
                       mean(class_one$Total_phenols),
                       mean(class_two$Total_phenols),
                       mean(class_three$Total_phenols)),

Mean_Flavanoids = c(mean(wines_data$Flavanoids),
                    mean(class_one$Flavanoids),
                    mean(class_two$Flavanoids),
                    mean(class_three$Flavanoids)),

Mean_Nonflavanoid_phenols = c(
    mean(wines_data$Nonflavanoid_phenols),
    mean(class_one$Nonflavanoid_phenols),
    mean(class_two$Nonflavanoid_phenols),
    mean(class_three$Nonflavanoid_phenols)),

Mean_Proanthocyanins = c(mean(wines_data$Proanthocyanins),
                          mean(class_one$Proanthocyanins),
                          mean(class_two$Proanthocyanins),
                          mean(class_three$Proanthocyanins)),

Mean_Color_intensity = c(mean(wines_data$Color_intensity),
                         mean(class_one$Color_intensity),
                         mean(class_two$Color_intensity),
                         mean(class_three$Color_intensity)),

Mean_Hue = c(mean(wines_data$Hue), mean(class_one$Hue),
             mean(class_two$Hue), mean(class_three$Hue)),

```

```

Mean_OD280_OD315_of_diluted_wines = c(
  mean(wines_data$OD280_OD315_of_diluted_wines),
  mean(class_one$OD280_OD315_of_diluted_wines),
  mean(class_two$OD280_OD315_of_diluted_wines),
  mean(class_three$OD280_OD315_of_diluted_wines)),

Mean_Proline = c(mean(wines_data$Proline),
  mean(class_one$Proline),
  mean(class_two$Proline),
  mean(class_three$Proline)))

average_table

```

	id	class_number	Mean_Alcohol	Mean_Malic_acid	Mean_Ash	Mean_Alcalinity_of_ash
1	1	overall	13.00062	2.336348	2.366517	19.49494
2	2	1	13.74475	2.010678	2.455593	17.03729
3	3	2	12.27873	1.932676	2.244789	20.23803
4	4	3	13.15375	3.333750	2.437083	21.41667
			Mean_Magnesium	Mean_Total_phenols	Mean_Flavanoids	Mean_Nonflavanoid_phenols
1			99.74157	2.295112	2.0292697	0.3618539
2			106.33898	2.840169	2.9823729	0.2900000
3			94.54930	2.258873	2.0808451	0.3636620
4			99.31250	1.678750	0.7814583	0.4475000
			Mean_Proanthocyanins	Mean_Color_intensity	Mean_Hue	
1			1.590899	5.058090	0.9574494	
2			1.899322	5.528305	1.0620339	
3			1.630282	3.086620	1.0562817	
4			1.153542	7.396250	0.6827083	
			Mean_OD280_OD315_of_diluted_wines	Mean_Proline		
1			2.611685	746.8933		
2			3.157797	1115.7119		
3			2.785352	519.5070		
4			1.683542	629.8958		

e

Since there are three different classes, one will need to do 3 comparisons, class 1 vs. class 2, class 1 vs class 3 and class 2 vs class 3. Firstly, extracting the data of level of phenols of each classes:

```
class_one_phenols <- class_one['Total_phenols']
class_two_phenols <- class_two['Total_phenols']
class_three_phenols <- class_three['Total_phenols']
```

For existing R function.

```
t_test_1_2 <- t.test(class_one_phenols, class_two_phenols)
t_test_1_2
```

Welch Two Sample t-test

```
data: class_one_phenols and class_two_phenols
t = 7.4206, df = 119.14, p-value = 1.889e-11
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.4261870 0.7364055
sample estimates:
mean of x mean of y
 2.840169  2.258873
```

```
t_test_1_3 <- t.test(class_one_phenols, class_three_phenols)
t_test_1_3
```

Welch Two Sample t-test

```
data: class_one_phenols and class_three_phenols
t = 17.12, df = 98.356, p-value < 2.2e-16
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 1.026801 1.296038
sample estimates:
mean of x mean of y
 2.840169  1.678750
```

```
t_test_2_3 <- t.test(class_two_phenols, class_three_phenols)
t_test_2_3
```

Welch Two Sample t-test

```
data: class_two_phenols and class_three_phenols
t = 7.0125, df = 116.91, p-value = 1.622e-10
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 0.4162855 0.7439610
sample estimates:
mean of x mean of y
 2.258873  1.678750
```

For manually conducting the t-test.

Then, calculating the mean, variance for each groups:

```
mean_one <- mean(class_one_phenols[,])
mean_one
```

```
[1] 2.840169
```

```
variance_one <- var(class_one_phenols[,])
variance_one
```

```
[1] 0.1148948
```

```
mean_two <- mean(class_two_phenols[,])
mean_two
```

```
[1] 2.258873
```

```
variance_two <- var(class_two_phenols[,])
variance_two
```

```
[1] 0.2974187
```

```
mean_three <- mean(class_three_phenols[,])
mean_three
```

```
[1] 1.67875
```



```
variance_three <- var(class_three_phenols[,])
variance_three
```

```
[1] 0.1274282
```

For different comparisons, assuming that the variances are different, first compute the t-statistics with formula: $t = \frac{(\hat{X}_1 - \hat{X}_2) - (\mu_1 - \mu_2)}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}}$, where \hat{X}_1 and \hat{X}_2 are the sample means, μ_1 and μ_2 are the means, S_1^2 and S_2^2 are the sample variances, n_1 and n_2 are the sizes.

Since the null hypothesis is that there is no difference between each class, $\mu_1 - \mu_2 = 0$, thus the t-statistics are:

```
t_1_2 <- (mean_one - mean_two) /
  (sqrt((variance_one / num_class_one) + variance_two / num_class_two))
t_1_2
```

```
[1] 7.420649
```

```
t_1_3 <- (mean_one - mean_three) /
  (sqrt((variance_one / num_class_one) + variance_three / num_class_three))
t_1_3
```

```
[1] 17.12025
```

```
t_2_3 <- (mean_two - mean_three) /
  (sqrt((variance_two / num_class_two) + variance_three / num_class_three))
t_2_3
```

```
[1] 7.012505
```

Next the degrees of freedom are defined as $\nu = \frac{\left(\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}\right)^2}{\frac{\left(\frac{S_1^2}{n_1}\right)}{n_1-1} + \frac{\left(\frac{S_2^2}{n_2}\right)}{n_2-1}}$, then rounding it down to find the degree of freedom. The results are:

```

nu_1_2 <- floor(
  (variance_one / num_class_one + variance_two / num_class_two)^2 /
  ((variance_one / num_class_one)^2 / (num_class_one - 1) +
   (variance_two / num_class_two)^2 / (num_class_two - 1))
)
nu_1_2

```

[1] 119

```

nu_1_3 <- floor(
  (variance_one / num_class_one + variance_three / num_class_three)^2 /
  ((variance_one / num_class_one)^2 / (num_class_one - 1) +
   (variance_three / num_class_three)^2 / (num_class_three - 1))
)
nu_1_3

```

[1] 98

```

nu_2_3 <- floor(
  (variance_two / num_class_two + variance_three / num_class_three)^2 /
  ((variance_two / num_class_two)^2 / (num_class_two - 1) +
   (variance_three / num_class_three)^2 / (num_class_three - 1))
)
nu_2_3

```

[1] 116

Define a function to manually compute the p-value of give t-statistics and degree of freedom:

```

#' Function used to compute the two-tail p-values
#' @param t_statistics, numeric, the t-statistics value
#' @param df, numeric, the degree of freedom of the model
#' @return p_value_two_tail, numeric, the derived p-value
compute_two_tail_p_value <- function(t_statistics, df){

  #' Function used to compute the probability density function of t-student distribution
  #' @param x, numeric, variable
  #' @param df, numeric, degree of freedom
  #' @return the probability density function

```

```

t_pdf <- function(x, df){
  return(gamma((df+1)/2) / (sqrt(df*pi) * gamma(df/2)) * (1 + (x^2)/df)^(-(df+1)/2))
}

p_value_two_tail <- 2 * integrate(t_pdf, t_statistics, Inf, df = df)$value

return(p_value_two_tail)
}
p_1_2 <- compute_two_tail_p_value(t_1_2, nu_1_2)
p_1_2

```

```
[1] 1.897952e-11
```

```

p_1_3 <- compute_two_tail_p_value(t_1_3, nu_1_3)
p_1_3

```

```
[1] 3.267661e-31
```

```

p_2_3 <- compute_two_tail_p_value(t_2_3, nu_2_3)
p_2_3

```

```
[1] 1.664716e-10
```

Through calculation, one can observe that the p-values of all three comparisons are extremely small. Thus one can argue that there is extremely strong evidence against the null hypothesis for each pairwise comparison. The differences in phenol levels between all the classes are statistically significant.

Problem 2

a

Import the data as `raw_table`.

```

raw_table <- read.table("AskAManager.csv", sep = ",", header = TRUE)
head(raw_table)

```

	X	Timestamp	How.old.are.you.	What.industry.do.you.work.in.
1	1	4/27/2021 11:02:10	25-34	Education (Higher Education)
2	2	4/27/2021 11:02:22	25-34	Computing or Tech
3	3	4/27/2021 11:02:38	25-34	Accounting, Banking & Finance
4	4	4/27/2021 11:02:41	25-34	Nonprofits
5	5	4/27/2021 11:02:42	25-34	Accounting, Banking & Finance
6	6	4/27/2021 11:02:46	25-34	Education (Higher Education)

	Job.title
1	Research and Instruction Librarian
2	Change & Internal Communications Manager
3	Marketing Specialist
4	Program Manager
5	Accounting Manager
6	Scholarly Publishing Librarian

If.your.job.title.needs.additional.context..please.clarify.here.

1
2
3
4
5
6

What.is.your.annual.salary...You.ll.indicate.the.currency.in.a.later.question..If.you.are.]

1
2
3
4
5
6

How.much.additional.monetary.compensation.do.you.get..if.any..for.example..bonuses.or.over

1
2
3
4
5
6

Please.indicate.the.currency If..Other...please.indicate.the.currency.here..

1	USD
2	GBP
3	USD
4	USD
5	USD
6	USD

If.your.income.needs.additional.context..please.provide.it.here.

1
2
3
4
5
6

What.country.do.you.work.in.

1 United States
2 United Kingdom
3 US
4 USA
5 US
6 USA

If.you.re.in.the.U.S...what.state.do.you.work.in. What.city.do.you.work.in.

1 Massachusetts Boston
2 Cambridge
3 Tennessee Chattanooga
4 Wisconsin Milwaukee
5 South Carolina Greenville
6 New Hampshire Hanover

How.many.years.of.professional.work.experience.do.you.have.overall.

1 5-7 years
2 8 - 10 years
3 2 - 4 years
4 8 - 10 years
5 8 - 10 years
6 8 - 10 years

How.many.years.of.professional.work.experience.do.you.have.in.your.field.

1 5-7 years
2 5-7 years
3 2 - 4 years
4 5-7 years
5 5-7 years
6 2 - 4 years

What.is.your.highest.level.of.education.completed. What.is.your.gender.

1 Master's degree Woman
2 College degree Non-binary
3 College degree Woman
4 College degree Woman
5 College degree Woman
6 Master's degree Man

What.is.your.race...Choose.all.that.apply..

1 White

2	White
3	White
4	White
5	White
6	White

b

In order to clean up the variable names, a rename will be conducted. The new variable names will be `id`, `timestamp`, `age`, `work_industry`, `job`, `job_context`, `annual_salary`, `compensation`, `currency`, `other_currency`, `income_context`, `country`, `state`, `city`, `overall_work_years`, `specific_work_years`, `education`, `gender`, `race`.

```
colnames(raw_table) <- c('id',
  'timestamp',
  'age',
  'work_industry',
  'job',
  'job_context',
  'annual_salary',
  'compensation',
  'currency',
  'other_currency',
  'income_context',
  'country', 'state',
  'city', 'overall_work_years',
  'specific_work_years',
  'education',
  'gender',
  'race')
head(raw_table)
```

	id	timestamp	age	work_industry
1	1	4/27/2021 11:02:10	25-34	Education (Higher Education)
2	2	4/27/2021 11:02:22	25-34	Computing or Tech
3	3	4/27/2021 11:02:38	25-34	Accounting, Banking & Finance
4	4	4/27/2021 11:02:41	25-34	Nonprofits
5	5	4/27/2021 11:02:42	25-34	Accounting, Banking & Finance
6	6	4/27/2021 11:02:46	25-34	Education (Higher Education)

	job	job_context	annual_salary
1	Research and Instruction Librarian		55000

2	Change & Internal Communications Manager	54600
3	Marketing Specialist	34000
4	Program Manager	62000
5	Accounting Manager	60000
6	Scholarly Publishing Librarian	62000
	compensation currency other_currency income_context country	
1	0 USD	United States
2	4000 GBP	United Kingdom
3	NA USD	US
4	3000 USD	USA
5	7000 USD	US
6	NA USD	USA
	state city overall_work_years specific_work_years	
1	Massachusetts Boston 5-7 years 5-7 years	
2	Cambridge 8 - 10 years 5-7 years	
3	Tennessee Chattanooga 2 - 4 years 2 - 4 years	
4	Wisconsin Milwaukee 8 - 10 years 5-7 years	
5	South Carolina Greenville 8 - 10 years 5-7 years	
6	New Hampshire Hanover 8 - 10 years 2 - 4 years	
	education gender race	
1	Master's degree Woman White	
2	College degree Non-binary White	
3	College degree Woman White	
4	College degree Woman White	
5	College degree Woman White	
6	Master's degree Man White	

c

In order to restrict the data to those being paid in USD, a logistical judgment has been down, which will yield the index of entries whose currency is USD or they have USD as their other_currency. After that, using mask to get the restricted table which is `usd_table`.

```
usd_table <- raw_table[raw_table['currency'] == 'USD'
                      | raw_table['other_currency'] == 'USD', ]
head(usd_table)
```

	id	timestamp	age	work_industry
1	1	4/27/2021 11:02:10	25-34	Education (Higher Education)
3	3	4/27/2021 11:02:38	25-34	Accounting, Banking & Finance
4	4	4/27/2021 11:02:41	25-34	Nonprofits

```

5 5 4/27/2021 11:02:42 25-34 Accounting, Banking & Finance
6 6 4/27/2021 11:02:46 25-34 Education (Higher Education)
7 7 4/27/2021 11:02:51 25-34 Publishing
                                job job_context annual_salary compensation
1 Research and Instruction Librarian                    55000             0
3           Marketing Specialist                        34000             NA
4           Program Manager                            62000             3000
5           Accounting Manager                         60000             7000
6     Scholarly Publishing Librarian                    62000             NA
7           Publishing Assistant                       33000             2000
currency other_currency income_context          country          state
1      USD                               United States Massachusetts
3      USD                               US              Tennessee
4      USD                               USA              Wisconsin
5      USD                               US South Carolina
6      USD                               USA New Hampshire
7      USD                               USA South Carolina
city overall_work_years specific_work_years      education gender
1    Boston           5-7 years           5-7 years Master's degree Woman
3 Chattanooga       2 - 4 years           2 - 4 years College degree Woman
4    Milwaukee       8 - 10 years           5-7 years College degree Woman
5    Greenville       8 - 10 years           5-7 years College degree Woman
6    Hanover          8 - 10 years           2 - 4 years Master's degree  Man
7    Columbia        2 - 4 years           2 - 4 years College degree Woman
race
1 White
3 White
4 White
5 White
6 White
7 White

```

For the number of observation:

```

total_num <- nrow(raw_table)
total_num

```

```
[1] 28062
```

```

usd_num <- nrow(usd_table)
usd_num

```



```
[1] 23382
```

```
diff_num <- total_num - usd_num  
diff_num
```

```
[1] 4680
```

By restricting the data to those being paid in USD, the number of observations decreases by 4680.

d

Assume everyone starts working at least they are 18. The impossible entry is that the maximum possible value of its age minus the lowest value in its years of experience in their field, and years of experience total respectively. If the result smaller than 18, this entry will be seen as impossible.

```
larger_age <- unlist(lapply(usd_table$age,  
                           function(x) max(  
                             as.numeric(  
                               unlist(  
                                 regmatches(  
                                   x, gregexpr("\\d+", x))))))  
  
smaller_overall_work <- unlist(lapply(usd_table$overall_work_years,  
                                     function(x) min(  
                                       as.numeric(  
                                         unlist(  
                                           regmatches(  
                                             x, gregexpr("\\d+", x))))))  
  
smaller_specific_work <- unlist(lapply(usd_table$specific_work_years,  
                                       function(x) min(  
                                         as.numeric(  
                                           unlist(  
                                             regmatches(  
                                               x, gregexpr("\\d+", x))))))
```

Thus the impossible index are as following, where TRUE means impossible.

```

overall_diff <- larger_age - smaller_overall_work
specific_diff <- larger_age - smaller_specific_work
overall_impossible <- overall_diff < 18
specific_impossible <- specific_diff < 18
impossible_index <- overall_impossible | specific_impossible
head(impossible_index)

```

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

Then the cleaned table is:

```

possible_usd_table <- usd_table[!impossible_index, ]
head(possible_usd_table)

```

	id	timestamp	age	work_industry
1	1	4/27/2021 11:02:10	25-34	Education (Higher Education)
3	3	4/27/2021 11:02:38	25-34	Accounting, Banking & Finance
4	4	4/27/2021 11:02:41	25-34	Nonprofits
5	5	4/27/2021 11:02:42	25-34	Accounting, Banking & Finance
6	6	4/27/2021 11:02:46	25-34	Education (Higher Education)
7	7	4/27/2021 11:02:51	25-34	Publishing

	job	job_context	annual_salary	compensation
1	Research and Instruction Librarian		55000	0
3		Marketing Specialist	34000	NA
4		Program Manager	62000	3000
5		Accounting Manager	60000	7000
6	Scholarly Publishing Librarian		62000	NA
7		Publishing Assistant	33000	2000

	currency	other_currency	income_context	country	state
1	USD			United States	Massachusetts
3	USD			US	Tennessee
4	USD			USA	Wisconsin
5	USD			US	South Carolina
6	USD			USA	New Hampshire
7	USD			USA	South Carolina

	city	overall_work_years	specific_work_years	education	gender
1	Boston	5-7 years	5-7 years	Master's degree	Woman
3	Chattanooga	2 - 4 years	2 - 4 years	College degree	Woman
4	Milwaukee	8 - 10 years	5-7 years	College degree	Woman
5	Greenville	8 - 10 years	5-7 years	College degree	Woman
6	Hanover	8 - 10 years	2 - 4 years	Master's degree	Man

	Columbia	2 - 4 years	2 - 4 years	College degree	Woman
race					
1	White				
3	White				
4	White				
5	White				
6	White				
7	White				

For the number of observations:

```
possible_num <- nrow(possible_usd_table)
possible_num
```

```
[1] 23321
```

```
diff_possible_num <- usd_num - possible_num
diff_possible_num
```

```
[1] 61
```

By restricting the data to those being paid in USD, the number of observations decreases by 61.

e

In this section, the IQR(interquartile range) will be used to identify the outliers, which means that the data fall below $Q1 - 1.5 \text{ IQR}$ or above $Q3 + 1.5 \text{ IQR}$ will be considered as outliers, then removed.

First, sorting the salary in ascending order, then calculating the Q1 and Q3. Finally, using $Q3 - Q1$ to get IQR.

```
sorted_salary <- sort(possible_usd_table$annual_salary)
Q_1 <- (sorted_salary[floor(1 + (possible_num - 1) / 4)] +
        sorted_salary[ceiling(1 + (possible_num - 1) / 4)]) / 2
Q_3 <- (sorted_salary[floor(1 + (possible_num - 1) * 3 / 4)] +
        sorted_salary[ceiling(1 + (possible_num - 1) * 3 / 4)]) / 2
IQR <- Q_3 - Q_1
IQR
```

```
[1] 55840
```

Then one can use this IQR to find the outliers:

```
min_salary <- Q_1 - 1.5 * IQR
max_salary <- Q_3 + 1.5 * IQR
final_table <- possible_usd_table[
  possible_usd_table['annual_salary'] >= min_salary &
  possible_usd_table['annual_salary'] <= max_salary, ]
head(final_table)
```

	id	timestamp	age	work_industry
1	1	4/27/2021 11:02:10	25-34	Education (Higher Education)
3	3	4/27/2021 11:02:38	25-34	Accounting, Banking & Finance
4	4	4/27/2021 11:02:41	25-34	Nonprofits
5	5	4/27/2021 11:02:42	25-34	Accounting, Banking & Finance
6	6	4/27/2021 11:02:46	25-34	Education (Higher Education)
7	7	4/27/2021 11:02:51	25-34	Publishing

	job	job_context	annual_salary	compensation
1	Research and Instruction Librarian		55000	0
3	Marketing Specialist		34000	NA
4	Program Manager		62000	3000
5	Accounting Manager		60000	7000
6	Scholarly Publishing Librarian		62000	NA
7	Publishing Assistant		33000	2000

	currency	other_currency	income_context	country	state
1	USD			United States	Massachusetts
3	USD			US	Tennessee
4	USD			USA	Wisconsin
5	USD			US	South Carolina
6	USD			USA	New Hampshire
7	USD			USA	South Carolina

	city	overall_work_years	specific_work_years	education	gender
1	Boston	5-7 years	5-7 years	Master's degree	Woman
3	Chattanooga	2 - 4 years	2 - 4 years	College degree	Woman
4	Milwaukee	8 - 10 years	5-7 years	College degree	Woman
5	Greenville	8 - 10 years	5-7 years	College degree	Woman
6	Hanover	8 - 10 years	2 - 4 years	Master's degree	Man
7	Columbia	2 - 4 years	2 - 4 years	College degree	Woman

	race
1	White
3	White

```
4 White
5 White
6 White
7 White
```

For the final sample size:

```
final_num <- nrow(final_table)
final_num
```

```
[1] 22407
```

Problem 3

a

```
#' Check the given number if it is a palindromic number or not
#' @param positive_int, numeric, a positive integer to be checked
#' @return result, list(logical, numeric), (isPalindromic, reserve)
isPalindromic <- function(positive_int){
  if(!is.numeric(positive_int)){

    warning("Input must be numeric.
            Attempting to convert to numeric...")

    positive_int <- as.numeric(positive_int)

    if(all(is.na(positive_int))){

      stop("Conversion to numeric failed")

    }
  }
  if(positive_int <= 0){
    stop("Input number is not positive")
  }

  digits <- as.numeric(unlist(strsplit(as.character(positive_int), "")))
  total_length <- length(digits)
```

```

mid_index <- total_length %/% 2
is_Palindromic <- TRUE
for(i in 1: mid_index){
  j <- total_length + 1 - i
  left_digits <- digits[i]
  right_digits <- digits[j]
  if (left_digits != right_digits){
    is_Palindromic <- FALSE
    break
  }
}
result <- list(isPalindromic=is_Palindromic, reserve=positive_int)
return(result)
}

result <- isPalindromic(728827)
result$isPalindromic

```

```
[1] TRUE
```

```
result$reserve
```

```
[1] 728827
```

b

```

#' For any given number, find the next palindromic number
#' @param positive_int, numeric, the give number
#' @return new_palindromic, numeric, the next palindromic number
nextPalindrome <- function(positive_int){
  if(!is.numeric(positive_int)){
    warning("Input must be numeric.
            Attempting to convert to numeric...")

    positive_int <- as.numeric(positive_int)

    if(all(is.na(positive_int))){

```

```

    stop("Conversion to numeric failed")
  }
}
if(positive_int <= 0){
  stop("Input number is not positive")
}

digits <- as.numeric(unlist(strsplit(as.character(positive_int), "")))
total_length <- length(digits)
if(total_length == 1){
  return(1 + positive_int)
}
mid_index <- total_length %/% 2
if (total_length %% 2 == 0){
  left_part <- digits[1: mid_index]
}else{
  index <- mid_index + 1
  left_part <- digits[1: index]
}
re_left <- rev(left_part[1: mid_index])
new_palindromic <- as.numeric(paste(c(left_part, re_left), collapse=""))
while (new_palindromic <= positive_int){
  left_part <- as.numeric(paste(left_part, collapse = ""))
  left_part <- left_part + 1
  left_part <- as.numeric(unlist(strsplit(as.character(left_part), "")))
  re_left <- rev(left_part[1: mid_index])
  new_palindromic <- as.numeric(paste(c(left_part, re_left), collapse=""))
}
return(new_palindromic)
}
nextPalindrome(7152)

```

```
[1] 7227
```

```
nextPalindrome(765431537)
```

```
[1] 765434567
```

c

i

```
nextPalindrome(391)
```

```
[1] 393
```

ii

```
nextPalindrome(9928)
```

```
[1] 9999
```

iii

```
nextPalindrome(19272719)
```

```
[1] 19277291
```

iv

```
nextPalindrome(109)
```

```
[1] 111
```

v

```
nextPalindrome(2)
```

```
[1] 3
```