

R Final Project : Breast Cancer Classification :: Cancer Cases

Utpal Mishra - 20207425
25 December 2020

Import Libraries

```
library(psych)

## Warning: package 'psych' was built under R version 3.6.3
```

Import Data

```
library(readxl)

data <- read_csv("E:/UCD/Lectures/Semester 1/Data Programming with R/Final Project/CancerCases.csv")
data

##           Cancer    Cases
## 1      Breast 2261419
## 2      Lung 2366771
## 3 Colorectum 1931590
## 4   Prostate 1414259
## 5    Stomach 1089103
## 6     Liver  905677
## 7 Cervix uteri 604127
## 8 Other cancers 8879843
```

Data Analysis

```
summary(data)

##           Cancer    Cases
## Breast      :1 Min.   : 604127
## Cervix uteri:1 1st Qu.:1043246
## Colorectum  :1 Median :1672934
## Liver       :1 Mean   :2411599
## Lung        :1 3rd Qu.:2220433
## Other cancers:1 Max.    :8879843
## (Other)     :2

describe(data)

##      vers n      mean      sd      median trimmed      mad      min
## Cancer*   1 8        4.5      2.45         4.5         4.5      2.97      1
## Cases     2 8 2411598.6 2683588.65 1672924.5 2411598.6 869037.85 604127
##          max      range skew kurtosis      se
## Cancer*   8      8275716 -1.65      0.87      0.87
## Cases     8 8879843 8275716 1.67      1.27 948791.87
```

Boxplot

Frequency plot for all the cancer cases being witnessed and as can be seen, breast cancer occupies the second spot.

```
library(ggplot2) #using ggplot2 to plot the frequency plot for the cancer cases

## Warning: package 'ggplot2' was built under R version 3.6.3

##
## Attaching package: 'ggplot2'

## The following objects are masked from 'package:psych':
##
##      %%, alpha

p <- ggplot(data, aes(x = Cancer, y = Cases, fill = Cases), main = "Frequency plot for the Cancer Cases", xlab = "Frequency", ylab = "Cancer") +
  geom_bar(stat="identity") + theme_minimal()
#ggplot(data = data, aes(x = Cancer, y = Cases, fill = Cancer)) +
  geom_bar(stat="identity") + guides(fill=FALSE)
p
```

