



SRI RAMACHANDRA
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SRI RAMACHANDRA ENGINEERING AND TECHNOLOGY

CSE-320 DATA MINING

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COMPUTER SCIENCE AND ENGINEERING

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INTRODUCTION:

Heart Dataset is from Kaggle and has various factors and measures in heart. It has 303 rows and 14 col.

This report provides an analysis and evaluation of the factors that causes heart disease and how much prone it is if some of the factors is misleading and improper. So 14 different factors with proper information in dataset will help us to derive different types of conclusions.

AIM:

The goal of the project is to predict whether a person is prone to heart disease and why he or she having heart disease

PROBLEM STATEMENT:

The dataset had 14 variables which are listed in the later part of the report. Using R programming and powerful libraries like "tidyverse" is used to do proper analysis of dataset and derive proper conclusion.

ATTRIBUTES IN DATASET :

1. **AGE** : Age of the person
2. **SEX** : Gender of the person took test
3. **CP** : Chest pain type (0 – 4)
4. **TRESTPBS** : Resting blood pressure (blood pressure at resting position)
5. **CHOL** : Cholesterol Level
6. **FBS** : Fasting Blood Sugar (Sugar level after fasting) $120 > fbs - 0$ $120 < fbs - 1$
7. **RESTECG** : Resting ElectroCardiographic result (0,1,2)
8. **THALACH** : Maximum heart rate achieved (Beats per minute)
9. **EXANG** : Exercise induced angina (exercise induced chest pain)
10. **OLDPEAK** : A measure of abnormality in electrocardiograms.
11. **SLOPE** : Quality of blood flow to heart
12. **CA** : Cardiography Results
13. **THAL** : Thallium stress test measuring blood flow to heart.
14. **TARGET** : Having heart disease or not (0 – negative 1 – positive)

```
In [ ]: # Heart disease analysis and visualization
```

```
In [1]: data = read.csv("heart.csv")
```

```
In [3]: head(data)
```

age	sex	cp	trestbps	chol	fb	restecg	thalach	exang	oldpeak	slope	ca	thal	target
63	1	3	145	233	1	0	150	0	2.3	0	0	1	1
37	1	2	130	250	0	1	187	0	3.5	0	0	2	1
41	0	1	130	204	0	0	172	0	1.4	2	0	2	1
56	1	1	120	236	0	1	178	0	0.8	2	0	2	1
57	0	0	120	354	0	1	163	1	0.6	2	0	2	1
57	1	0	140	192	0	1	148	0	0.4	1	0	1	1

This is head of the dataset and values and attributes will be in this format.

```
In [6]: library(tidyverse)
```

```
Registered S3 methods overwritten by 'ggplot2':
```

```
method      from  
[.quosures  rlang  
c.quosures  rlang  
print.quosures rlang
```

```
Registered S3 method overwritten by 'rvest':
```

```
method      from  
read_xml.response xml2
```

```
-- Attaching packages ----- tidyverse 1.2.1 --
```

```
-
```

```
v ggplot2 3.1.1      v purrr   0.3.2  
v tibble  2.1.1      v dplyr   0.8.0.1  
v tidyr   0.8.3      v stringr 1.4.0  
v readr   1.3.1      v forcats 0.4.0
```

```
-- Conflicts ----- tidyverse_conflicts() --
```

```
-
```

```
x dplyr::filter() masks stats::filter()
```

```
x dplyr::lag()    masks stats::lag()
```

Installing tidyverse library for enhanced analysis .

```
In [7]: glimpse(data)
```

```
Observations: 303
Variables: 14
$ i..age <int> 63, 37, 41, 56, 57, 57, 56, 44, 52, 57, 54, 48, 49, 64, 58...
$ sex <int> 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 1, 0, 1, 1, 0, 0, 0, 0, 1, 0...
$ cp <int> 3, 2, 1, 1, 0, 0, 1, 1, 2, 2, 0, 2, 1, 3, 3, 2, 2, 3, 0, 3...
$ trestbps <int> 145, 130, 130, 120, 120, 140, 140, 120, 172, 150, 140, 130...
$ chol <int> 233, 250, 204, 236, 354, 192, 294, 263, 199, 168, 239, 275...
$ fbs <int> 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0...
$ restecg <int> 0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 1, 1, 1, 1, 1...
$ thalach <int> 150, 187, 172, 178, 163, 148, 153, 173, 162, 174, 160, 139...
$ exang <int> 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0...
$ oldpeak <dbl> 2.3, 3.5, 1.4, 0.8, 0.6, 0.4, 1.3, 0.0, 0.5, 1.6, 1.2, 0.2...
$ slope <int> 0, 0, 2, 2, 2, 1, 1, 2, 2, 2, 2, 2, 2, 1, 2, 1, 2, 0, 2, 2...
$ ca <int> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2...
$ thal <int> 1, 2, 2, 2, 2, 1, 2, 3, 3, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2...
$ target <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1...
```

```
In [7]: ncol(data)
```

```
14
```

```
In [8]: nrow(data)
```

```
303
```

```
In [9]: colnames(data)
```

```
'i..age' 'sex' 'cp' 'trestbps' 'chol' 'fbs' 'restecg' 'thalach' 'exang' 'oldpeak' 'slope'
'ca' 'thal' 'target'
```

Basic glimpse of the dataset and some info on number of rows and columns.

```
In [8]: summary(data)
```

i.age	sex	cp	trestbps
Min. :29.00	Min. :0.0000	Min. :0.000	Min. : 94.0
1st Qu.:47.50	1st Qu.:0.0000	1st Qu.:0.000	1st Qu.:120.0
Median :55.00	Median :1.0000	Median :1.000	Median :130.0
Mean :54.37	Mean :0.6832	Mean :0.967	Mean :131.6
3rd Qu.:61.00	3rd Qu.:1.0000	3rd Qu.:2.000	3rd Qu.:140.0
Max. :77.00	Max. :1.0000	Max. :3.000	Max. :200.0

chol	fbs	restecg	thalach
Min. :126.0	Min. :0.0000	Min. :0.0000	Min. : 71.0
1st Qu.:211.0	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:133.5
Median :240.0	Median :0.0000	Median :1.0000	Median :153.0
Mean :246.3	Mean :0.1485	Mean :0.5281	Mean :149.6
3rd Qu.:274.5	3rd Qu.:0.0000	3rd Qu.:1.0000	3rd Qu.:166.0
Max. :564.0	Max. :1.0000	Max. :2.0000	Max. :202.0

exang	oldpeak	slope	ca
Min. :0.0000	Min. :0.00	Min. :0.000	Min. :0.0000
1st Qu.:0.0000	1st Qu.:0.00	1st Qu.:1.000	1st Qu.:0.0000
Median :0.0000	Median :0.80	Median :1.000	Median :0.0000
Mean :0.3267	Mean :1.04	Mean :1.399	Mean :0.7294
3rd Qu.:1.0000	3rd Qu.:1.60	3rd Qu.:2.000	3rd Qu.:1.0000
Max. :1.0000	Max. :6.20	Max. :2.000	Max. :4.0000

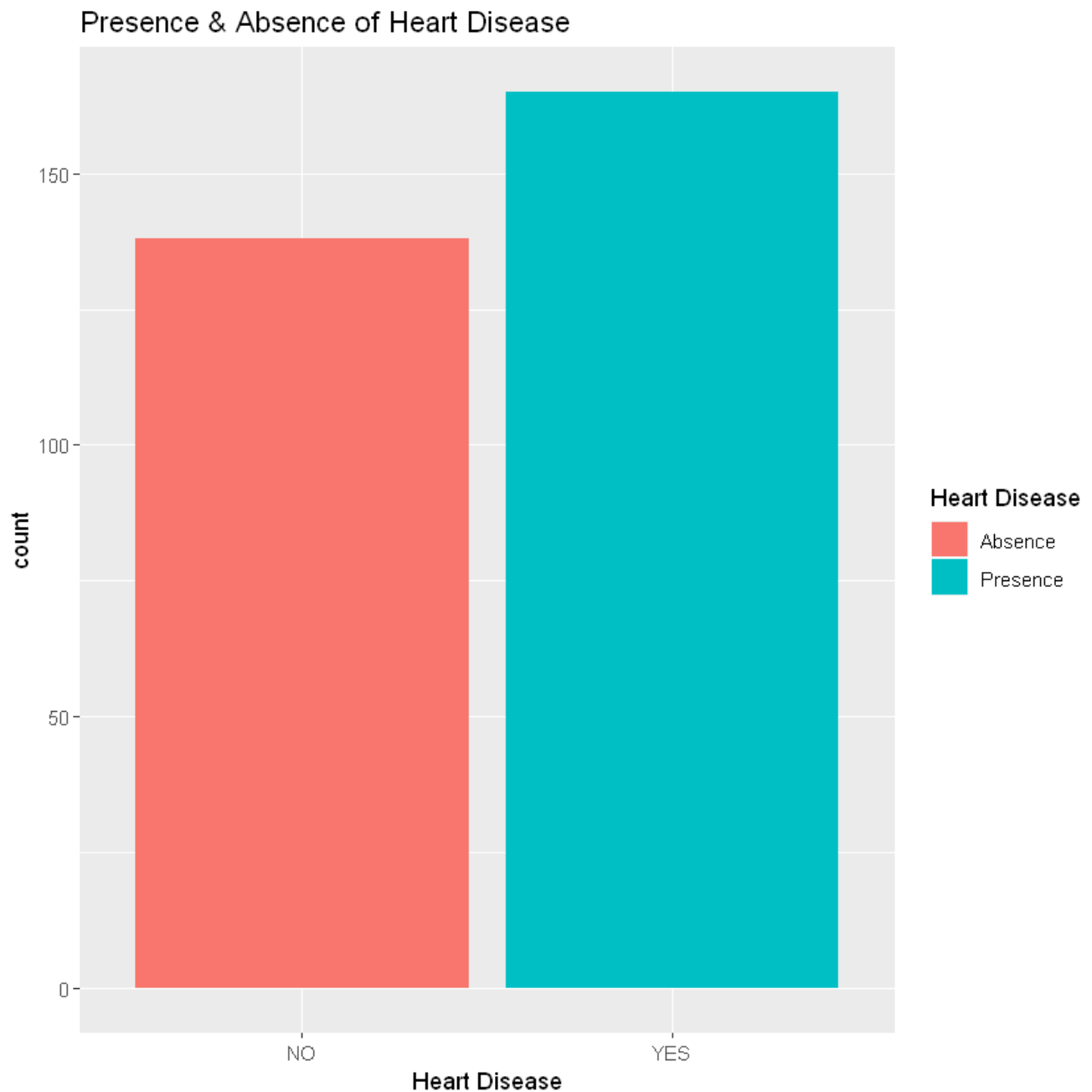
thal	target
Min. :0.000	Min. :0.0000
1st Qu.:2.000	1st Qu.:0.0000
Median :2.000	Median :1.0000
Mean :2.314	Mean :0.5446
3rd Qu.:3.000	3rd Qu.:1.0000
Max. :3.000	Max. :1.0000

Summary of the dataset and its attributes with mean and median and much more variables describing the dataset.

```
In [9]: #Data transformation
data2 <- data %>%
  mutate(sex = if_else(sex == 1, "MALE", "FEMALE"),
         fbs = if_else(fbs == 1, ">120", "<=120"),
         exang = if_else(exang == 1, "YES", "NO"),
         cp = if_else(cp == 1, "ATYPICAL ANGINA",
                      if_else(cp == 2, "NON-ANGINAL PAIN", "ASYMPTOMATIC")),
         restecg = if_else(restecg == 0, "NORMAL",
                           if_else(restecg == 1, "ABNORMALITY", "PROBABLE OR DEFINITE")),
         slope = as.factor(slope),
         ca = as.factor(ca),
         thal = as.factor(thal),
         target = if_else(target == 1, "YES", "NO")
  ) %>%
  mutate_if(is.character, as.factor) %>%
  dplyr::select(target, sex, fbs, exang, cp, restecg, slope, ca, thal, everything())
```

Data transformation for making user friendly analysis on further graphs .

```
In [10]: #Data Visualization
#Bar plot for Target (heart disease)
ggplot(data2, aes(x=data2$target, fill=data2$target))+
  geom_bar()+
  xlab("Heart Disease")+
  ylab("count")+
  ggtitle("Presence & Absence of Heart Disease")+
  scale_fill_discrete(name= 'Heart Disease', labels =c("Absence", "Presence"))
```



Presence of heart disease in this data set to see whether it is biased or natural and proper dataset.

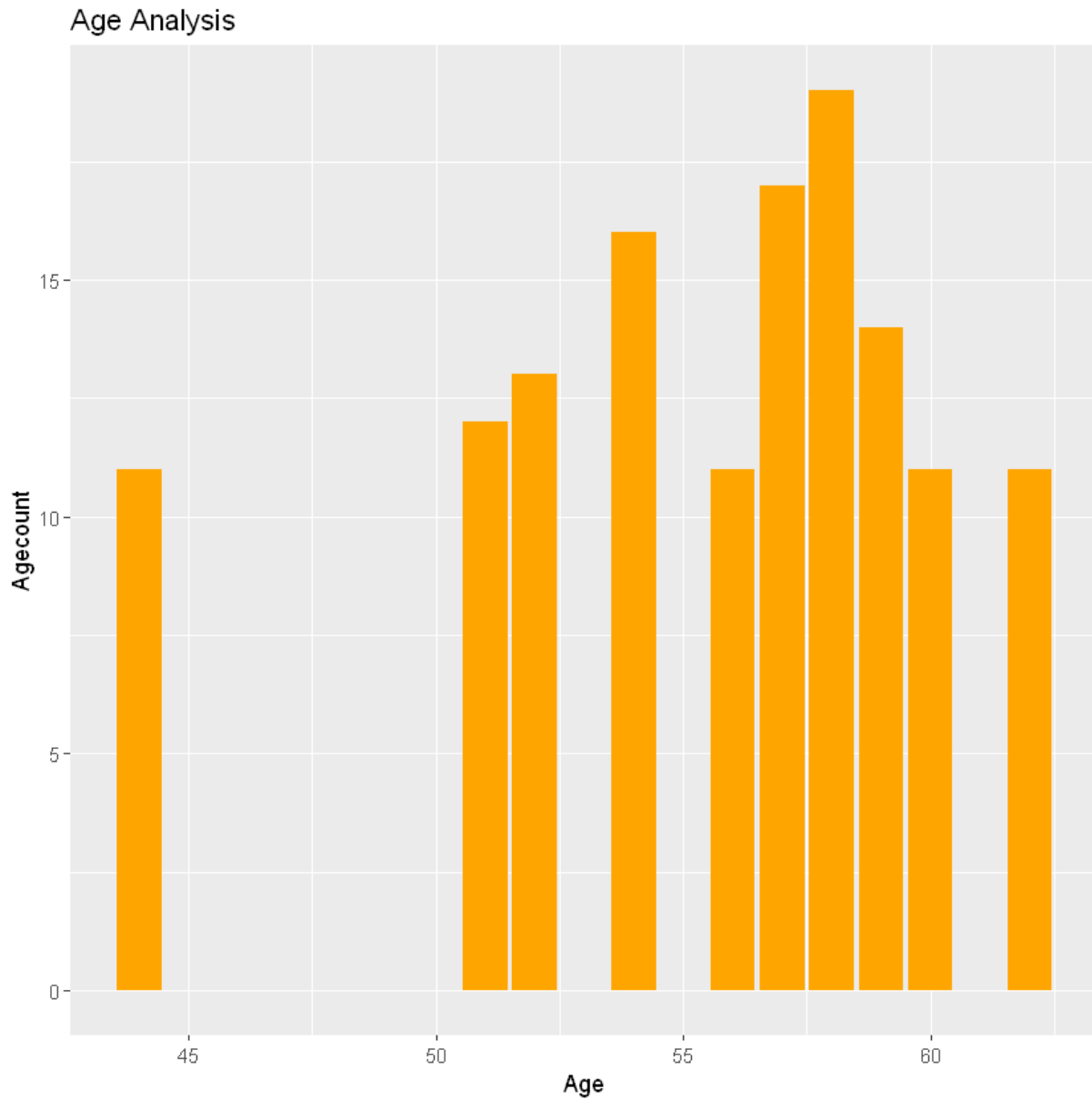
```
In [12]: #proportion
prop.table(table(data2$target))
```

```
      NO      YES
0.4554455 0.5445545
```

This proportion confirms our above statement.

```
In [13]: # count the frequency of the values of age

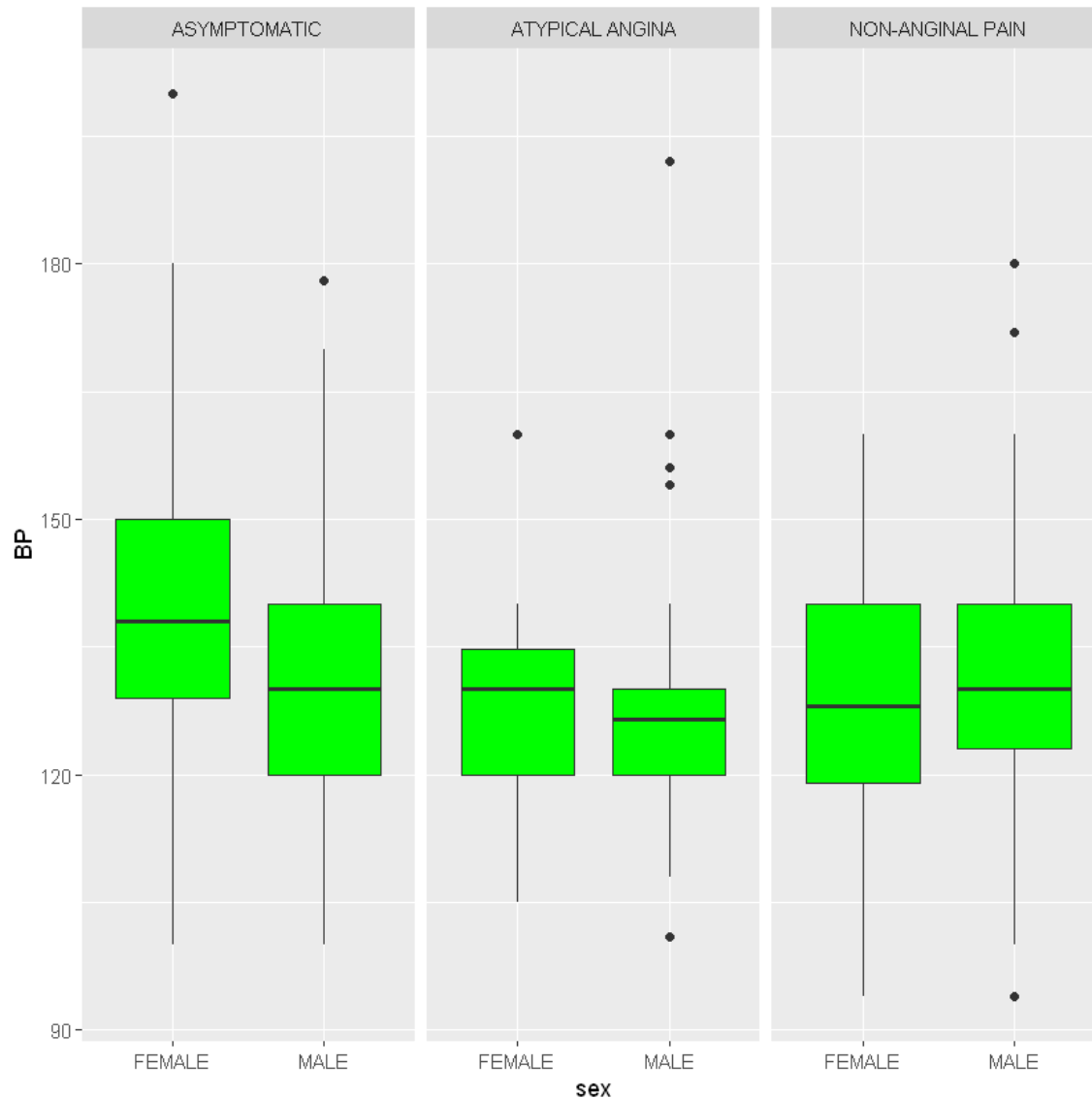
data2 %>%
  group_by(i..age) %>%
  count() %>%
  filter(n>10) %>%
  ggplot()+
  geom_col(aes(i..age, n), fill = 'orange')+
  ggtitle("Age Analysis")+
  xlab("Age")+
  ylab("Agecount")
```



Above analysis shows how age factor affects the heart disease and we can see 45 to 50 has big gap not having any heart disease and below 30 lots of heart disease.

In [20]: # compare blood pressure across the chest pain

```
data2 %>%  
  ggplot(aes(x=sex, y=trestbps))+  
  geom_boxplot(fill='green')+  
  xlab('sex')+  
  ylab('BP')+  
  facet_grid(~cp)
```



Male having many outliers shows higher blood pressure for them and females are having not so higher than males so males are prone to heart disease .

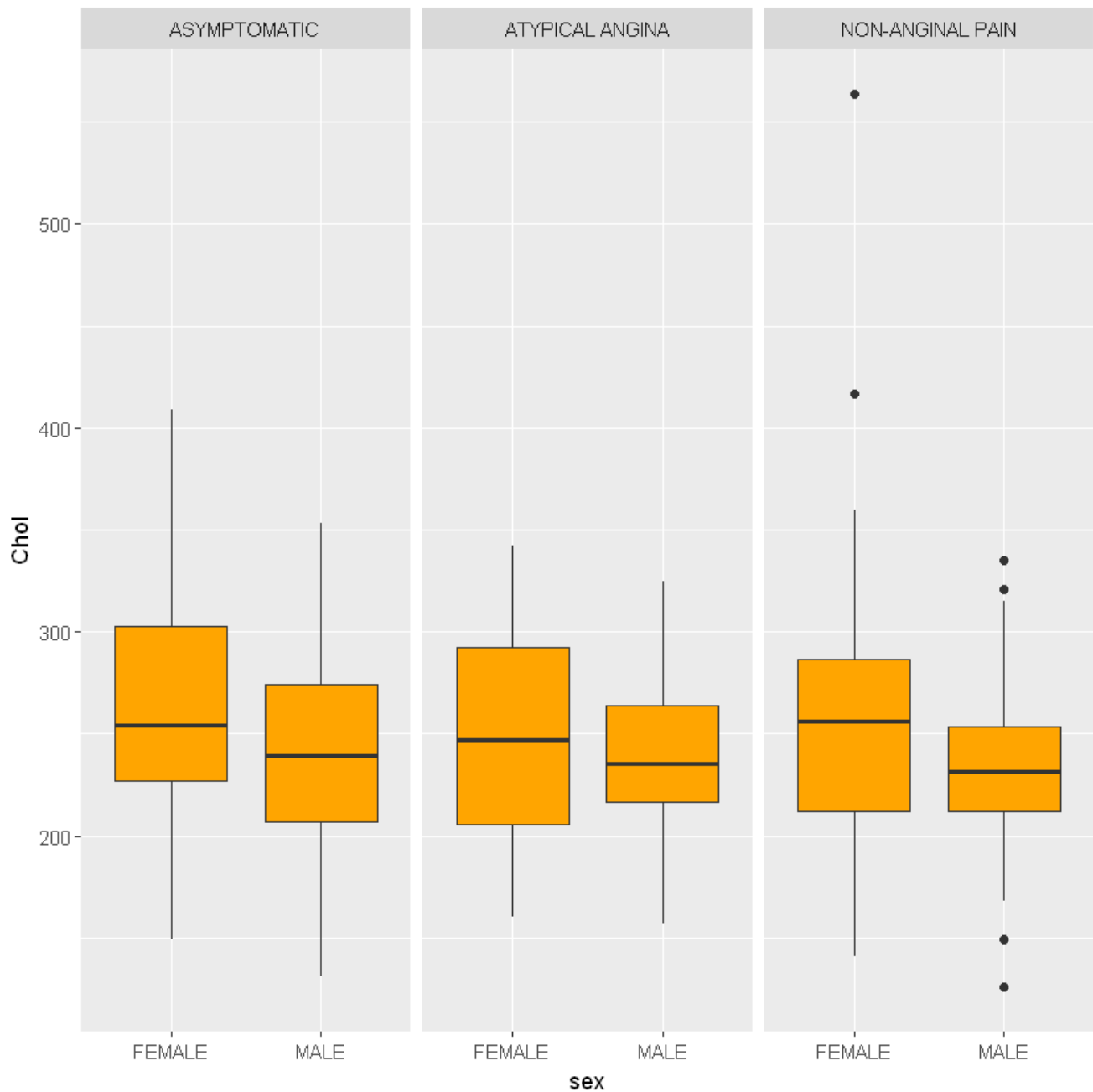
sex

Box plot showing the distribution of blood pressure (BP) across four groups (0, 1, 2, 3) for two sexes (0: female, 1: male). The y-axis represents BP (ranging from 90 to 180). The x-axis represents sex (0: female, 1: male). The plot displays the median, quartiles, and range (whiskers) for each sex within each group, with individual data points overlaid. Red boxes represent the interquartile range, and black lines represent the median. Outliers are shown as black dots.

Group	Sex	Median BP	Q1 BP	Q3 BP	Min BP (Whisker)	Max BP (Whisker)	Outliers BP
0	0 (female)	~130	~120	~140	~100	~170	~175, ~178, ~180, ~185
	1 (male)	~130	~120	~135	~100	~140	~155, ~158, ~160, ~170
1	0 (female)	~130	~120	~140	~100	~160	~175, ~180
	1 (male)	~130	~120	~135	~100	~140	~155, ~158, ~160, ~170
2	0 (female)	~130	~120	~140	~100	~160	~175, ~180
	1 (male)	~130	~120	~140	~100	~140	~155, ~158, ~160, ~170
3	0 (female)	~140	~120	~150	~100	~170	~175, ~180
	1 (male)	~130	~120	~140	~100	~140	~155, ~158, ~160, ~170

Here we are used data which is not transformed data , so from graph we can clearly see that without transforming how difficult is to do analysis and we cant able to derive any conclusions.

```
In [24]: data2 %>%
  ggplot(aes(x=sex, y=chol))+
  geom_boxplot(fill='orange')+
  xlab('sex')+
  ylab('chol')+
  facet_grid(~cp)
```



We can clearly see that female are showing higher spike than males which draws conclusion that they are prone to heart disease because of higher cholesterol. So they have to aware of that.

CORRELATION :

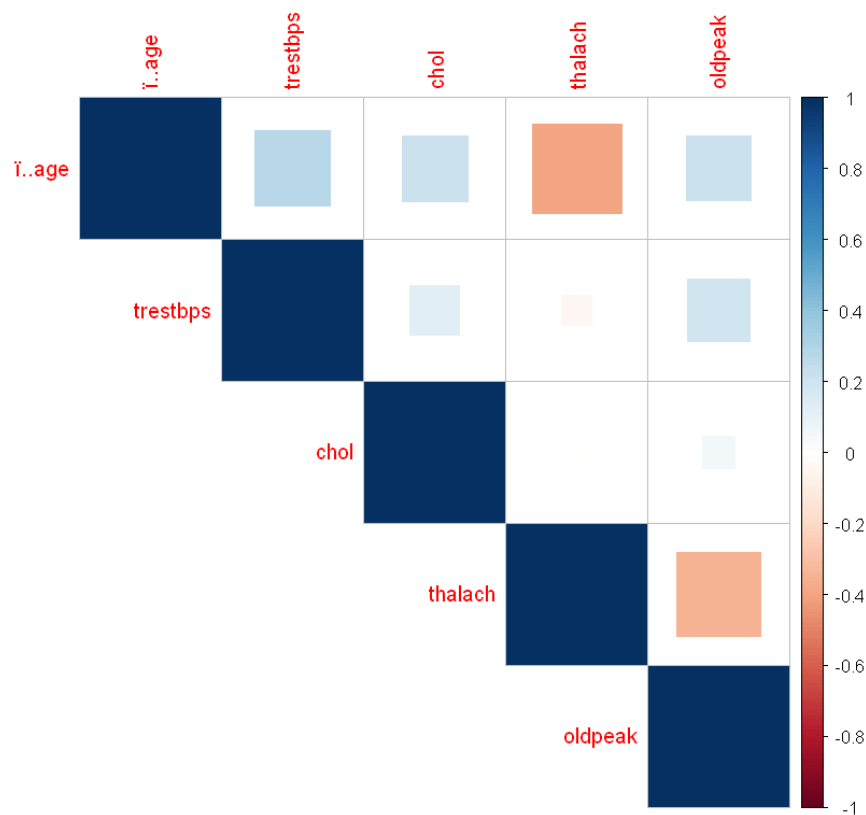
```
In [26]: library(corrplot)
library(ggplot2)

corrplot 0.90 loaded
```

```
In [27]: cor_heart <- cor(data2[, 10:14])
cor_heart

corrplot(cor_heart, method = 'square', type = 'upper')
```

	i..age	trestbps	chol	thalach	oldpeak
i..age	1.0000000	0.27935091	0.213677957	-0.398521938	0.21001257
trestbps	0.2793509	1.00000000	0.123174207	-0.046697728	0.19321647
chol	0.2136780	0.12317421	1.000000000	-0.009939839	0.05395192
thalach	-0.3985219	-0.04669773	-0.009939839	1.000000000	-0.34418695
oldpeak	0.2100126	0.19321647	0.053951920	-0.344186948	1.00000000



As our dataset is small here the correlation not showing any significant relation between any of the attributes , may be in future dataset with more attributes may show relation between the attributes.

CONCLUSION :

From our dataset we have done some analysis on factors which leads to heart disease and how it affects male and female on different factors .

For example males are more prone because of high blood pressure and females are to higher cholesterol levels .

And at end we didn' t see any significant relation between the attributes.

