

# FLS301\_2021

## Inferential Stats with R

### Empowerment for Local People Foundation

#### Lagos: Dec 8-9, 2021

Congratulation for making the leap. If you enjoy solving problems, you will enjoy R. But first let us enjoy a `garri` solution using 1 bowl and then 2 bowls of `water` by adding `garri` and `sugar`.

R simply works as a programming language that let us create objects and use reuse them as we want in subsequent iterations.

```
sugar <- 1+2

garri <- 8-2

water <- 1 #1 bowl

garri_solution_one <-sugar + garri + water
garri_solution_two <-sugar + garri + (2*water)
garri_solution_one
```

```
## [1] 10
```

```
garri_solution_two
```

```
## [1] 11
```

In R, we have to work in a specific folder our system called working directory. That is where everything happens!

Lets take a quick look at our current working directory.

```
getwd()
```

```
## [1] "/cloud/project"
```

You can use the function `setwd` to change/set a new working directory

```
setwd("/cloud/project")
```

We can set path directly. The easiest way to do this is to a set default working directory: `Session > Set Working Directory`.

## R Data Structures

**Vector:** A vector is simply a list of items that are of the same type. They are six types of atomic vectors-logical, integer, character, raw, double, and complex.

**Matrices:** A matrix is a two dimensional data set with columns and rows.

**List:** A list in R can contain many different data types inside it. A list is a collection of data which is ordered and changeable.

**Data Frames:** Data Frames are data displayed in a format as a table.

**Factors:** Factors are used to categorize data.

## OTHER R Syntax and Keywords

Objects: vector, list, matrix, array, factor, and data frame.

Functions.

Rows, Columns.

Method.

Loops.

Packages.

Working Directory.

## R Operators

Arithmetic operators (+, -, /, ^, x %% y) Assignment operators (<-) Comparison operators (==, !=, >=)  
Logical operators (&, |, !) Miscellaneous operators (%in%)

## Create dataset for our analysis

Here we want to create a dataset of 6 variables consisting data about 20 staff in organization. The variables are Gender, Weight, income, rating, marital status and whether staff stays in the city central.

```
Gender <- c("Male", "Male", "Male", "Male", "Male",  
           "Male", "Male", "Male", "Male", "Male",  
           "FeMale", "FeMale", "FeMale", "FeMale", "FeMale",  
           "FeMale", "FeMale", "FeMale", "FeMale", "FeMale")  
weight <- c(89, 75, 88, 75, 49, 89, 110, 120, 89, 75,  
           75, 76, 87, 110, 67, 76, 43, 55, 59, 60)  
income <- c(50000, 95000, 120000, 800000, 650000, 92000, 94000, 222000, 543000, 75000,  
           63000, 40000, 99000, 450000, 180000, 190000, 96000, 780000, 150000, 342000)  
rating <- c(5, 1, 2, 4, 9, 9, 8, 1, 9, 7,  
           5, 6, 6, 1, 1, 1, 3, 6, 9, 4)  
Marstatus <- c("Married", "Married", "Single", "Single", "Single",  
              "Single", "Divorced", "Single", "Married", "Single",  
              "Married", "Single", "Single", "Divorced", "Single",  
              "Single", "Divorced", "Single", "Married", "Divorced")  
CityCentral <- c("Yes", "No", "Yes", "Yes", "Yes", "Yes", "Yes", "Yes", "No", "Yes",  
               "No", "No", "Yes", "Yes", "No", "Yes", "No", "Yes", "No", "No")
```

## Binding two columns - cbind

Concatenate which is 'c' allows us to group different things into one object.

Next, we are taking 2 objects into a column and we telling R to use `cbind` to take these different columns and merge them as one data frame is an object in R.

```
officew <- as.data.frame(cbind(Gender, weight, income, rating, Marstatus,
                               CityCentral))
officew
```

```
##      Gender weight income rating Marstatus CityCentral
## 1    Male      89  50000      5   Married      Yes
## 2    Male      75  95000      1   Married      No
## 3    Male      88 120000      2    Single      Yes
## 4    Male      75  8e+05      4    Single      Yes
## 5    Male      49 650000      9    Single      Yes
## 6    Male      89  92000      9    Single      Yes
## 7    Male     110  94000      8 Divorced      Yes
## 8    Male     120 222000      1    Single      Yes
## 9    Male      89 543000      9   Married      No
## 10   Male      75  75000      7    Single      Yes
## 11 FeMale      75  63000      5   Married      No
## 12 FeMale      76  40000      6    Single      No
## 13 FeMale      87  99000      6    Single      Yes
## 14 FeMale     110 450000      1 Divorced      Yes
## 15 FeMale      67 180000      1    Single      No
## 16 FeMale      76 190000      1    Single      Yes
## 17 FeMale      43  96000      3 Divorced      No
## 18 FeMale      55 780000      6    Single      Yes
## 19 FeMale      59 150000      9   Married      No
## 20 FeMale      60 342000      4 Divorced      No
```

Factor is another way of calling categorical variable in R. The `as.data.frame` changes the factor (categorical) into a data frame without necessarily changing the class. The `c` only works if the number of rows in each variable is the same.

```
meanincome <- mean(officew$income)
```

```
## Warning in mean.default(officew$income): argument is not numeric or logical:
## returning NA
```

```
modeincome <- mode(officew$income)
modeincome
```

```
## [1] "character"
```

```
officew$income <- as.numeric(as.character(officew$income))
officew$Marstatus <- as.factor(as.character(officew$Marstatus))
sdincome <- sd(officew$income)
varincome <- var(officew$income)
varincome
```

```
## [1] 62240786842
```

```
class(officew$income)
```

```
## [1] "numeric"
```

```
head(officew, n = 5)
```

```
##      Gender weight income rating Marstatus CityCentral
## 1    Male      89  50000      5   Married      Yes
## 2    Male      75  95000      1   Married      No
## 3    Male      88 120000      2    Single      Yes
## 4    Male      75 800000      4    Single      Yes
```

```
## 5    Male      49 650000      9    Single      Yes
```

### summary stats

Let us use rbind (rowbind) to bind the rows of two different dataset together. The row names of the two datasets must be same for it to work

Create dataset for men with 3 variables

```
Gender1 <- c("Male", "Male", "Male", "Male", "Male",
             "Male", "Male", "Male", "Male", "Male", "Male")
weight1 <- c(89, 75, 88, 75, 49, 89, 110, 120, 89, NA, 75)

rating1 <- c(5, 1, 2, 4, 9, 9, 8, 1, 9, NA, 7)

officew_men22<- as.data.frame(cbind(Gender1, weight1, rating1))
officew_men22
```

```
##      Gender1 weight1 rating1
## 1      Male      89        5
## 2      Male      75        1
## 3      Male      88        2
## 4      Male      75        4
## 5      Male      49        9
## 6      Male      89        9
## 7      Male     110        8
## 8      Male     120        1
## 9      Male      89        9
## 10     Male    <NA>    <NA>
## 11     Male      75        7
```

Create dataset for women with 3 variables

```
Gender1 <- c("FeMale","FeMale", "FeMale","FeMale", "FeMale",
             "FeMale","FeMale","FeMale","FeMale", "FeMale", "FeMale")
weight1 <- c(75, 76, 87, 110, 67, 76, 43, NA, 55, 59, 60)

rating1 <- c( 4, 6, 4, 1, 1, 4, 3, 6, NA, 9, 4)

officew_women22<- as.data.frame(cbind(Gender1, weight1, rating1))
officew_women22
```

```
##      Gender1 weight1 rating1
## 1    FeMale      75        4
## 2    FeMale      76        6
## 3    FeMale      87        4
## 4    FeMale     110        1
## 5    FeMale      67        1
## 6    FeMale      76        4
## 7    FeMale      43        3
## 8    FeMale    <NA>        6
## 9    FeMale      55    <NA>
## 10   FeMale      59        9
## 11   FeMale      60        4
```

## Row Bind

We are binding both female and male dataset with `rbind`. Since they have the same number of rows, we can bind:

```
officew_full <- rbind(officew_men22, officew_women22)
officew_full
```

```
##      Gender1 weight1 rating1
## 1      Male      89        5
## 2      Male      75        1
## 3      Male      88        2
## 4      Male      75        4
## 5      Male      49        9
## 6      Male      89        9
## 7      Male     110        8
## 8      Male     120        1
## 9      Male      89        9
## 10     Male    <NA>    <NA>
## 11     Male      75        7
## 12  FeMale      75        4
## 13  FeMale      76        6
## 14  FeMale      87        4
## 15  FeMale     110        1
## 16  FeMale      67        1
## 17  FeMale      76        4
## 18  FeMale      43        3
## 19  FeMale    <NA>        6
## 20  FeMale      55    <NA>
## 21  FeMale      59        9
## 22  FeMale      60        4
```

## REMOVING NAs

```
officew_women22_nona <- officew_women22[!is.na(officew_women22$rating1)
                                          &!is.na(officew_women22$weight1), ]
officew_women22_nona
```

```
##      Gender1 weight1 rating1
## 1    FeMale      75        4
## 2    FeMale      76        6
## 3    FeMale      87        4
## 4    FeMale     110        1
## 5    FeMale      67        1
## 6    FeMale      76        4
## 7    FeMale      43        3
## 10   FeMale      59        9
## 11   FeMale      60        4
```

The new dataset `officew_women22_nona` has no missing values(NAs)

#Exporting files

```
library(openxlsx)# export to excel
library(haven)

write.csv(officew, "officew.csv") #export to csv
write_sav(officew, "officew.sav")#export to spss

#Importing files

library(readr)
officew_wd <- read_csv("officew.csv")

#Importing files from Github
#install.packages("readr")
#library(readxl)

library(openxlsx)# export to excel
library(RCurl)
x <- getURL("https://raw.githubusercontent.com/abiola1864/FLS301/main/officew.csv")
officew<- read.csv(text = x)
head(officew)
```

```
##      X Gender weight income rating Marstatus CityCentral
## 1 1   Male      89  50000      5   Married           Yes
## 2 2   Male      75  95000      1   Married           No
## 3 3   Male      88 120000      2   Single            Yes
## 4 4   Male      75 800000      4   Single            Yes
## 5 5   Male      49 650000      9   Single            Yes
## 6 6   Male      89  92000      9   Single            Yes
```

## Subsetting and Filtering

You will find two ways you can subset a data using base R. Additionally, with the subset and select functions, subset both Gender variable and select as the required row.

```
officew_women1 <- officew[officew$Gender == "FeMale",]
officew_women2 <- subset(officew, Gender == "FeMale")
officew_women3 <- subset(officew, Gender == "FeMale", select =
  c("weight"))
```

```
officew_women2
```

```
##      X Gender weight income rating Marstatus CityCentral
## 11 11 FeMale      75  63000      5   Married           No
## 12 12 FeMale      76  40000      6   Single           No
## 13 13 FeMale      87  99000      6   Single            Yes
## 14 14 FeMale     110 450000      1  Divorced            Yes
## 15 15 FeMale      67 180000      1   Single           No
## 16 16 FeMale      76 190000      1   Single            Yes
## 17 17 FeMale      43  96000      3  Divorced           No
## 18 18 FeMale      55 780000      6   Single            Yes
## 19 19 FeMale      59 150000      9   Married           No
## 20 20 FeMale      60 342000      4  Divorced           No
```

```
officew_women3
```

```
##      weight
## 11      75
## 12      76
## 13      87
## 14     110
## 15      67
## 16      76
## 17      43
## 18      55
## 19      59
## 20      60
```

```
# male
```

```
officew_men1 <- officew[officew$Gender == "Male",]
officew_men2 <- subset(officew, Gender == "Male")
officew_men3 <- subset(officew, Gender == "Male", select =
                      c("weight"))
```

We will be using several tools from the tidyr and dplyr packages to achieve data wrangling. Remember we already know some functions from this packages: drop\_na, etc...

```
summary(officew_men3$weight)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      49.0   75.0   88.5   85.9   89.0   120.0
```

```
summary(officew_women3$weight)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      43.00  59.25   71.00   70.80   76.00   110.00
```

standard deviation:

```
sd(officew$weight, na.rm = T)
```

```
## [1] 20.25957
```

For the sample size, we need to omit all missing values. the length(), which() and is.na() functions can help us:

```
length(which(!is.na(officew$weight)))
```

```
## [1] 20
```

Let us select only the 3rd and 5th variable (Income and marital status)

```
officew_3_5 <- officew[c(3,5)]
officew_3_5
```

```
##      weight rating
## 1      89      5
## 2      75      1
## 3      88      2
## 4      75      4
## 5      49      9
## 6      89      9
## 7     110      8
## 8     120      1
```

```
## 9      89      9
## 10     75      7
## 11     75      5
## 12     76      6
## 13     87      6
## 14    110      1
## 15     67      1
## 16     76      1
## 17     43      3
## 18     55      6
## 19     59      9
## 20     60      4
```

Let us exclude 3rd and 5th variable (Income and marital status)

```
officew_1_2_4 <- officew[c(-3,-5)]
officew_1_2_4
```

```
##      X Gender income Marstatus CityCentral
## 1  1  Male  50000   Married         Yes
## 2  2  Male  95000   Married         No
## 3  3  Male 120000   Single          Yes
## 4  4  Male 800000   Single          Yes
## 5  5  Male 650000   Single          Yes
## 6  6  Male  92000   Single          Yes
## 7  7  Male  94000   Divorced        Yes
## 8  8  Male 222000   Single          Yes
## 9  9  Male 543000   Married         No
## 10 10 Male  75000   Single          Yes
## 11 11 FeMale 63000   Married         No
## 12 12 FeMale 40000   Single          No
## 13 13 FeMale 99000   Single          Yes
## 14 14 FeMale 450000   Divorced        Yes
## 15 15 FeMale 180000   Single          No
## 16 16 FeMale 190000   Single          Yes
## 17 17 FeMale  96000   Divorced        No
## 18 18 FeMale 780000   Single          Yes
## 19 19 FeMale 150000   Married         No
## 20 20 FeMale 342000   Divorced        No
```

Here is another way of writing what we wrote above (including and not excluding)

```
office_1_2_4B <- officew[c(1,2,4)]
office_1_2_4B
```

```
##      X Gender income
## 1  1  Male  50000
## 2  2  Male  95000
## 3  3  Male 120000
## 4  4  Male 800000
## 5  5  Male 650000
## 6  6  Male  92000
## 7  7  Male  94000
## 8  8  Male 222000
## 9  9  Male 543000
## 10 10 Male  75000
```



```
## 11 11 FeMale 63000
## 12 12 FeMale 40000
## 13 13 FeMale 99000
## 14 14 FeMale 450000
## 15 15 FeMale 180000
## 16 16 FeMale 190000
## 17 17 FeMale 96000
## 18 18 FeMale 780000
## 19 19 FeMale 150000
## 20 20 FeMale 342000
```

Include 1st and 2nd variable(column), and 4 and the 4th and 5th observation(row)

```
officew_weight_inc_ <- officew[c(1:2),c(4:5)]
officew_weight_inc_
```

```
## income rating
## 1 50000 5
## 2 95000 1
```

#Conditional Subsetting

In R, | means it returns TRUE if one of the statement is TRUE. In R & means it returns TRUE if both elements are TRUE

We want to subset a dataframe of female staff whose income or if any staff earn N100,000 and above (AND). Returns the result for any of the conditions met.

```
office_f_hincome <- subset(officew, income >=100000 | Gender %in% "FeMale",
                           select=c(1:5))
office_f_hincome
```

```
## X Gender weight income rating
## 3 3 Male 88 120000 2
## 4 4 Male 75 800000 4
## 5 5 Male 49 650000 9
## 8 8 Male 120 222000 1
## 9 9 Male 89 543000 9
## 11 11 FeMale 75 63000 5
## 12 12 FeMale 76 40000 6
## 13 13 FeMale 87 99000 6
## 14 14 FeMale 110 450000 1
## 15 15 FeMale 67 180000 1
## 16 16 FeMale 76 190000 1
## 17 17 FeMale 43 96000 3
## 18 18 FeMale 55 780000 6
## 19 19 FeMale 59 150000 9
## 20 20 FeMale 60 342000 4
```

We want to subset a dataframe of female staff whose income is N100,000 and above (AND). The two conditions have to be met here.

```
office_f_hincome1 <- subset(officew, income >=100000 & Gender %in% "FeMale",
                           select=c(1:5))
office_f_hincome1
```

```
## X Gender weight income rating
## 14 14 FeMale 110 450000 1
```

```
## 15 15 FeMale      67 180000      1
## 16 16 FeMale      76 190000      1
## 18 18 FeMale      55 780000      6
## 19 19 FeMale      59 150000      9
## 20 20 FeMale      60 342000      4
```

```
#DPLYR
```

If you have not installed the packages, you have to install them first by removing the ash symbols!

```
# install.packages("tidyr")
# install.packages("dplyr")
library(tidyr)
library(dplyr)
```

We are not getting accurate interval level summary because R see it as a factor.

```
class(officew_women3$weight)
```

```
## [1] "integer"
```

```
class(officew_men3$weight)
```

```
## [1] "integer"
```

Let us change the class from character to numeric or interval

```
officew_women3$weight <-
  as.numeric(as.character(officew_women3$weight))
```

```
officew_men3$weight <-
  as.numeric(as.character(officew_men3$weight))
```

Let us check us again, great!

```
class(officew_women3$weight)
```

```
## [1] "numeric"
```

```
class(officew_men3$weight)
```

```
## [1] "numeric"
```

We can now compare the means of gender:

```
summary(officew_women3$weight)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  43.00   59.25   71.00   70.80   76.00  110.00
```

## Sampling Inference with t and Z test

#What is the probability that a random male staff will weigh above 125.3

```
rand_mstaff <- (125.3118 - mean(officew_men3$weight))/sd(officew_men3$weight)
1-pnorm(rand_mstaff)
```

```
## [1] 0.02275009
```

```
1- pnorm(2)
```

```
## [1] 0.02275013
```

What is the probability that a random male staff will be obese (weigh above 100) It is 2.3 percent

```
rand_menObestaff <- (100 - mean(officew_men3$weight))/sd(officew_men3$weight)
1-pnorm(rand_menObestaff)
```

```
## [1] 0.2371433
```

What is the probability that a random female staff will be . overweight (weigh above 100 is overweight) It is 0.05 percent (less than 1%)

```
rand_womenObestaff <- (100 - mean(officew_women3$weight))/
  sd(officew_women3$weight)
1-pnorm(rand_womenObestaff)
```

```
## [1] 0.05968218
```

## HYPOTHESIS TESTING

We know that the probability that the company will hire men with obesity is 2.3%, and for women is about 1%. It is clear that the company hires more men with obesity than women. However, what we do not know is if this difference is due to error in our random sampling or truly reflects the differences in the entire staff. We are going to use HYPOTHESIS TESTING. By assuming first, that the difference between them is zero referred to as NULL

#Method 1 and 2: P-value & T-test Do the t-test

```
t.test(officew_women3$weight, officew_men3$weight)
```

```
##
## Welch Two Sample t-test
##
## data: officew_women3$weight and officew_men3$weight
## t = -1.7555, df = 17.956, p-value = 0.09621
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -33.173987 2.973987
## sample estimates:
## mean of x mean of y
## 70.8 85.9
```

### Alternative check of the t-statistic:

This is similar to what we did earlier with sampling inference. HERE, the mean is Zero, and the figure (Weight) we want to get the probability for is the mean difference. We want to know the probability of the mean difference. H1 is the mean difference. H0 is the NULL

#6 steps to understanding the P-value

- 1 We want to test the probability that makes us believe that an effect (15) or difference between two groups is NOT happening by random chance
2. We assume the mean effect between these groups is zero meaning we assume that there is no effect
3. We run a ttest to get the prob of that effect happening and check the equivalent Probability. Note that we still assume our mean difference is zero
4. We can say that if that probability we get is 5% or less, then that is probability of that effect occurring at an assumption of a mean of zero.

5. It means the probability of that effect happening if we set our mean difference to zero is very low. Since the probability that it will occur is very low, we should not accept that our mean difference is zero.

6. So we reject that our mean difference is zero, and take the alternative hypothesis.

Remember that if the T-value is 2 (or 1.96) or more, it means the probability is 2.2% and 5% (two-tail) or less. This means we have 5% or less probability, that we will by chance get the difference in mean (15.1 -the effect) with the assumption that the null hypothesis is true (mean is zero). We are assuming that there is no difference, but that assumption will only occur 5% or less if the difference is -15.1. So due to this, we reject the null hypothesis and accept the alternative

## Method 1

First: Calculate Standard Errors for both groups

```
se.women <- sd(officew_women3$weight) / sqrt(length(officew_women3$weight))
se.women
```

```
## [1] 5.928837
```

```
se.men <- sd(officew_men3$weight) / sqrt(length(officew_men3$weight))
se.men
```

```
## [1] 6.231551
```

sum (standardized version of) both standard errors:

```
se.diff <- sqrt((se.women^2 + se.men^2))
se.diff
```

```
## [1] 8.601356
```

then calculate confidence intervals:

```
#  $t = (H1 - H0) / sem.diff$ 
```

```
mean.diff <- mean(officew_women3$weight) - mean(officew_men3$weight)
mean.diff
```

```
## [1] -15.1
```

The t-value is

```
t <- (mean.diff - mean(0)) / se.diff
t # bigger than 1.96?
```

```
## [1] -1.755537
```

calculating t-value at 95% confidence interval and 18 degree of freedom

```
qt(0.975, 18)
```

```
## [1] 2.100922
```

I use a critical t value for 0.05 significance and 18 degrees of freedom The degree of freedom is calculated by:

```
dof <- nrow(officew_women3$weight) + nrow(officew_men3$weight) - 2
dof
```

```
## numeric(0)
```

the confidence interval

```

(qt(0.975, 18) * se.diff)

## [1] 18.07078
upper.ci <- mean.diff + (qt(0.975, 18) * se.diff)

lower.ci <- mean.diff - (qt(0.975, 18) * se.diff)
lower.ci

## [1] -33.17078
upper.ci

## [1] 2.970779
#Quiz Example
weight_men <- c(89, 75, 88, 75, 49, 89, 110, 120, 89, 75)

weight_women <- c(75, 76, 87, 110, 67, 76, 43, 55, 59, 60)
weight_men

## [1] 89 75 88 75 49 89 110 120 89 75
weight_women

## [1] 75 76 87 110 67 76 43 55 59 60
#Method 1
t.test(weight_men,weight_women)

##
## Welch Two Sample t-test
##
## data: weight_men and weight_women
## t = 1.7555, df = 17.956, p-value = 0.09621
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -2.973987 33.173987
## sample estimates:
## mean of x mean of y
## 85.9 70.8
#Method 2 (by hand)
se.quiz.women <- sd(weight_women) / sqrt(length(weight_women ))
se.quiz.women

## [1] 5.928837
se.quiz.men <- sd(weight_men) / sqrt(length(weight_men ))
se.quiz.men

## [1] 6.231551
Let us check the mean and standard deviation of men and women
mean(weight_men )

## [1] 85.9

```

```
sd(weight_men)
```

```
## [1] 19.70589
```

```
mean(weight_women )
```

```
## [1] 70.8
```

```
sd(weight_women)
```

```
## [1] 18.74863
```

We then need to the standardized difference between the two standard errors

```
se.diff_quiz <- sqrt((se.quiz.men^2 + se.quiz.women^2 ))  
se.diff_quiz
```

```
## [1] 8.601356
```

The difference in the mean of the two gender

```
mean.diff55 <- mean(weight_men) - mean(weight_women )  
mean.diff55
```

```
## [1] 15.1
```

The upper bound of the confidence interval

```
upper.ci_quiz <- mean.diff55 + (qt(0.975, 18) * se.diff_quiz)  
upper.ci_quiz
```

```
## [1] 33.17078
```

The lower bound of the confidence interval

```
lower.ci_quiz <- mean.diff55 - (qt(0.975, 18) * se.diff_quiz)  
lower.ci_quiz
```

```
## [1] -2.970779
```

tvalue

```
tvaluequiz <- (mean.diff55 - mean(0)) / se.diff_quiz #t value  
tvaluequiz
```

```
## [1] 1.755537
```

Men weighing above 100 kg

```
obess_quizm<- (100 - mean(weight_men))/sd(weight_men)  
1-pnorm(obess_quizm)
```

```
## [1] 0.2371433
```

Wwomen weighing above 100 kg

```
obess_quizf<- (100 - mean(weight_women))/sd(weight_women)  
1-pnorm(obess_quizf)
```

```
## [1] 0.05968218
```

For question of proportion in the quiz

```

p <- 0.22
n <- 1200
se.prop<-sqrt(p*(1-p))/sqrt (n)

upperCI.prop <- p +(qt(0.975, (n-1))*se.prop)
upperCI.prop

## [1] 0.2434614

lowerCI.prop <- p -(qt(0.975, (n-1))*se.prop)
lowerCI.prop

## [1] 0.1965386

```

## Interpretation? Can we reject H0?

No, we cannot reject the null hypothesis that the difference in the mean of both gender is zero at 95% confidence. We are 95% confident that the difference in mean of both gender is zero. We can also say that we cannot reject the null hypothesis that the difference in the mean of both gender will happen by random chance 5 times or more out of every 100 occurrence.

Using the First Method (t-value test), the t-statistic is 1.96, and our t-value is -1.75. If we plot this in a graph, 1.75 falls within regions lower than 1.96 (0.05 p-value), but we were only ready to accept region at -1.96 and above it

Using the second method (p-value test), our p-value as well is higher than 0.05. Our P-value is the probability of getting a result as extreme as our test statistic, assuming our NULL hypothesis is true that there is no difference in the mean.

Using the third method (CI test), our confidence interval is -33 to 2.with the mean as 15.1. Zero (0)is within the confidence interval that we are 95 % confident the difference in mean between both gender can also be 0. We cannot reject the null hypothesis in this regard.

## ANOTHER EXAMPLE OF T-TEST WITH NORMAL DISTRIBUTION

Make simulations replicable:

```
set.seed(101112)
```

disable scientific notation:

```
options(scipen=999)
```

We start by creating two different normally distributed variables:

```

var1 <- rnorm(50, mean = 0, sd = 1)
var1

## [1] -0.75886627  1.35359814 -0.20107037 -0.44020778  1.29733664 -1.77972690
## [7] -0.83939342  0.95828995  0.42984356  0.57647495  0.02182424  0.05155345
## [13] -1.44215582  3.09711393  0.60303556  1.56984330 -0.30714872 -0.87877014
## [19]  0.99365026  1.10491075  0.05902839 -0.72572592  0.61683306  0.76522140
## [25]  1.12996161  0.04939211  0.64416160 -0.20130345  0.79669099  0.84365116
## [31] -1.07974123  0.91799752 -1.82842883 -0.16075665  0.14761583 -1.46805313
## [37]  1.52113035  0.97308093  0.02674715 -0.08527182 -2.18919556 -0.09255031
## [43]  1.12027192  0.36227175 -1.08154375  2.31478977 -1.77177149 -1.66071484

```

```
## [49] -0.76352840 0.56684880
var2 <- rnorm(100, mean = 0.5, sd = 3)
var2

## [1] 0.50209787 0.90784791 -0.22941116 3.51909605 1.28583811 3.24834278
## [7] 0.48008027 8.41920197 -0.93041455 4.62418359 2.88894985 -1.32263923
## [13] 3.31639820 -3.02120655 1.41778850 -0.63817913 1.07666747 1.78747131
## [19] 6.60213712 -4.44306297 3.67788246 3.30703994 -4.62364943 2.03825362
## [25] 4.84540154 -1.72471828 0.41937474 -2.84170751 3.65945560 -5.21100921
## [31] 4.37137774 -4.94866039 -5.60707819 -0.88447897 -0.52563202 -4.07731505
## [37] -0.29093857 2.05032961 6.91266138 0.02084897 -2.78871444 1.17091235
## [43] 1.35525897 3.02618274 -1.70754625 -1.62039019 6.22818944 -1.25320076
## [49] 0.34662568 2.01195870 0.38273763 -1.89338378 1.46254231 0.48318039
## [55] 1.15986634 -5.60941130 2.54299540 3.30143024 -1.77764401 -0.74128823
## [61] -2.51898754 0.52648564 -3.82201313 -3.18110417 -0.39275039 3.04378387
## [67] -3.34115600 -3.67115026 1.65222304 2.20707289 0.39242646 -4.34476888
## [73] 0.47902270 -2.37498287 -2.36238327 -0.29233467 0.44808159 4.60699570
## [79] 0.05713016 -0.77166218 2.89230534 1.50039635 1.43260044 -0.69679826
## [85] 4.09618435 -2.92627607 -5.84466681 0.49394818 -0.90040102 -1.41043344
## [91] 0.40839684 0.91381982 2.22293636 2.51697495 2.24030033 2.37997298
## [97] 0.87324808 4.85636485 -0.59974494 0.12610685
```

What are their means?

```
mean(var1)
```

```
## [1] 0.1031449
```

```
mean(var2)
```

```
## [1] 0.3705409
```

Is there a significant difference?

```
t.test(var1, var2)
```

```
##
## Welch Two Sample t-test
##
## data: var1 and var2
## t = -0.80081, df = 140.91, p-value = 0.4246
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.9275146 0.3927225
## sample estimates:
## mean of x mean of y
## 0.1031449 0.3705409
```

## Chi-square

Chi-Square test in R is a statistical method which used to determine if two categorical variables have a significant correlation between them. The difference with  $\chi^2$  is between the observed frequency ( $f_o$ ) and the expected frequency ( $f_e$ ).

$H_0$ : every i.v. category should have the same distribution across the d.v. as the total, i.e. i.v. doesn't matter.

Let us assume you that in the process of the review, an argument from one of your HR staff is that men are more single than women in the organization.



We are interested in knowing whether gender affect being single We want to know if whether either you are a male of female has an effect on the marital status

Here gender is the IV and Marital Status is the DV

STEP 1- DERIVE A CONTINGENCY TABLE Let us first divide our marital status into two concrete divisions -

Single Vs Not Single

```
table(officew$Marstatus)

##
## Divorced   Married   Single
##          4         5        11

table(officew$Marstatus=="Single")

##
## FALSE   TRUE
##       9    11

officew$NewMarStatus<- ifelse(officew$Marstatus=="Single",
                              "Single", "Not Single")
officew$NewMarStatus

## [1] "Not Single" "Not Single" "Single"      "Single"      "Single"
## [6] "Single"        "Not Single" "Single"      "Not Single" "Single"
## [11] "Not Single"   "Single"      "Single"      "Not Single" "Single"
## [16] "Single"        "Not Single" "Single"      "Not Single" "Not Single"
```

We then use the table function to show the cross tab

Converting NewMarStatus to a Factor

```
officew$NewMarStatus <- as.factor(as.character(officew$NewMarStatus))
officew$NewMarStatus

## [1] Not Single Not Single Single      Single      Single      Single
## [7] Not Single Single      Not Single Single      Not Single Single
## [13] Single      Not Single Single      Single      Not Single Single
## [19] Not Single Not Single
## Levels: Not Single Single

table(officew$Gender, officew$NewMarStatus)
```

```
##
##           Not Single Single
## FeMale           5        5
## Male             4        6
```

To get the percentages, we use prop.table function

```
prop.table(table( officew$NewMarStatus,officew$Gender), 2)

##
##           FeMale Male
## Not Single    0.5  0.4
## Single        0.5  0.6
```

50 percent of female are single, and 60 percent of males are single. We can say the effect of being a male is 10 percentage point higher for men than women.

```
prop.table(table(officew$NewMarStatus, officew$Gender), 1)
```

```
##
##           FeMale      Male
## Not Single 0.5555556 0.4444444
##   Single   0.4545455 0.5454545
```

```
#ALTERNATIVE 2: Install Gmodel package
```

```
install.packages("gmodels")
library(gmodels)
CrossTable(officew$Gender, officew$NewMarStatus)
```

```
##
##
##   Cell Contents
## |-----|
## |                N |
## | Chi-square contribution |
## |      N / Row Total |
## |      N / Col Total |
## |      N / Table Total |
## |-----|
##
##
## Total Observations in Table:  20
##
##
##           | officew$NewMarStatus
## officew$Gender | Not Single |      Single | Row Total |
## -----|-----|-----|-----|
##           FeMale |          5 |          5 |          10 |
##           |          0.056 |          0.045 |          |
##           |          0.500 |          0.500 |          0.500 |
##           |          0.556 |          0.455 |          |
##           |          0.250 |          0.250 |          |
## -----|-----|-----|-----|
##           Male |          4 |          6 |          10 |
##           |          0.056 |          0.045 |          |
##           |          0.400 |          0.600 |          0.500 |
##           |          0.444 |          0.545 |          |
##           |          0.200 |          0.300 |          |
## -----|-----|-----|-----|
## Column Total |          9 |          11 |          20 |
##           |          0.450 |          0.550 |          |
## -----|-----|-----|-----|
##
##
```

Let us assume we want to CONTROL for location. We think we can also use the location to staff (either they stay in the central city across both genders to know whether they are single or not. You just need to insert the new variable to the TABLE function

```
table(officew$Gender, officew$NewMarStatus, officew$CityCentral)
```

```
## , , = No
```

```
##
##
##      Not Single Single
## FeMale      4      2
## Male        2      0
##
## , , = Yes
##
##
##      Not Single Single
## FeMale      1      3
## Male        2      6
prop.table (table(officew$Gender, officew$NewMarStatus, officew$CityCentral), 3)

## , , = No
##
##
##      Not Single      Single
## FeMale 0.50000000 0.25000000
## Male   0.25000000 0.00000000
##
## , , = Yes
##
##
##      Not Single      Single
## FeMale 0.08333333 0.25000000
## Male   0.16666667 0.50000000
```

It might be more convenient to create two subsets of the data one for those who live in Central Area, and one for those who don't.

For people who live in the central area

```
officew_central <- officew[officew$CityCentral=="Yes",]
officew_central
```

```
##      X Gender weight income rating Marstatus CityCentral NewMarStatus
## 1  1  Male      89  50000      5  Married      Yes  Not Single
## 3  3  Male     88 120000      2   Single      Yes   Single
## 4  4  Male     75 800000      4   Single      Yes   Single
## 5  5  Male     49 650000      9   Single      Yes   Single
## 6  6  Male     89  92000      9   Single      Yes   Single
## 7  7  Male    110  94000      8 Divorced     Yes  Not Single
## 8  8  Male    120 222000      1   Single      Yes   Single
## 10 10 Male     75  75000      7   Single      Yes   Single
## 13 13 FeMale    87  99000      6   Single      Yes   Single
## 14 14 FeMale   110 450000      1 Divorced     Yes  Not Single
## 16 16 FeMale    76 190000      1   Single      Yes   Single
## 18 18 FeMale    55 780000      6   Single      Yes   Single
```

For people who DO NOT live in the central area

```
officew_Nocentral <- officew[officew$CityCentral=="No",]
officew_Nocentral
```

```
##      X Gender weight income rating Marstatus CityCentral NewMarStatus
## 2  2  Male     75  95000      1  Married      No  Not Single
```

```
## 9 9 Male 89 543000 9 Married No Not Single
## 11 11 FeMale 75 63000 5 Married No Not Single
## 12 12 FeMale 76 40000 6 Single No Single
## 15 15 FeMale 67 180000 1 Single No Single
## 17 17 FeMale 43 96000 3 Divorced No Not Single
## 19 19 FeMale 59 150000 9 Married No Not Single
## 20 20 FeMale 60 342000 4 Divorced No Not Single
```

With these subsets, you can obtain the cross-tabulations separately and in percentage form  
For people who live in the central area.

```
prop.table (table(officew_central$Gender, officew_central$NewMarStatus),2)
```

```
##
##      Not Single    Single
## FeMale 0.3333333 0.3333333
## Male   0.6666667 0.6666667
```

For people who DO NOT live in the central area

```
prop.table (table(officew_Nocentral$Gender, officew_Nocentral$NewMarStatus),2)
```

```
##
##      Not Single    Single
## FeMale 0.6666667 1.0000000
## Male   0.3333333 0.0000000
```

STEP 2: CONDUCT a t-test and check the chi square(x2) and p value

```
chisq.test(officew$Gender,officew$NewMarStatus,correct=FALSE)
```

```
## Warning in chisq.test(officew$Gender, officew$NewMarStatus, correct = FALSE):
## Chi-squared approximation may be incorrect
##
## Pearson's Chi-squared test
##
## data:  officew$Gender and officew$NewMarStatus
## X-squared = 0.20202, df = 1, p-value = 0.6531
```

It gave the warning because many of the expected values will be very small and therefore the approximations of p may not be right.

In R you can use `chisq.test(a, simulate.p.value = TRUE)` to use simulate p values.

```
chisq.test(officew$Gender,officew$NewMarStatus, simulate.p.value = TRUE)
```

```
##
## Pearson's Chi-squared test with simulated p-value (based on 2000
## replicates)
##
## data:  officew$Gender and officew$NewMarStatus
## X-squared = 0.20202, df = NA, p-value = 1
```

However, with such small cell sizes, all estimates will be poor. It might be good to just test pass vs. fail (deleting “no show”).

Either with chi-square or logistic regression. Indeed, since it is pretty clear that the pass/fail grade is a dependent variable, logistic regression might be better

## Correlation

packages:

```
install.packages("corrplot") # Install the corrplot library, for nice-looking
# correlation plots. Do this once.
install.packages("ggplot2") # Install the ggplot2 package, for high-quality
# graphs
install.packages("cowplot") # Install the cowplot package, to arrange plots
# into a grid
install.packages("ggpubr")
```

```
library(corrplot) # Plotting nice correlation matrix
library(cowplot) # arranging plots into a grid
library(ggplot2) # high-quality graphs
```

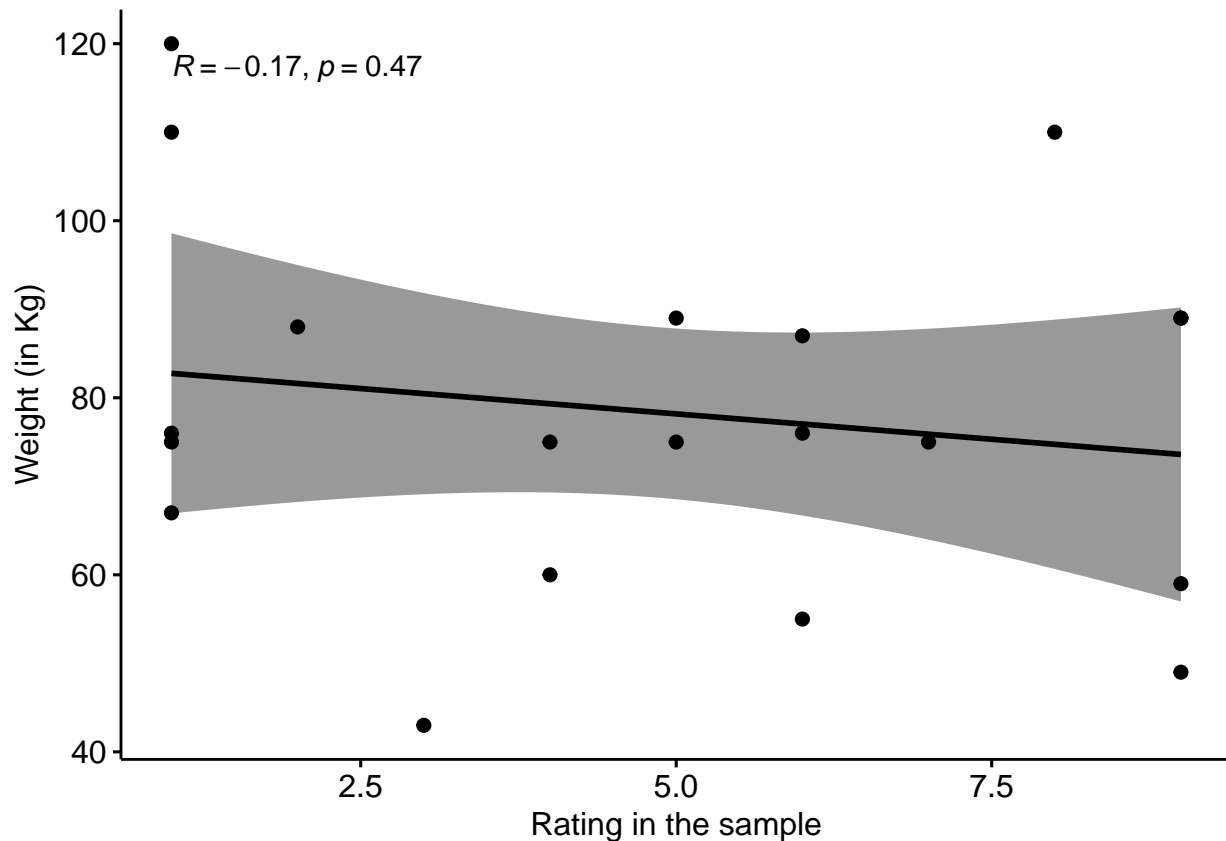
**Optional packages:** check the data properties

```
str(officew)
```

```
## 'data.frame': 20 obs. of 8 variables:
## $ X : int 1 2 3 4 5 6 7 8 9 10 ...
## $ Gender : chr "Male" "Male" "Male" "Male" ...
## $ weight : int 89 75 88 75 49 89 110 120 89 75 ...
## $ income : num 50000 95000 120000 800000 650000 92000 94000 222000 543000 75000 ...
## $ rating : int 5 1 2 4 9 9 8 1 9 7 ...
## $ Marstatus : chr "Married" "Married" "Single" "Single" ...
## $ CityCentral : chr "Yes" "No" "Yes" "Yes" ...
## $ NewMarStatus: Factor w/ 2 levels "Not Single","Single": 1 1 2 2 2 2 1 2 1 2 ...
```

plot the graph , use y as income

```
library("ggpubr")
ggscatter(officew, x = "rating", y = "weight",
          add = "reg.line", conf.int = TRUE,
          cor.coef = TRUE, cor.method = "pearson",
          xlab = "Rating in the sample", ylab = "Weight (in Kg)")
```



```
cor(officew$weight,officew$rating)
```

```
## [1] -0.1713778
```

```
str(officew)
```

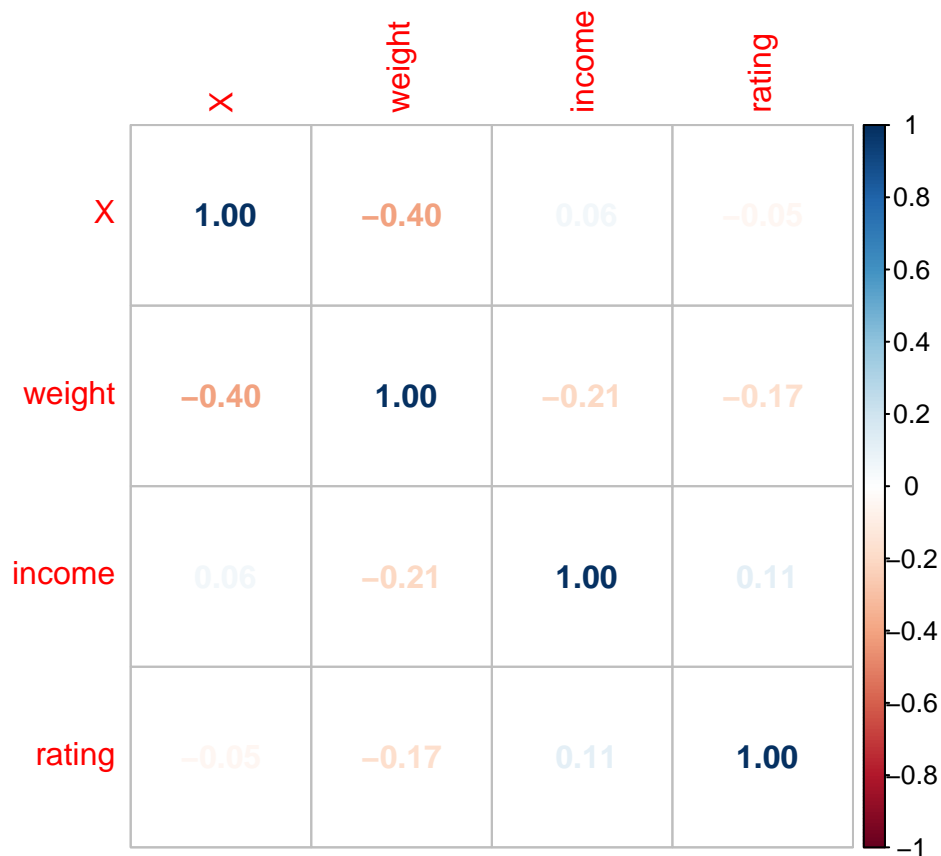
```
## 'data.frame':  20 obs. of  8 variables:
## $ X          : int  1 2 3 4 5 6 7 8 9 10 ...
## $ Gender      : chr  "Male" "Male" "Male" "Male" ...
## $ weight      : int  89 75 88 75 49 89 110 120 89 75 ...
## $ income      : num  50000 95000 120000 800000 650000 92000 94000 222000 543000 75000 ...
## $ rating      : int  5 1 2 4 9 9 8 1 9 7 ...
## $ Marstatus   : chr  "Married" "Married" "Single" "Single" ...
## $ CityCentral : chr  "Yes" "No" "Yes" "Yes" ...
## $ NewMarStatus: Factor w/ 2 levels "Not Single","Single": 1 1 2 2 2 2 1 2 1 2 ...
```

```
corr1<- cor(officew[c(-2,-6,-7,-8)]) #we do it without the
#last non-numeric variable "type" which are indexed in 3, 4 and 5
corr1
```

```
##           X      weight      income      rating
## X          1.00000000 -0.4037699  0.05917688 -0.04549472
## weight -0.40376994  1.00000000 -0.20964475 -0.17137780
## income  0.05917688 -0.2096447  1.00000000  0.11266232
## rating -0.04549472 -0.1713778  0.11266232  1.00000000
```

Nice correlation matrix ?corrplot

```
library(corrplot)
plot1<-corrplot(corr1, method = "number") # Try with different methods!
```



plot1

```
## $corr
##           X      weight      income      rating
## X      1.00000000 -0.4037699  0.05917688 -0.04549472
## weight -0.40376994  1.00000000 -0.20964475 -0.17137780
## income  0.05917688 -0.2096447  1.00000000  0.11266232
## rating -0.04549472 -0.1713778  0.11266232  1.00000000
##
## $corrPos
##      xName  yName x y      corr
## 1      X      X 1 4  1.00000000
## 2      X weight 1 3 -0.40376994
## 3      X income 1 2  0.05917688
## 4      X rating 1 1 -0.04549472
## 5 weight      X 2 4 -0.40376994
## 6 weight weight 2 3  1.00000000
## 7 weight income 2 2 -0.20964475
## 8 weight rating 2 1 -0.17137780
## 9 income      X 3 4  0.05917688
## 10 income weight 3 3 -0.20964475
## 11 income income 3 2  1.00000000
## 12 income rating 3 1  0.11266232
## 13 rating      X 4 4 -0.04549472
## 14 rating weight 4 3 -0.17137780
## 15 rating income 4 2  0.11266232
## 16 rating rating 4 1  1.00000000
```

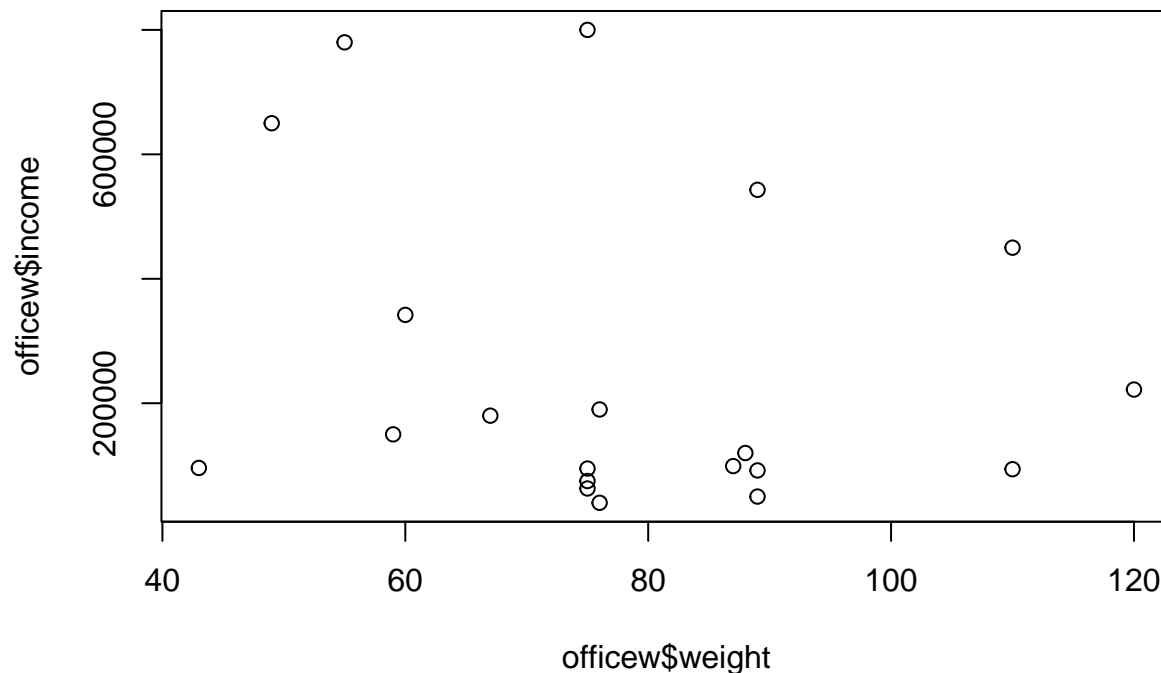
```
##
## $arg
## $arg$type
## [1] "full"
```

## REGRESSION

### STEP 1, CHECK THE PLOT

How does this look? Let's plot the data!

```
plot2<-plot(officew$weight, officew$income) # First x axis, then y axis:
```



```
plot2
```

```
## NULL
```

Our independent variable is income

### STEP 2 Run your first (bivariate) regression

```
myfirstreg <- lm(weight ~ income, data= officew) #First you run it
summary(myfirstreg) #Then you see the output
```

```
##
## Call:
## lm(formula = weight ~ income, data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -38.083 -13.100  -4.760   7.456  41.062
##
```



```
## Coefficients:
##           Estimate Std. Error t value      Pr(>|t|)
## (Intercept) 82.71765777  6.61544011   12.50 0.00000000026 ***
## income      -0.00001702  0.00001872   -0.91      0.375
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 20.35 on 18 degrees of freedom
## Multiple R-squared:  0.04395,    Adjusted R-squared:  -0.009163
## F-statistic: 0.8275 on 1 and 18 DF,  p-value: 0.375
```

```
confint(myfirstreg, level=0.99)
```

```
##           0.5 %          99.5 %
## (Intercept) 63.67550720175 101.75980833575
## income      -0.00007089542   0.00003684625
```

```
#STEP 3 PRINT your Result
```

```
install.packages("stargazer") # Install stargazer, for nice-looking regression
# tables. Do this once
library(stargazer)
# Lets get a nice table out of it
stargazer(myfirstreg, title="Regression Results", out="reg.txt")
```

```
##
## % Table created by stargazer v.5.2.2 by Marek Hlavac, Harvard University. E-mail: hlavac at fas.harvard.edu
## % Date and time: Tue, Dec 07, 2021 - 23:33:34
## \begin{table}[!htbp] \centering
##   \caption{Regression Results}
##   \label{}
##   \begin{tabular}{@{\extracolsep{5pt}}lc}
##     \hline
##     \hline \hline
##     & \multicolumn{1}{c}{\textit{Dependent variable:}} & \\
##     \cline{2-2}
##     \hline & weight & \\
##     \hline \hline
##     income & $-0.00002$ & \\
##     & (0.00002) & \\
##     & & \\
##     Constant & 82.718*** & \\
##     & (6.615) & \\
##     & & \\
##     \hline \hline
##     Observations & 20 & \\
##     R2 & 0.044 & \\
##     Adjusted R2 & -0.009 & \\
##     Residual Std. Error & 20.352 (df = 18) & \\
##     F Statistic & 0.827 (df = 1; 18) & \\
##     \hline
##     \hline \hline
##     \textit{Note:} & \multicolumn{1}{r}{*} & p < 0.1; **} & p < 0.05; ***} & p < 0.01 \\
##     \end{tabular}
##   \end{table}
```

#STEP 4 Interpret the result

```
options(scipen=999) #run this once to turn off scientific notation in
#your reg output
myfirstreg2 <- lm(weight ~ income+rating, data= officew) #First you run it
summary(myfirstreg2) #Then you see the output
```

```
##
## Call:
## lm(formula = weight ~ income + rating, data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -39.714 -14.604  -4.316   8.731  37.258
##
## Coefficients:
##              Estimate Std. Error t value    Pr(>|t|)
## (Intercept) 87.21803671  9.77667182   8.921 0.0000000803 ***
## income      -0.00001566  0.00001916  -0.817    0.425
## rating      -1.00034050  1.57672115  -0.634    0.534
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 20.7 on 17 degrees of freedom
## Multiple R-squared:  0.06606,    Adjusted R-squared:  -0.04381
## F-statistic: 0.6013 on 2 and 17 DF,  p-value: 0.5594
```

## MULTICOLLINEARITY (VIF & TOLERANCE) & Post Treatment Effect

```
install.packages("carData")
install.packages("car")
library(car)
library(carData)
```

VIF and Tolerance Let us test for the multicollinearity of both rating and income on our dependent variable  
'VIF test The square root of the VIF tells us by which factor at which the standard error for the coefficient of the IV will be larger than if that if it had 0 correlation with other independent variables.

?vif

```
vif(myfirstreg2)
```

```
##      income      rating
## 1.012856 1.012856
```

The Standard Error of the CE of income will be inflated by 1.012 if we include it in the model.

Tolerance is proportion of the model's independent variables not explained by other independent variables. The tolerance is the inverse of the vif and is the Percent of variance in the predictor that cannot be accounted for by other predictors.

```
1/vif(myfirstreg2)
```

```
##      income      rating
## 0.9873072 0.9873072
```

For example, if you run the VIF, 98 percent of the variance of income cannot be explained by other. This is where there is no correlation. It is what is unique to this variable income, that can't be explained by any other in the set

## Post Treatment Bias

```
library(AER)
data("Fatalities")

fatal <- lm(fatal~beertax + youngdrivers + miles + pop, data = Fatalities)
summary(fatal) # model with a number of covariates to isolate effect of drunk driving
```

```
##
## Call:
## lm(formula = fatal ~ beertax + youngdrivers + miles + pop, data = Fatalities)
##
## Residuals:
```

	Min	1Q	Median	3Q	Max
	-1196.13	-110.66	-7.06	116.84	1355.82

```
##
## Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-811.074536041	146.194713239	-5.548	0.000000059272093 ***
beertax	225.858365309	30.185471310	7.482	0.000000000000664 ***
youngdrivers	1133.885402562	591.746236039	1.916	0.0562 .
miles	0.065143934	0.009817656	6.635	0.000000000132276 ***
pop	0.000182334	0.000002887	63.148	< 0.0000000000000002 ***

```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 252.4 on 331 degrees of freedom
## Multiple R-squared:  0.9279, Adjusted R-squared:  0.927
## F-statistic: 1064 on 4 and 331 DF, p-value: < 0.00000000000000022

fatal.ptb <- lm(fatal ~ beertax + youngdrivers + miles + pop + spirits,
               data = Fatalities)
summary(fatal.ptb) # adding control for the mechanism (spirits consumption)
```

```
##
## Call:
## lm(formula = fatal ~ beertax + youngdrivers + miles + pop + spirits,
##     data = Fatalities)
##
## Residuals:
```

	Min	1Q	Median	3Q	Max
	-1191.92	-110.87	-5.48	118.65	1355.78

```
##
## Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-773.900635635	154.946661376	-4.995	0.000000095728464 ***
beertax	224.347841268	30.278022192	7.410	0.000000000000107 ***
youngdrivers	1105.951671017	593.407198244	1.864	0.0632 .
miles	0.064623804	0.009850549	6.560	0.00000000020754 ***
pop	0.000182124	0.000002904	62.719	< 0.0000000000000002 ***

```
## spirits      -14.863322043   20.407940241  -0.728           0.4669
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 252.6 on 330 degrees of freedom
## Multiple R-squared:  0.928, Adjusted R-squared:  0.9269
## F-statistic: 850.3 on 5 and 330 DF,  p-value: < 0.00000000000000022
stargazer(fatal, fatal.ptb, type = "text", data = Fatalities,
          out = "fatal.reg.txt")
```

```
##
## =====
##                               Dependent variable:
##                               -----
##                               fatal
##                               (1)                (2)
## -----
## beertax                225.858***            224.348***
##                        (30.185)              (30.278)
##
## youngdrivers            1,133.885*            1,105.952*
##                        (591.746)              (593.407)
##
## miles                   0.065***             0.065***
##                        (0.010)              (0.010)
##
## pop                    0.0002***             0.0002***
##                        (0.00000)            (0.00000)
##
## spirits                -14.863
##                        (20.408)
##
## Constant               -811.075***           -773.901***
##                        (146.195)            (154.947)
## -----
## Observations              336                336
## R2                       0.928                0.928
## Adjusted R2              0.927                0.927
## Residual Std. Error    252.400 (df = 331)    252.579 (df = 330)
## F Statistic            1,064.206*** (df = 4; 331) 850.264*** (df = 5; 330)
## =====
## Note:                    *p<0.1; **p<0.05; ***p<0.01
##
```

```
## =====
## Statistic    N      Mean      St. Dev.      Min      Pctl(25)      Pctl(75)      Max
## -----
## spirits      336      1.754      0.684      0.790      1.300      2.012      4.900
## unemp        336      7.347      2.533      2.400      5.475      8.900     18.000
## income       336    13,880.180    2,253.046    9,513.762    12,085.850    15,175.120    22,193.460
## emppop       336     60.806      4.722     42.993     57.691     64.413     71.269
## beertax      336      0.513      0.478      0.043      0.209      0.652      2.721
## baptist      336      7.157      9.763      0.000      0.627     13.127     30.356
## mormon       336      2.802      9.665      0.100      0.272      0.629     65.916
```

## drinkage	336	20.456	0.899	18	20	21	21
## dry	336	4.267	9.501	0.000	0.000	2.425	45.792
## youngdrivers	336	0.186	0.025	0.073	0.170	0.202	0.282
## miles	336	7,890.754	1,475.659	4,576.346	7,182.539	8,504.015	26,148.270
## fatal	336	928.664	934.051	79	293.8	1,063.5	5,504
## nfatal	336	182.583	188.431	13	53.8	212	1,049
## sfatal	336	109.949	108.540	8	35	131	603
## fatal1517	336	62.610	55.729	3	25.8	77	318
## nfatal1517	336	12.262	12.253	0	4	15.2	76
## fatal1820	336	106.661	104.224	7	38	130.2	601
## nfatal1820	336	33.527	33.238	0	11	44	196
## fatal2124	336	126.872	131.789	12	42	150.5	770
## nfatal2124	336	41.378	42.930	1	13	49	249
## afatal	336	293.333	303.581	24.600	90.498	363.958	2,094.900
## pop	336	4,930,272.000	5,073,704.000	478,999.700	1,545,251.000	5,751,735.000	28,314,028.000
## pop1517	336	230,815.500	229,896.300	21,000.020	71,749.930	270,500.200	1,172,000.000
## pop1820	336	249,090.400	249,345.600	20,999.960	76,962.120	308,311.400	1,321,004.000
## pop2124	336	336,389.900	345,304.400	30,000.160	103,500.000	413,000.100	1,892,998.000
## milestot	336	37,101.490	37,454.370	3,993	11,691.5	44,139.8	241,575
## unempus	336	7.529	1.479	5.500	6.200	9.600	9.700
## emppopus	336	59.971	1.585	57.800	57.900	61.500	62.300
## gsp	336	0.025	0.043	-0.124	0.001	0.057	0.142
## -----							

Here controlling for the mechanism causes part of the effect of beertax to be mathematically “soaked up”. Admittedly, the effect is a little weak.

## Dummy Variable and Binomial Regression

Convert gender to dummy variable, where male is 1 and female is 0. Male is our baseline variable

```
officew$highincome<- ifelse(officew$income>120000, 1, 0)
officew$highincome
```

```
## [1] 0 0 0 1 1 0 0 1 1 0 0 0 0 1 1 1 0 1 1 1
```

```
officew$overweight<- ifelse(officew$weight >=100, 1, 0)
officew$overweight
```

```
## [1] 0 0 0 0 0 0 1 1 0 0 0 0 0 1 0 0 0 0 0 0
```

## Run regression

```
reg11 <- lm(overweight ~ Gender,data = officew)
summary (reg11)
```

```
##
## Call:
## lm(formula = overweight ~ Gender, data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
##    -0.2    -0.2    -0.1    -0.1     0.9
##
## Coefficients:
```

```
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.1000     0.1179   0.849   0.407
## GenderMale     0.1000     0.1667   0.600   0.556
##
## Residual standard error: 0.3727 on 18 degrees of freedom
## Multiple R-squared:  0.01961,    Adjusted R-squared:  -0.03486
## F-statistic:  0.36 on 1 and 18 DF,  p-value: 0.556

#INTEPRETE THE RESULT Men who are overweight will weigh 10 kg higher than the reference group -
females who are overweighted
```

## Run regression

```
options(scipen = 999)
reg10<- lm(overweight ~ Gender + officew$highincome,data = officew)
summary (reg10)

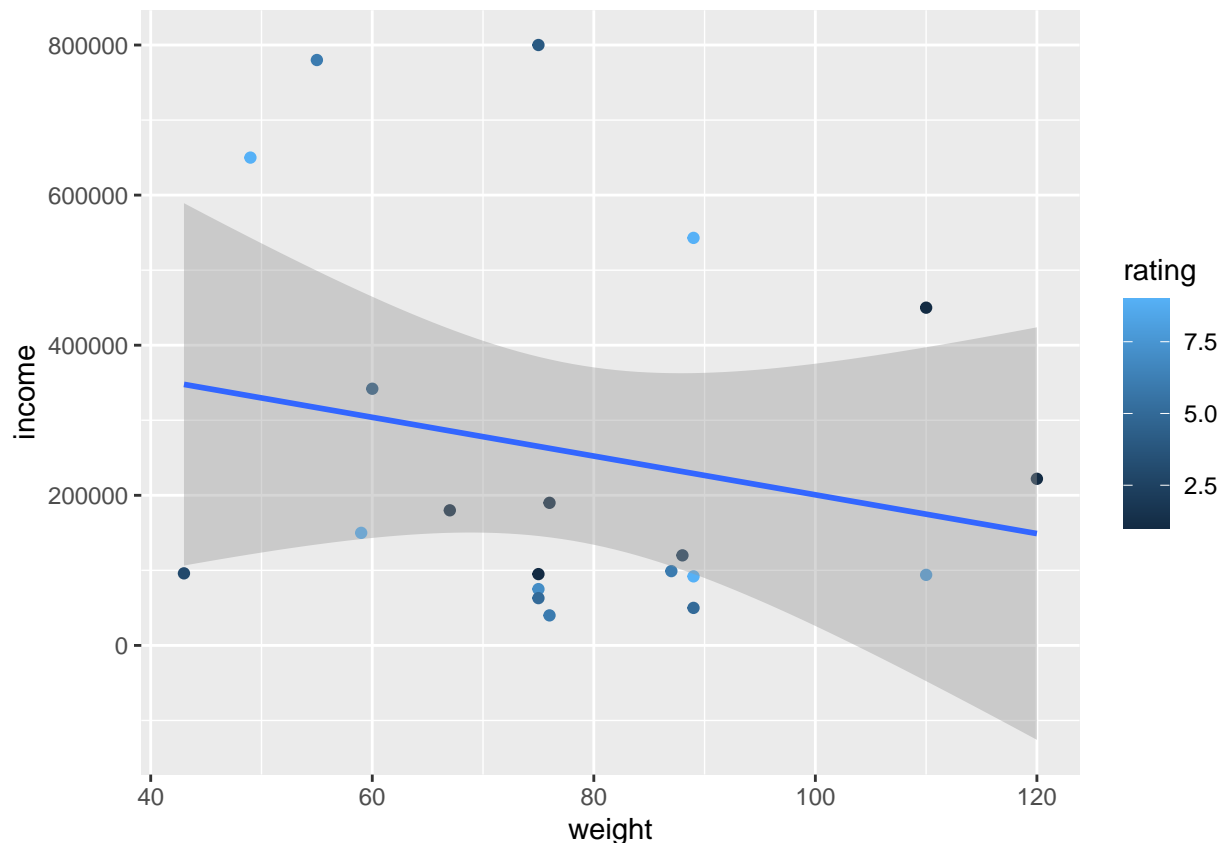
##
## Call:
## lm(formula = overweight ~ Gender + officew$highincome, data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.275 -0.150 -0.150 -0.025  0.850
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)         0.0250     0.1580   0.158   0.876
## GenderMale           0.1250     0.1724   0.725   0.478
## officew$highincome   0.1250     0.1724   0.725   0.478
##
## Residual standard error: 0.3777 on 17 degrees of freedom
## Multiple R-squared:  0.04902,    Adjusted R-squared:  -0.06286
## F-statistic: 0.4381 on 2 and 17 DF,  p-value: 0.6523
```

This suggests that, after effects of highincome are taken into account, men will weight 12kg higher than the reference group (women).

## GGPLOT and INTERACTION EFFECT

Graphically using ggplot

```
plot3<-ggplot(officew, aes(x = weight, y = income, colour = rating)) +
  geom_point() +
  geom_smooth(method = "lm")
plot3
```



What if we are interested on how the effect of `highincome` works across `Gender` which is the interaction effect of both variables. You will use \*

```
options(scripen = 100, "digits" = 3)
reg16 <- lm(overweight ~ Gender + highincome + Gender*highincome, data = officew)
summary(reg16)
```

```
##
## Call:
## lm(formula = overweight ~ Gender + highincome + Gender * highincome,
##     data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.250 -0.167 -0.167  0.000  0.833
##
## Coefficients:
##              Estimate      Std. Error t value
## (Intercept)  0.00000000000000621  0.1943203969393503261    0.00
## GenderMale   0.166666666666666019  0.2508665537248394584    0.66
## highincome   0.1666666666666665464  0.2508665537248394584    0.66
## GenderMale:highincome -0.0833333333333331761  0.3547788826234666293   -0.23
##
##              Pr(>|t|)
## (Intercept)      1.00
## GenderMale       0.52
## highincome       0.52
## GenderMale:highincome 0.82
##
```

```
## Residual standard error: 0.389 on 16 degrees of freedom
## Multiple R-squared: 0.0523, Adjusted R-squared: -0.125
## F-statistic: 0.294 on 3 and 16 DF, p-value: 0.829

reg15 <- lm(weight ~ Gender + highincome+ Gender*highincome,data = officew)
summary (reg15)
```

```
##
## Call:
## lm(formula = weight ~ Gender + highincome + Gender * highincome,
##     data = officew)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -34.25 -12.29   0.83    5.75   38.83
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      70.250     10.162   6.91 0.0000035 ***
## GenderMale       17.417     13.120   1.33    0.20
## highincome        0.917     13.120   0.07    0.95
## GenderMale:highincome -5.333     18.554  -0.29    0.78
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 20.3 on 16 degrees of freedom
## Multiple R-squared: 0.152, Adjusted R-squared: -0.00647
## F-statistic: 0.959 on 3 and 16 DF, p-value: 0.436
```

Interpretation.

You will also notice that the Rsquared has increased to 15 percent which means the model now account more variation of the dependent variable **weight**. That means explaining the effect of gender on **weight** works through the **income** staff receives.

The **weight** of men with **higherincome** is reduced by 5kg compared to women with **higherincome**. However, on average, men weigh (17.4kg -5.3kg) about 11.9kg more than effect of **income** held constant.

The average **weight** of men, the effect of **income** held constant, can still be derived as  $(70+17.4-(5.333)) = 82.1\text{kg}$ . The average **weight** of men = 70.2 kg (which is the intercept).

The **weight** of men over women  $(82.1 - 70.2)\text{kg}$  is 11.9kg which is what we got earlier.