R Notebook

Install and load helpful library for analysis.

6 0

normal No

```
# Install required library, need not to re-install if you already have
# install.packages("dplyr")
# install.packages("Hmisc")
# install.packages("ggplot2")

# Load the require library
# library(dplyr)
# library(Hmisc)
# library(ggplot2)
```

Data Loading & Exploratory data analysis

Load the data in csv file and store to variable name data.

```
path <- "https://raw.githubusercontent.com/pnhuy/datasets/master/heart_uci/heart.csv"
data <- read.csv(path)</pre>
```

Show some rows of data to get some insight of data.

```
head(data)
                 ChestPain RestBP Chol Fbs RestECG MaxHR ExAng Oldpeak Slope
    X Age Sex
## 1 1 63
                                   233
                                                     150
                                                            0
                                                                  2.3
                                                                          3
           1
                              145
                                         1
                                                 2
                   typical
                                                                          2
## 2 2
       67
            1 asymptomatic
                              160
                                   286
                                                    108
                                                                  1.5
## 3 3 67
           1 asymptomatic
                              120
                                   229
                                        0
                                                 2
                                                    129
                                                                  2.6
                                                                          2
                                                            1
## 4 4
       37
                nonanginal
                              130
                                   250
                                        0
                                                 0
                                                    187
                                                            0
                                                                  3.5
                                                                          3
            1
## 5 5 41
                                                2 172
                nontypical
                              130
                                   204
                                       0
                                                            0
                                                                  1.4
                                                                          1
            0
## 6 6 56
                              120
                                   236
                                                   178
                                                                  0.8
                                                                          1
                nontypical
    Ca
             Thal AHD
##
## 1 0
            fixed No
## 2 3
           normal Yes
## 3 2 reversable Yes
## 4 0
           normal No
## 5 0
           normal No
```

Remove the first columns because it is not useful for analysis.

```
data_col <- colnames(data)
data <- data[data_col[2:length(data_col)]]
head(data)</pre>
```

```
ChestPain RestBP Chol Fbs RestECG MaxHR ExAng Oldpeak Slope
    Age Sex
                           145
## 1 63
          1
                 typical
                                 233
                                      1
                                              2
                                                  150
                                                          0
                                                                2.3
                                                                        3
## 2 67
          1 asymptomatic
                            160
                                 286
                                      0
                                              2
                                                  108
                                                          1
                                                                1.5
                                                                        2
                            120 229
                                              2
                                                  129
                                                                2.6
                                                                        2
## 3 67
          1 asymptomatic
                                      0
                                                          1
## 4 37
                            130
                                 250
                                      0
                                                  187
                                                                3.5
                                                                        3
         1 nonanginal
                                              2
                                 204
                                                  172
## 5 41
         0 nontypical
                            130
                                      0
                                                          0
                                                                1.4
                                                                        1
```

```
## 6 56 1 nontypical 120 236 0 0 178
                                                      0.8
           Thal AHD
##
   Ca
## 1 0
          fixed No
## 2 3
         normal Yes
## 3 2 reversable Yes
## 4 0
         normal No
## 5 0
        normal No
## 6 0
         normal No
```

What was the average age?

```
mean(data$Age)
```

```
## [1] 54.43894
```

From sex column, create new variable gender which only have 'male', 'female'?

```
gender <- factor(data$Sex, levels=c(0,1), labels=c('female', 'male'))</pre>
```

How many percent of patient who was male were there in the data?

```
sum(data$Sex == 1)/length(data$Sex)*100
## [1] 67.9868
```

How many percent of male patient who suffered heart disease were there in the data? What about female?

```
print(sum(data[data$Sex == 1, 'AHD'] == 'Yes')/length(data[data$Sex == 1, 'AHD']))
## [1] 0.5533981
print(sum(data[data$Sex == 0, 'AHD'] == 'Yes')/length(data[data$Sex == 0, 'AHD']))
## [1] 0.257732
```

What was the range of RestBP?

```
print(c(min(data$RestBP), max(data$RestBP)))
## [1] 94 200
```

What was the distribution of AHD?

```
print(table(data$AHD))

##

## No Yes
## 164 139
```

Calculate some basic descriptive statistics

```
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
summary(data)
                                             {\tt ChestPain}
##
         Age
                         Sex
                                                             RestBP
           :29.00
                           :0.0000
                                      asymptomatic:144
                                                         Min.
                                                                : 94.0
##
   Min.
                    Min.
                    1st Qu.:0.0000
##
   1st Qu.:48.00
                                      nonanginal: 86
                                                         1st Qu.:120.0
                                                 : 50
  Median :56.00
                    Median :1.0000
                                      nontypical
                                                         Median :130.0
##
   Mean
           :54.44
                           :0.6799
                                                  : 23
                    Mean
                                      typical
                                                         Mean
                                                                :131.7
##
   3rd Qu.:61.00
                    3rd Qu.:1.0000
                                                         3rd Qu.:140.0
##
   Max.
          :77.00
                    Max.
                           :1.0000
                                                         Max.
                                                                :200.0
##
##
         Chol
                         Fbs
                                         RestECG
                                                           MaxHR
                           :0.0000
                                                             : 71.0
##
   Min.
           :126.0
                    Min.
                                     Min.
                                             :0.0000
                                                       Min.
                                      1st Qu.:0.0000
   1st Qu.:211.0
                    1st Qu.:0.0000
                                                       1st Qu.:133.5
   Median :241.0
                    Median :0.0000
                                     Median :1.0000
                                                       Median :153.0
          :246.7
                                             :0.9901
                                                             :149.6
##
   Mean
                    Mean
                           :0.1485
                                      Mean
                                                       Mean
##
   3rd Qu.:275.0
                    3rd Qu.:0.0000
                                      3rd Qu.:2.0000
                                                       3rd Qu.:166.0
                           :1.0000
                                             :2.0000
##
   Max.
           :564.0
                    Max.
                                     Max.
                                                       Max.
                                                              :202.0
##
##
        ExAng
                        Oldpeak
                                         Slope
                                                           Ca
##
   Min.
           :0.0000
                            :0.00
                                            :1.000
                                                     Min.
                                                            :0.0000
                     Min.
                                    Min.
   1st Qu.:0.0000
                     1st Qu.:0.00
                                     1st Qu.:1.000
                                                     1st Qu.:0.0000
  Median :0.0000
                     Median:0.80
                                    Median :2.000
                                                     Median :0.0000
##
   Mean
          :0.3267
                     Mean
                           :1.04
                                    Mean
                                          :1.601
                                                     Mean
                                                            :0.6722
##
   3rd Qu.:1.0000
                     3rd Qu.:1.60
                                     3rd Qu.:2.000
                                                     3rd Qu.:1.0000
   Max.
           :1.0000
                     Max.
                            :6.20
                                    Max. :3.000
                                                     Max.
                                                            :3.0000
##
                                                     NA's
                                                            :4
            Thal
                      AHD
##
##
                     No :164
   fixed
              : 18
  normal
              :166
                     Yes:139
  reversable:117
##
   NA's
              : 2
##
##
##
library(Hmisc)
## Loading required package: lattice
## Loading required package: survival
## Loading required package: Formula
## Loading required package: ggplot2
```

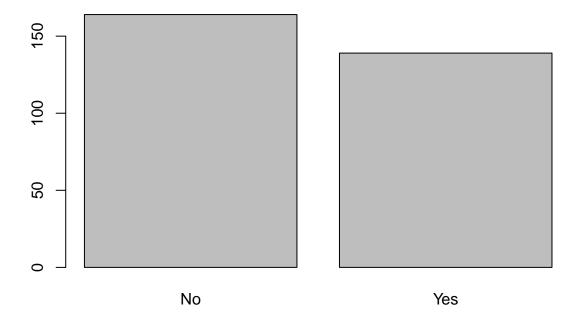
```
##
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:dplyr':
##
    src, summarize
## The following objects are masked from 'package:base':
##
     format.pval, units
describe(data)
## data
##
## 14 Variables 303 Observations
## Age
                                           .05
                                                  .10
     n missing distinct
                       Info Mean
                                     Gmd
                                   10.3
         0 41 0.999 54.44
##
     303
                                            40
                                                   42
               .75 .90
     .25
            .50
                             .95
      48
            56
                         66
##
                  61
## lowest : 29 34 35 37 38, highest: 70 71 74 76 77
     n missing distinct
                       Info Sum
                                    Mean
                                            Gmd
         0 2
                               206 0.6799 0.4367
##
     303
                       0.653
##
## -----
## ChestPain
## n missing distinct
##
     303 0 4
##
## Value asymptomatic nonanginal nontypical typical
           144
## Frequency
                     86
                              50
                                          0.076
## Proportion
             0.475
                       0.284
                                0.165
## RestBP
     n missing distinct Info Mean
                                    Gmd .05
##
                                                  .10
     303 0 50 0.995 131.7 19.41
##
                                            108
                                                   110
                  .75
##
     . 25
            .50
                       .90
                              .95
            130
##
     120
                 140
                         152
                                160
## lowest : 94 100 101 102 104, highest: 174 178 180 192 200
## Chol
##
     n missing distinct Info Mean Gmd .05
                                                  .10
                       1 246.7 55.91 175.1 188.8
     303 0 152
                  .75
                        .90
                               .95
##
     . 25
           .50
    211.0 241.0 275.0 308.8 326.9
##
##
## lowest : 126 131 141 149 157, highest: 394 407 409 417 564
## Fbs
  n missing distinct Info
                               Sum Mean
                                           Gmd
```

```
303 0 2 0.379 45 0.1485 0.2538
##
##
## -----
## RestECG
## n missing distinct Info Mean Gmd
## 303 0 3 0.76 0.9901 1.003
                                 Gmd
##
## Value
         0
              1 2
## Frequency 151 4 148
## Proportion 0.498 0.013 0.488
## -----
## MaxHR
    U 91 1 149.6 25.73 108.1 116.0 .25 .50 .75 .90 .95
  n missing distinct Info Mean
##
##
   133.5 153.0 166.0 176.6 181.9
##
##
## lowest : 71 88 90 95 96, highest: 190 192 194 195 202
## ExAng
  n missing distinct Info Sum Mean Gmd
303 0 2 0.66 99 0.3267 0.4414
##
                                       Gmd
##
## -----
## Oldpeak
  n missing distinct Info Mean
                                 \operatorname{\mathsf{Gmd}}
                                       .05
                                              .10
        0 40 0.964 1.04 1.225 0.0
.50 .75 .90 .95
##
     303
                                              0.0
         .50 .75
0.8 1.6
     . 25
##
                      2.8
     0.0
## lowest : 0.0 0.1 0.2 0.3 0.4, highest: 4.0 4.2 4.4 5.6 6.2
## -----
## n missing distinct Info Mean
     303 0 3 0.798 1.601 0.6291
##
##
## Value
          1 2 3
## Frequency 142 140 21
## Proportion 0.469 0.462 0.069
## n missing distinct Info Mean
     299 4 4 0.783 0.6722 0.9249
##
                  2
          0
              1
## Frequency 176 65 38 20
## Proportion 0.589 0.217 0.127 0.067
## -----
## Thal
## n missing distinct
##
     301 2 3
##
## Value fixed normal reversable
## Frequency 18 166 117
```

```
0.060
                         0.551
## Proportion
                                   0.389
##
## AHD
##
         n missing distinct
##
       303
##
## Value
               No
## Frequency
               164
                     139
## Proportion 0.541 0.459
```

Plot the distribution of AHD?

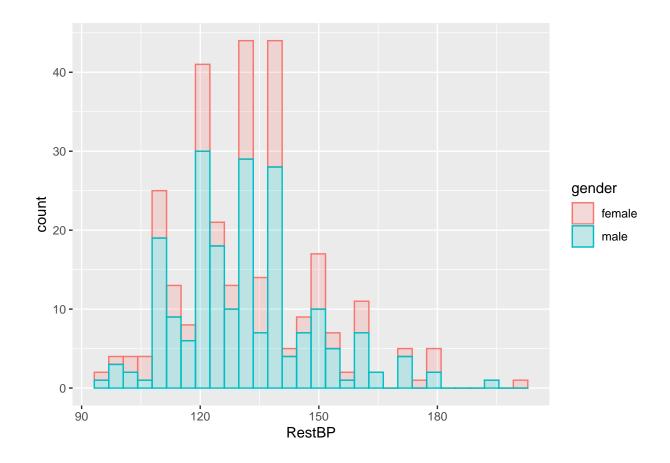
```
barplot(table(data$AHD))
```



Plot the distribution of RestBP per Sex?

```
# In this plot, the distribution need a Factor variable to split, so we input the `gender`
# instead of `Sex`
ggplot(data, aes(x=RestBP)) + geom_histogram(aes(color=gender, fill=gender), alpha=0.2)
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.



Illustrate the relationship between sex and AHD?

```
table(data$Sex, data$AHD)

##

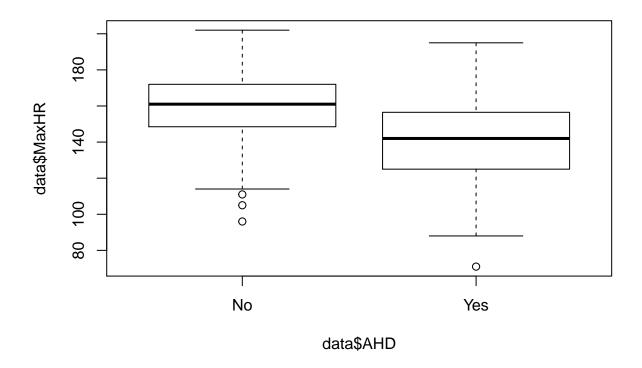
## No Yes

## 0 72 25

## 1 92 114
```

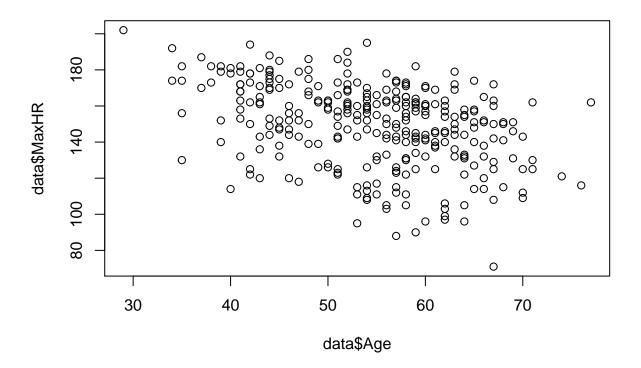
Illustrate the relationship between MaxHR and AHD?

```
boxplot(data$MaxHR ~ data$AHD)
```



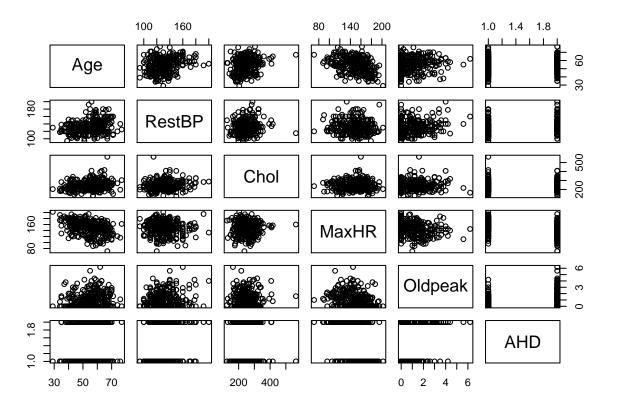
Illustrate the relationship between Age and MaxHR?

plot(data\$Age, data\$MaxHR)



Illustrate the relationship between the continuous variables and the target?

```
pairs(data[c('Age', 'RestBP', 'Chol', 'MaxHR', 'Oldpeak', 'AHD')])
```



Hypothesis testing

Compare the mean RestBP with normal BP (120)?

```
t.test(data$RestBP, mu=120)

##

## One Sample t-test

##

## data: data$RestBP

## t = 11.562, df = 302, p-value < 2.2e-16

## alternative hypothesis: true mean is not equal to 120

## 95 percent confidence interval:

## 129.7001 133.6794

## sample estimates:

## mean of x

## 131.6898</pre>
```

Compare mean RestBP by AHD?

```
RestBP_Yes = data[data$AHD == "Yes", "RestBP"]
RestBP_No = data[data$AHD == "No", "RestBP"]
t.test(RestBP_Yes, RestBP_No)
```

##

```
## Welch Two Sample t-test
##
## data: RestBP_Yes and RestBP_No
## t = 2.6152, df = 274.64, p-value = 0.009409
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 1.314915 9.321775
## sample estimates:
## mean of x mean of y
## 134.5683 129.2500
```

Test the independence between AHD and Sex?

```
chisq.test(data$Sex, data$AHD)

##

## Pearson's Chi-squared test with Yates' continuity correction

##

## data: data$Sex and data$AHD

## X-squared = 22.043, df = 1, p-value = 2.667e-06
```

Compare mean MaxHR by Thal

2 observations deleted due to missingness

ANOVA

```
anova <- aov(MaxHR ~ Thal, data=data)
summary(anova)

## Df Sum Sq Mean Sq F value Pr(>F)

## Thal 2 14396 7198 15.06 5.86e-07 ***

## Residuals 298 142389 478

## ---

## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Pairwise T-Test

```
pairwise.t.test(data$MaxHR, data$Thal)
##
```

```
## Pairwise comparisons using t tests with pooled SD
##
## data: data$MaxHR and data$Thal
##
## fixed normal
## normal 0.00036 -
## reversable 0.13359 1.5e-05
##
## P value adjustment method: holm
```

Tukey multiple pairwise-comparisons

```
TukeyHSD(anova)
```

```
##
     Tukey multiple comparisons of means
       95% family-wise confidence level
##
##
## Fit: aov(formula = MaxHR ~ Thal, data = data)
##
## $Thal
                           diff
                                       lwr
                                                  upr
                                                          p adj
## normal-fixed
                      20.587684
                                  7.810658 33.364710 0.0005225
## reversable-fixed
                       8.324786 -4.711349 21.360922 0.2903956
## reversable-normal -12.262898 -18.478131 -6.047665 0.0000150
```

Linear Regression

Build a model to predict MaxHR by Age?

```
model_1 <- lm(MaxHR ~ Age, data=data)</pre>
summary(model 1)
##
## Call:
## lm(formula = MaxHR ~ Age, data = data)
##
## Residuals:
      Min
                1Q Median
                                3Q
                                       Max
                    3.965 15.937 44.955
## -66.088 -12.040
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 203.8634
                            7.3991 27.553 < 2e-16 ***
                -0.9966
                            0.1341 -7.433 1.11e-12 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 21.06 on 301 degrees of freedom
## Multiple R-squared: 0.1551, Adjusted R-squared: 0.1523
## F-statistic: 55.25 on 1 and 301 DF, p-value: 1.109e-12
```

Build a model to predict MaxHR by Age & RestBP & Thal?

```
model_2 <- lm(MaxHR ~ Age + RestBP + Thal, data=data)</pre>
summary(model_2)
## Call:
## lm(formula = MaxHR ~ Age + RestBP + Thal, data = data)
## Residuals:
##
       Min
                1Q Median
                                 3Q
## -70.046 -12.269
                     3.579 14.086 52.824
##
## Coefficients:
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  172.57393
                              11.48221 15.030 < 2e-16 ***
```

```
## Age
                -0.98287
                            0.13478 -7.292 2.8e-12 ***
## RestBP
                 0.13314
                            0.06932 1.921 0.055720 .
## Thalnormal
                18.34722
                            5.04476 3.637 0.000325 ***
## Thalreversable 7.69466
                            5.11709 1.504 0.133721
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 20.19 on 296 degrees of freedom
    (2 observations deleted due to missingness)
## Multiple R-squared: 0.2301, Adjusted R-squared: 0.2197
## F-statistic: 22.12 on 4 and 296 DF, p-value: 5.425e-16
```