

# HEART

ilke

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```
getwd()
```

```
## [1] "/home/ilke/Documents/github/r"
```

```
setwd("/home/ilke/Downloads")
```

```
df<- read.csv("heart.csv",sep="," , header=TRUE,stringsAsFactors = FALSE)
```

```
head(df)
```

```
##   age sex cp trtbps chol fbs restecg thalach exng oldpeak slp caa thall output
## 1  63  1  3  145 233  1    0   150  0   2.3 0 0    1    1
## 2  37  1  2  130 250  0    1   187  0   3.5 0 0    2    1
## 3  41  0  1  130 204  0    0   172  0   1.4 2 0    2    1
## 4  56  1  1  120 236  0    1   178  0   0.8 2 0    2    1
## 5  57  0  0  120 354  0    1   163  1   0.6 2 0    2    1
## 6  57  1  0  140 192  0    1   148  0   0.4 1 0    1    1
```

exang =egzerzise baęlı engina: (1 = evet; 0 = hayır)

cp : Gs Aęrısı tipi Gs aęrısı tipi Deęer 1: tipik anjina Deęer 2: atipik anjina Deęer 3: anjinal olmayan aęrı Deęer 4: asemptomatik

trtbps : dinlenme kan basıncı (mm Hg olarak)

chol : BMI sensr aracılıęıyla alınan mg/dl cinsinden kolestoral

fbs : (alık kan řekeri > 120 mg/dl) (1 = doęru; 0 = yanlış

rest\_ecg : dinlenme elektrokardiyografik sonuları 0 deęeri: normal Deęer 1: ST-T dalga anormallięine sahip olmak (T dalgası inversiyonları ve/veya ST ykselmesi veya > 0.05 mV depresyonu) Deęer 2: Estes kriterlerine gre olası veya kesin sol ventrikl hipertrofisini gsteriyor

thalach: ulařılan maksimum kalp atıř hızı

target : 0= daha az kalp krizi geirme olasılıęı 1= daha fazla kalp krizi geirme olasılıęı

```
colnames(df)
```

```
## [1] "age"  "sex"  "cp"   "trtbps" "chol" "fbs"
## [7] "restecg" "thalachh" "exng" "oldpeak" "slp" "caa"
## [13] "thall" "output"
```

```
summary(df)
```

```
##      age      sex      cp      trtbps
## 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      thalachh
## 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
##      exng      oldpeak      slp      caa
## 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
##      thall      output
## 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
```

**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
```

**glimpse**(df)

```
## Rows: 303
## Columns: 14
## $ age      <int> 63, 37, 41, 56, 57, 57, 56, 44, 52, 57, 54, 48, 49, 64, 58, 5...
## $ sex      <int> 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 1, 0, 1, 1, 0, 0, 0, 1, 0, 1...
## $ cp       <int> 3, 2, 1, 1, 0, 0, 1, 1, 2, 2, 0, 2, 1, 3, 3, 2, 2, 3, 0, 3, 0...
## $ trtbps   <int> 145, 130, 130, 120, 120, 140, 140, 120, 172, 150, 140, 130, 1...
## $ chol     <int> 233, 250, 204, 236, 354, 192, 294, 263, 199, 168, 239, 275, 2...
## $ fbs      <int> 1, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 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...
## $ thalachh <int> 150, 187, 172, 178, 163, 148, 153, 173, 162, 174, 160, 139, 1...
## $ exng     <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, 0...
## $ slp      <int> 0, 0, 2, 2, 2, 1, 1, 2, 2, 2, 2, 2, 2, 1, 2, 1, 2, 0, 2, 2, 1...
## $ caa      <int> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 0...
## $ thall    <int> 1, 2, 2, 2, 2, 1, 2, 3, 3, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 3...
## $ output   <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1...
```

#kategorik verileri faktor türüne dönüştürme

```
df$sex <- as.factor(df$sex)
df$output <- as.factor(df$output)
df$caa <- as.factor(df$caa)
df$exng <- as.factor(df$exng)
df$thall <- as.factor(df$thall)
df$fbs <- as.factor(df$fbs)
df$restecg <- as.factor(df$restecg)
df$slp <- as.factor(df$slp)
df$cp <- as.factor(df$output)
```

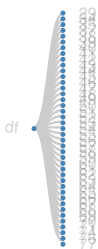
glimpse(df)

```
## Rows: 303
## Columns: 14
## $ age    <int> 63, 37, 41, 56, 57, 57, 56, 44, 52, 57, 54, 48, 49, 64, 58, 5...
## $ sex    <fct> 1, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 1, 1, 0, 0, 0, 0, 1, 0, 1...
## $ cp     <fct> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1...
## $ trtbps <int> 145, 130, 130, 120, 120, 140, 140, 120, 172, 150, 140, 130, 1...
## $ chol   <int> 233, 250, 204, 236, 354, 192, 294, 263, 199, 168, 239, 275, 2...
## $ fbs    <fct> 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0...
## $ restecg <fct> 0, 1, 0, 1, 1, 1, 0, 1, 1, 1, 1, 1, 0, 0, 1, 1, 1, 1, 1...
## $ thalachh <int> 150, 187, 172, 178, 163, 148, 153, 173, 162, 174, 160, 139, 1...
## $ exng    <fct> 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, 0...
## $ slp     <fct> 0, 0, 2, 2, 2, 1, 1, 2, 2, 2, 2, 2, 1, 2, 1, 2, 0, 2, 2, 1...
## $ caa     <fct> 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 0...
## $ thall   <fct> 1, 2, 2, 2, 1, 2, 3, 3, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 3...
## $ output  <fct> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1...
```

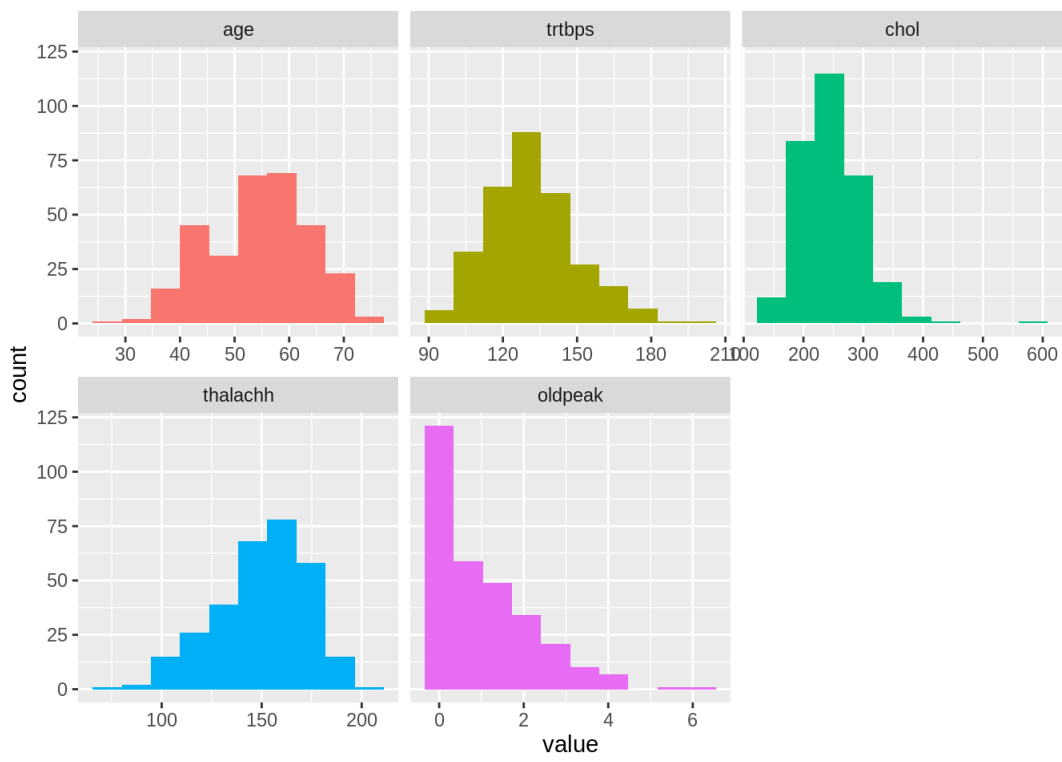
```
df[df==""] <- NA
sum(is.na(df))      #eksik gözlem yok. temiz veri seti
```

## [1] 0

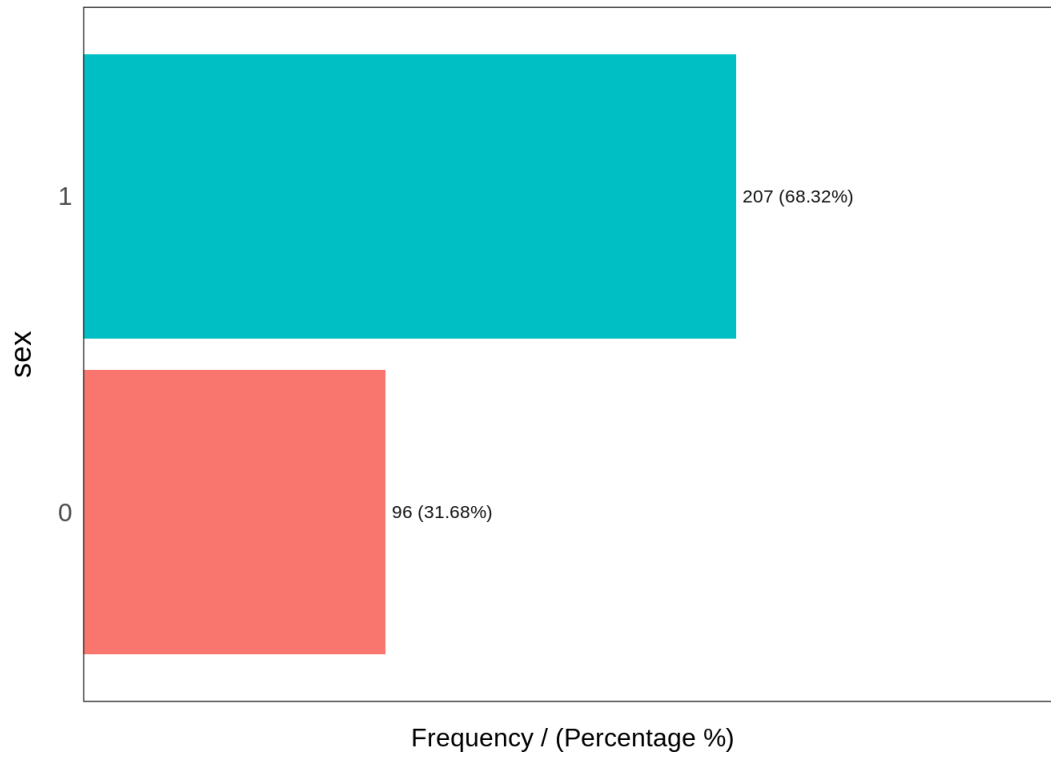
```
library(d3Tree)
d3tree(list(root = df2tree(rootname = 'df',
  struct = as.data.frame(df)),
  layout = 'collapse'))
```

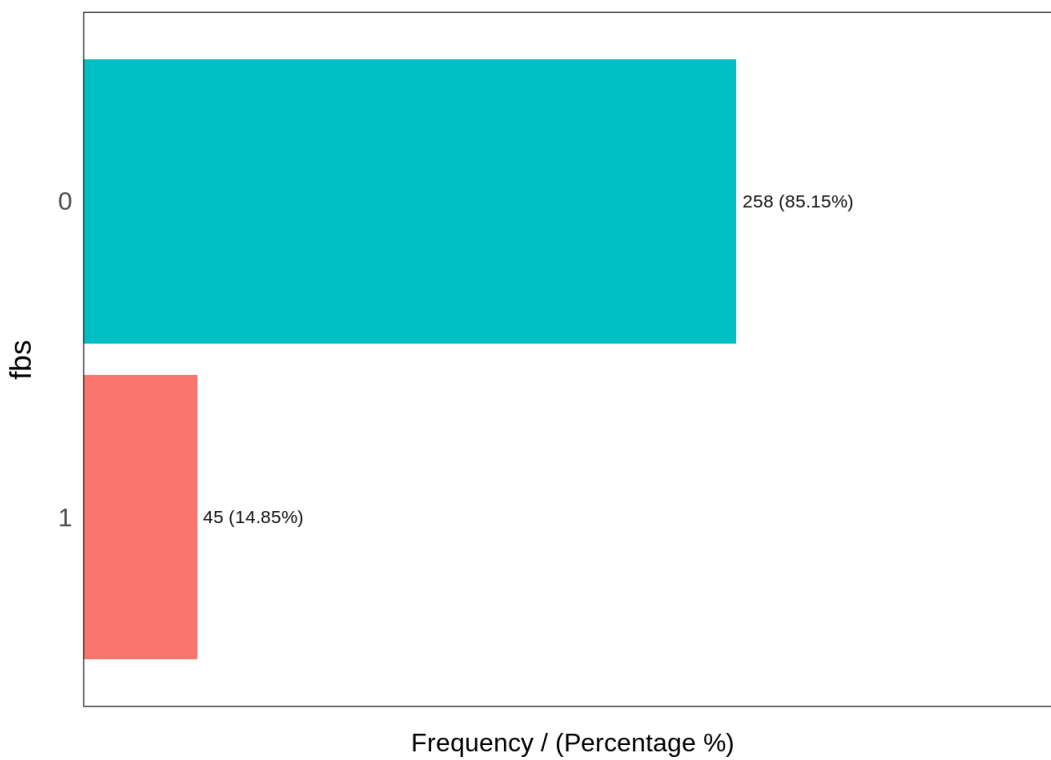
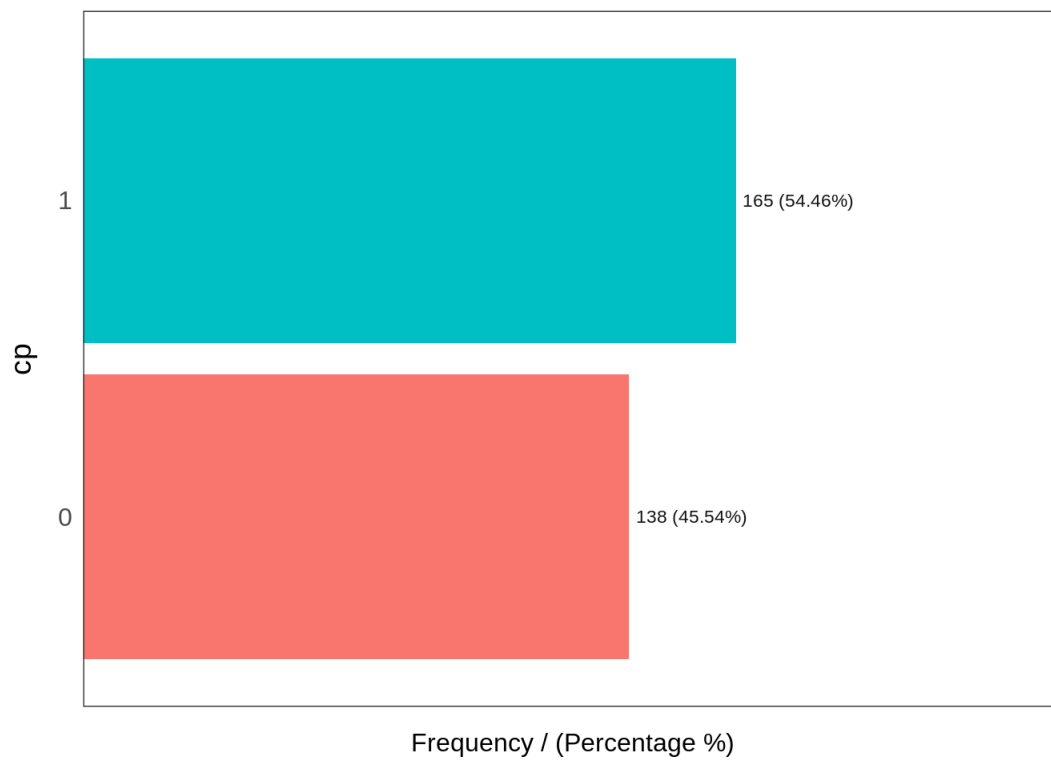


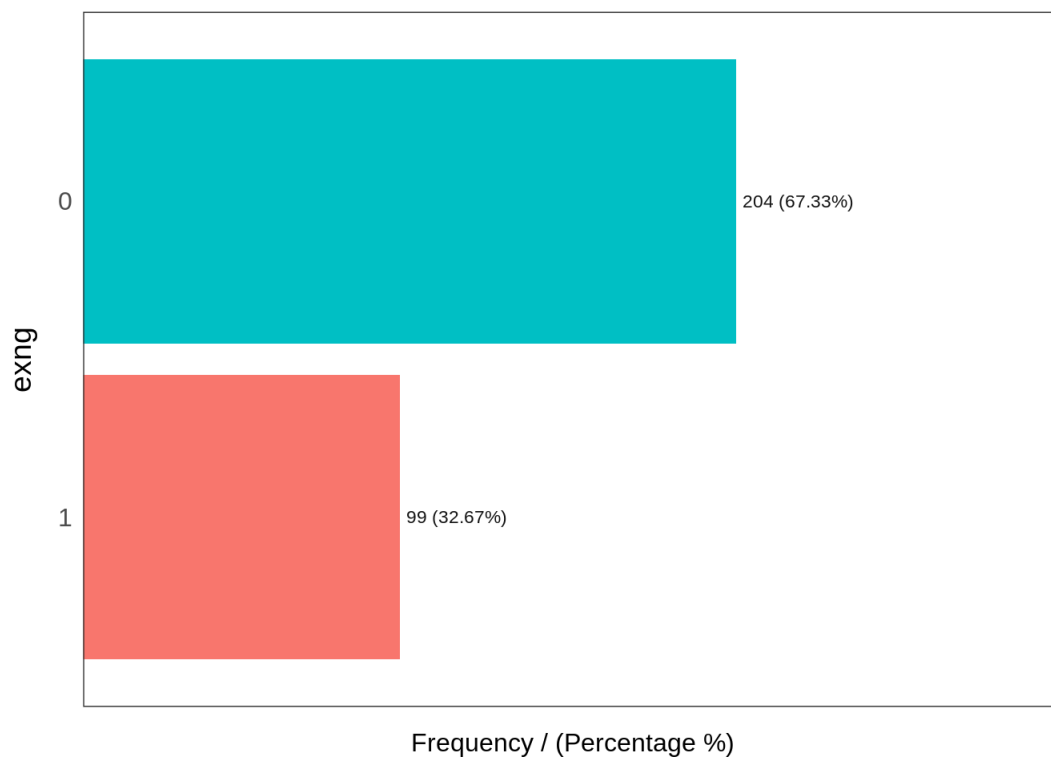
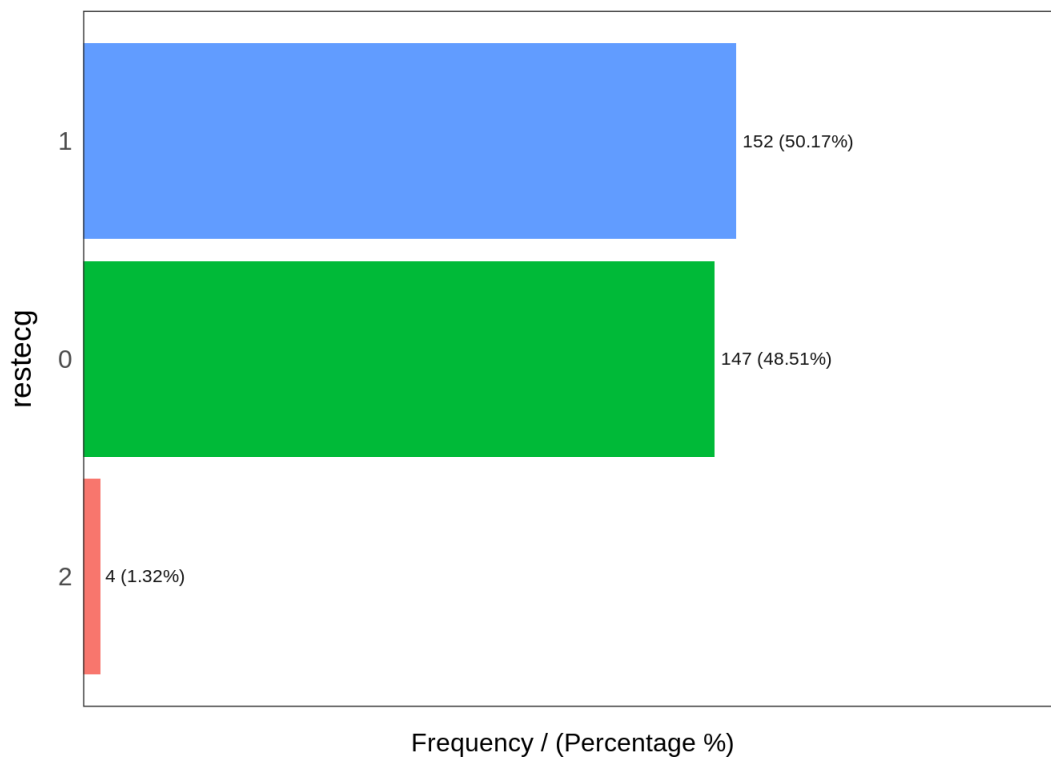
```
library(funModeling)
profiling_num(df) #sürekli değişkenlerin özet bazı istatistiklerini verir
plot_num(df)      #genel histogram
```

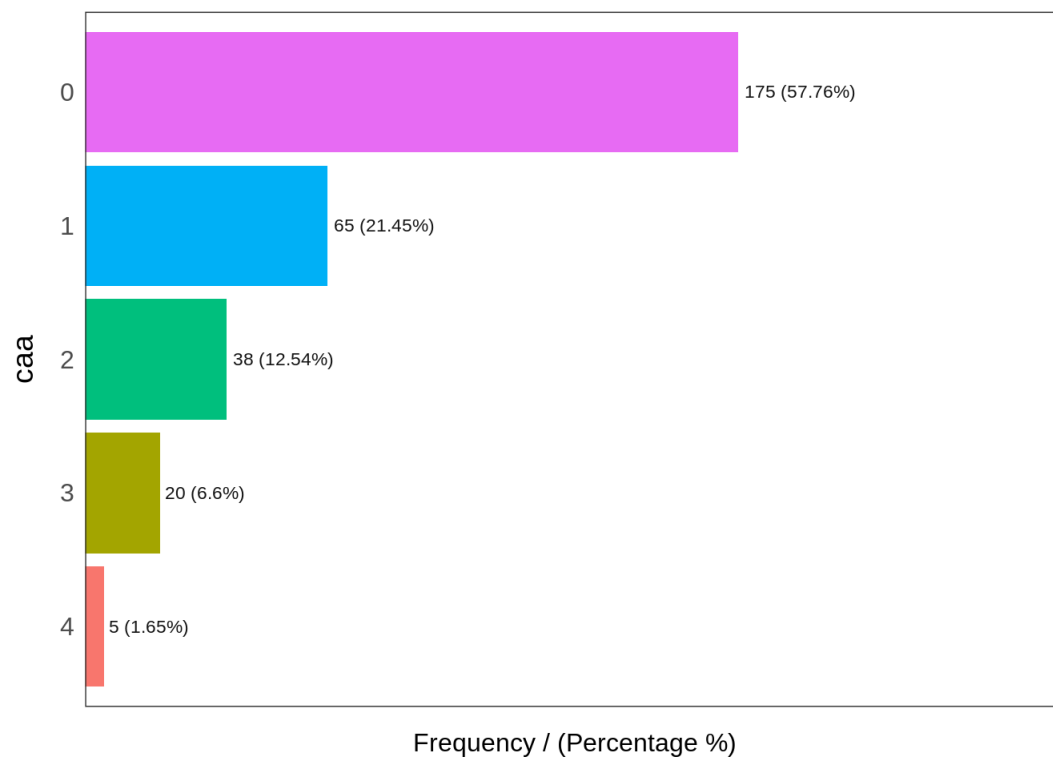
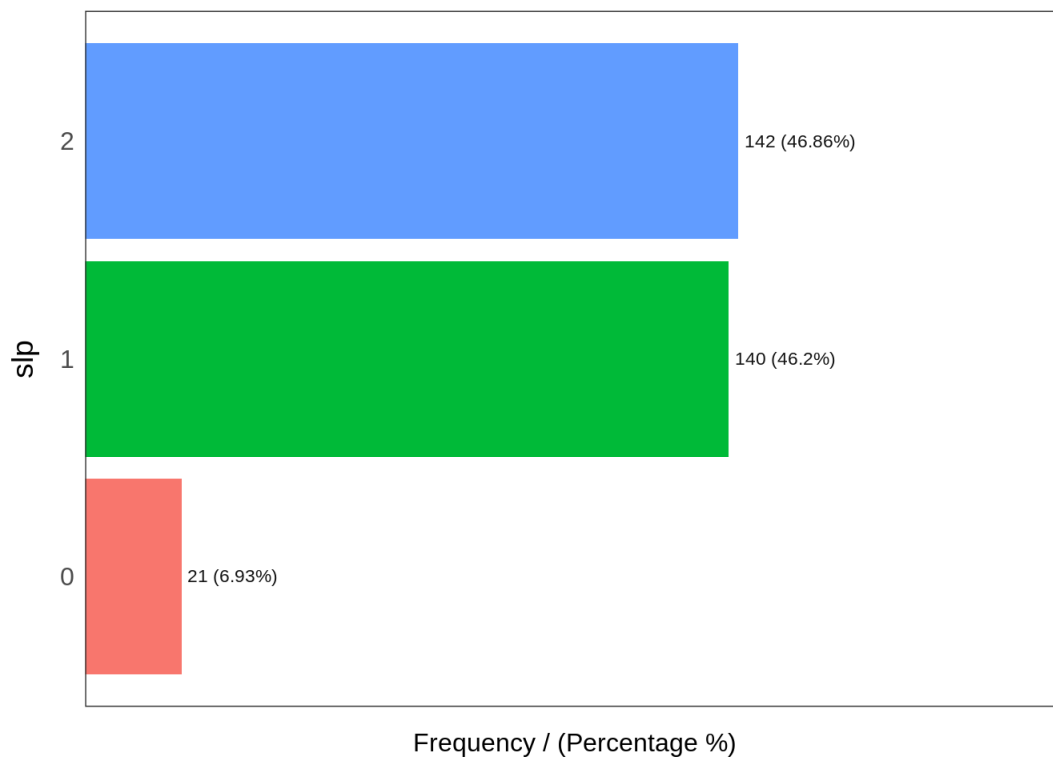


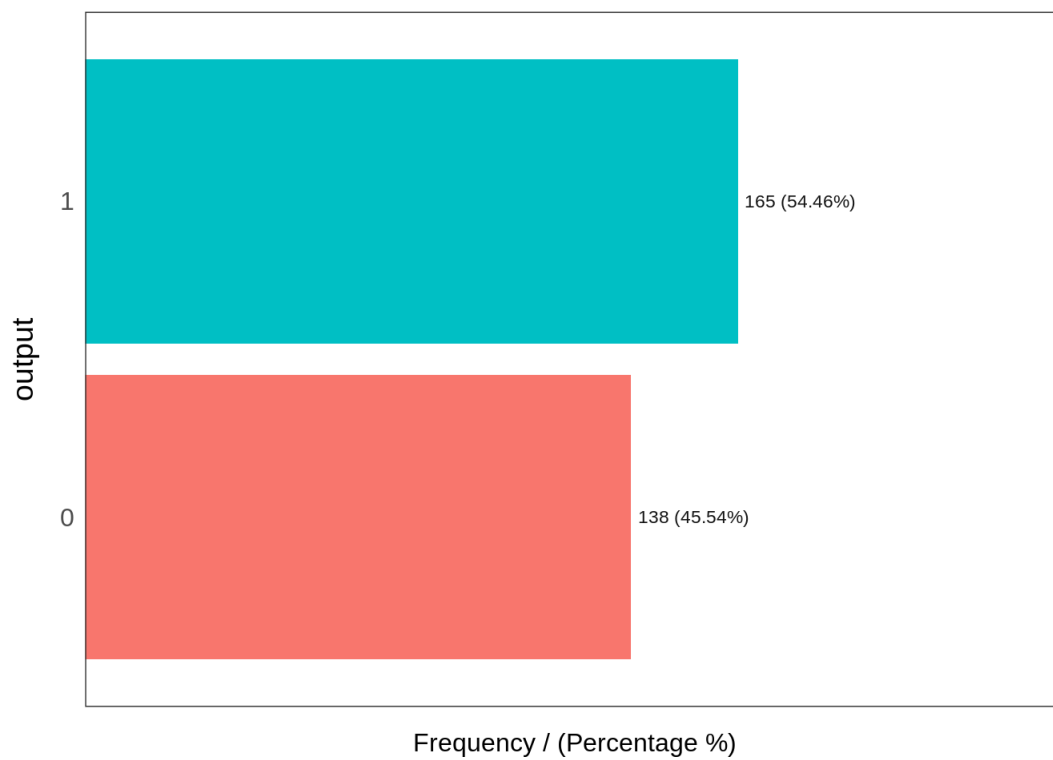
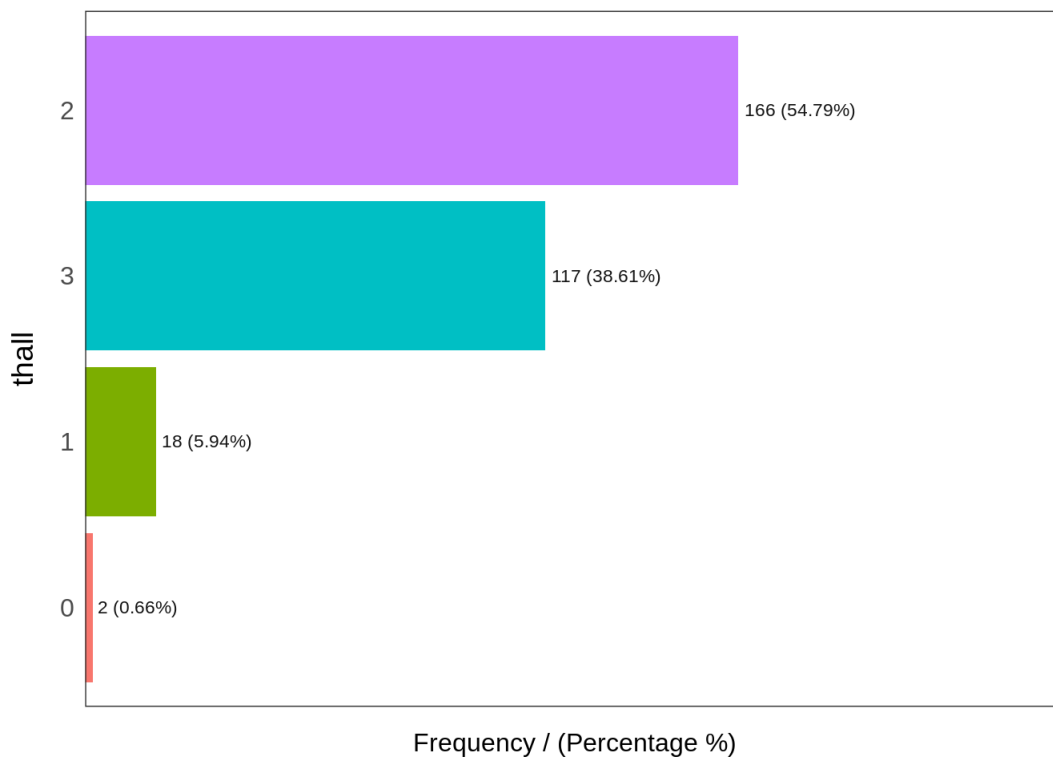
freq(df) #detaylı data frame,grafikler verir











```
library(tidyverse)
```

```
## — Attaching packages — tidyverse 1.3.2 —
## ✔ tibble 3.1.8   ✔ purrr 0.3.4
## ✔ tidyr 1.2.0   ✔ stringr 1.4.0
## ✔ readr 2.1.2   ✔ forcats 0.5.1
## — Conflicts — tidyverse_conflicts() —
## ✖ dplyr::filter() masks stats::filter()
## ✖ dplyr::lag() masks stats::lag()
## ✖ Hmisc::src() masks dplyr::src()
## ✖ Hmisc::summarize() masks dplyr::summarize()
```

```
library(ggplot2)
```

```
library(plotly)
```

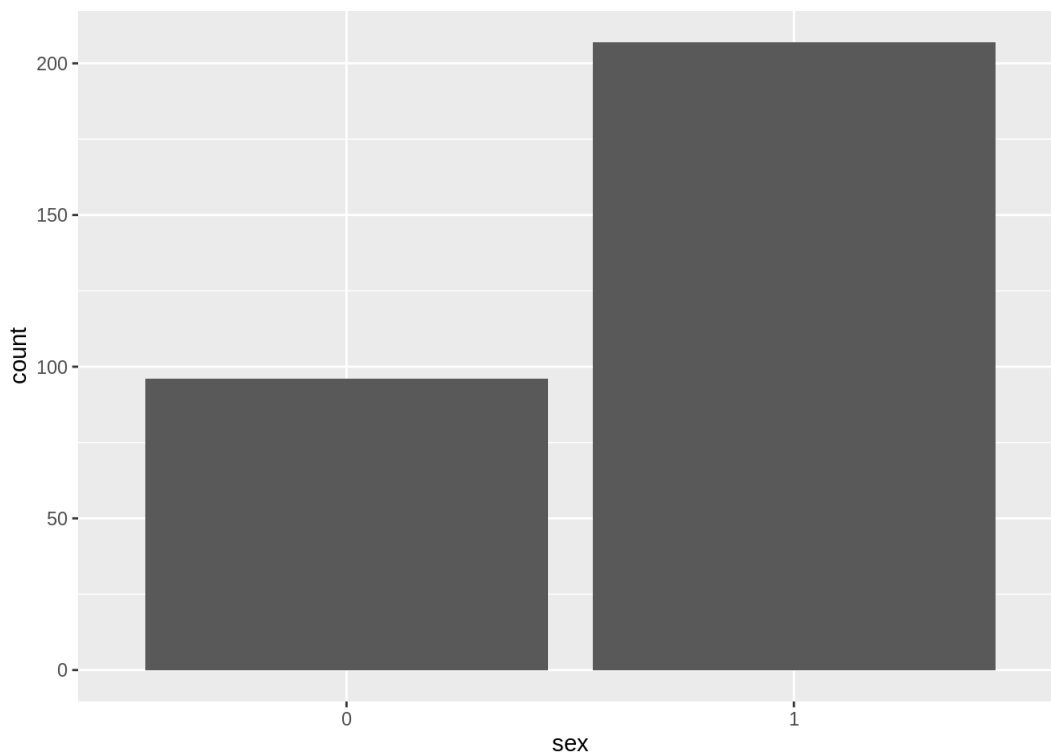


```
##  
## Attaching package: 'plotly'  
##  
## The following object is masked from 'package:Hmisc':  
##  
##   subplot  
##  
## The following object is masked from 'package:ggplot2':  
##  
##   last_plot  
##  
## The following object is masked from 'package:stats':  
##  
##   filter  
##  
## The following object is masked from 'package:graphics':  
##  
##   layout
```

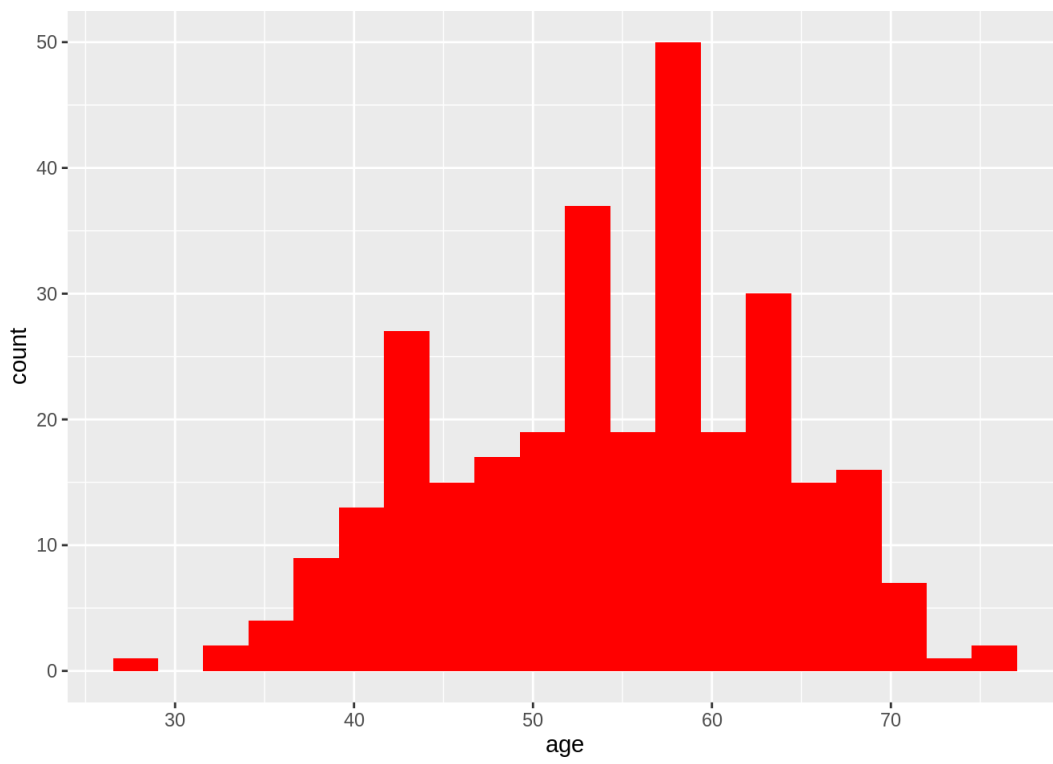
```
cinsiyet<- df %>% group_by(sex) %>% tally() %>% arrange(n, decreasing=T)  
cinsiyet
```

```
## # A tibble: 2 × 2  
##   sex     n  
##   <fct> <int>  
## 1 0      96  
## 2 1     207
```

```
ggplot(data = df) +  
  geom_bar(mapping = aes(x = sex))
```



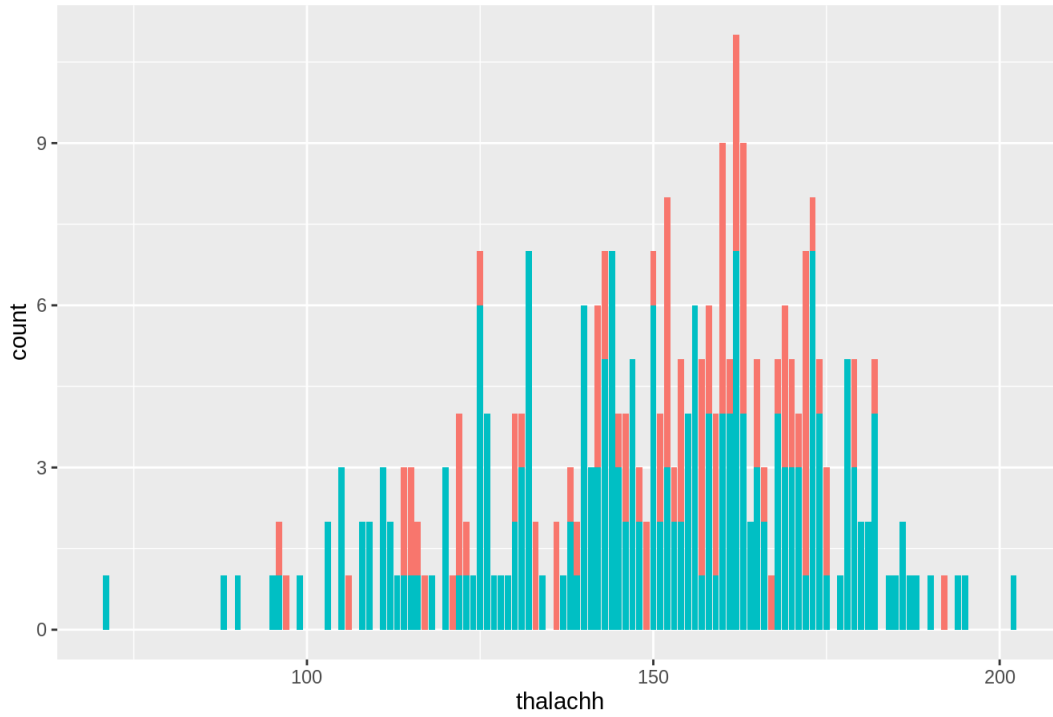
```
ggplot(df ,aes(x=age)) +geom_histogram(bins=20,fill='red')
```



```
#maksimum kalp ve cinsiyet
```

```
ggplot(df, aes(x = thalachh, fill=factor(sex)))+  
  geom_bar() +  
  theme(legend.position = "yaş") +  
  ggtitle("yaş ve kalp krizi geçirme olasılığı")
```

yaş ve kalp krizi geçirme olasılığı



```
library(gridExtra)
```

```
##  
## Attaching package: 'gridExtra'
```

```
## The following object is masked from 'package:dplyr':  
##  
## combine
```

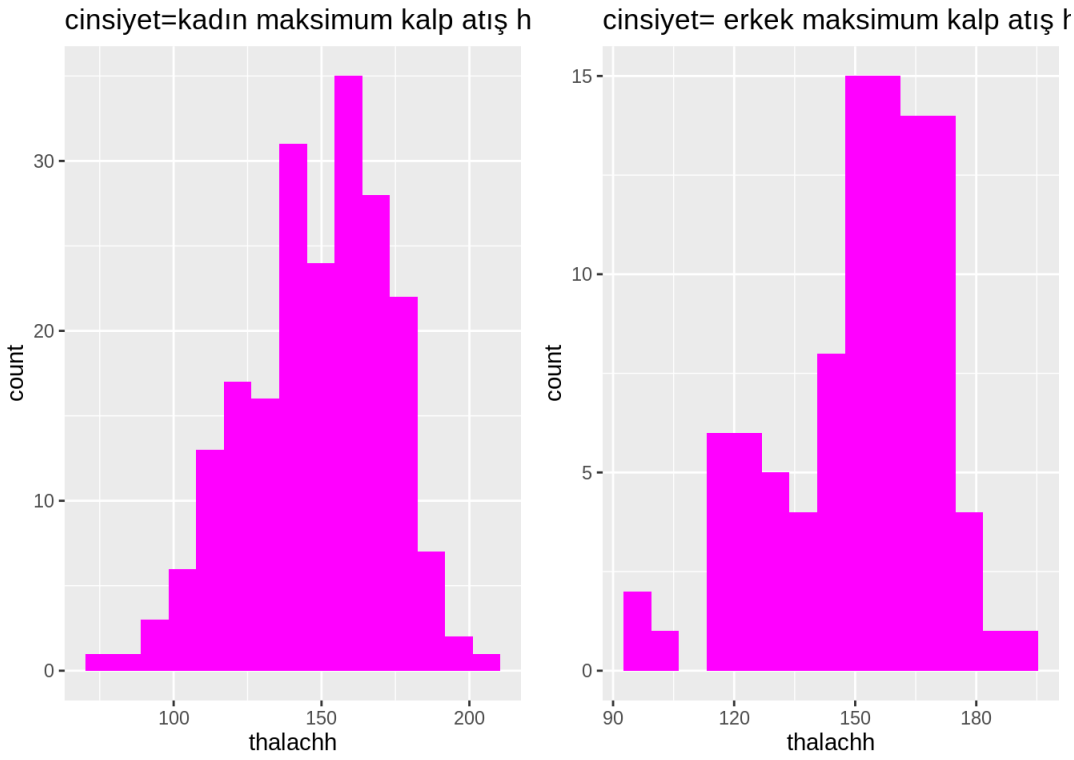
```
df1<- df %>% filter(sex == '1')
```

```
df2<- df %>% filter(sex == '0')
```

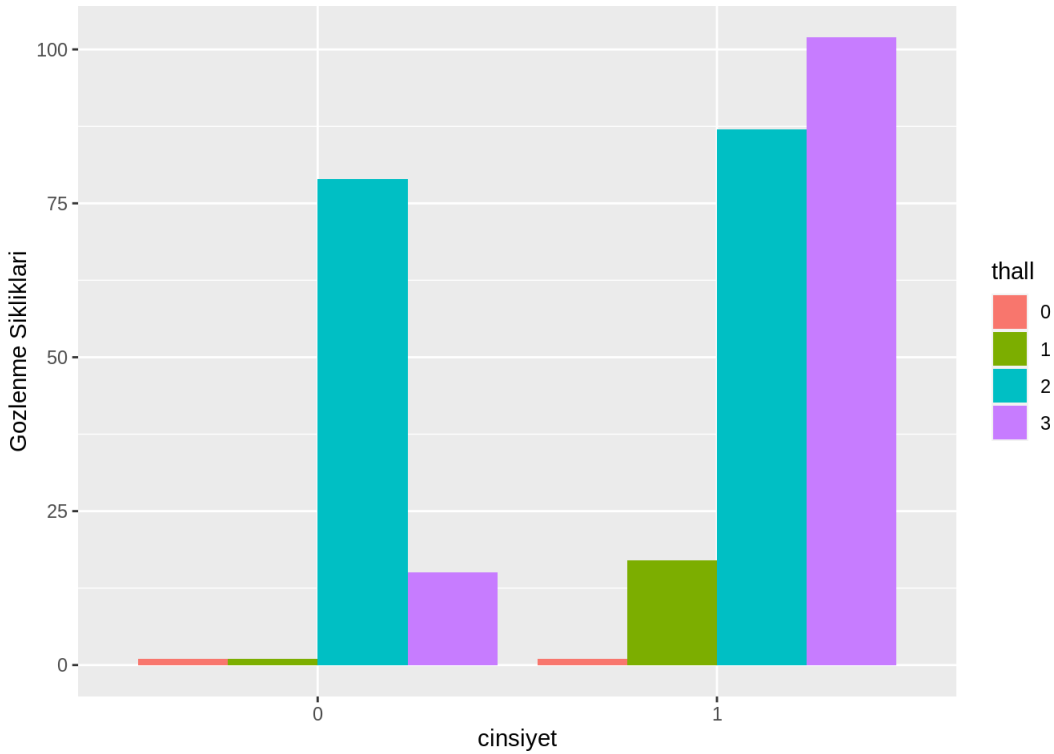
```
df11<- ggplot(df1,aes(x=thalachh))+geom_histogram(bins=15,fill='magenta') + ggtitle("cinsiyet=kadın maksimum kalp atış hızı")
```

```
df22<- ggplot(df2,aes(x=thalachh))+geom_histogram(bins=15,fill='magenta') + ggtitle("cinsiyet= erkek maksimum kalp atış hızı")
```

```
grid.arrange(df11,df22,nrow=1)
```

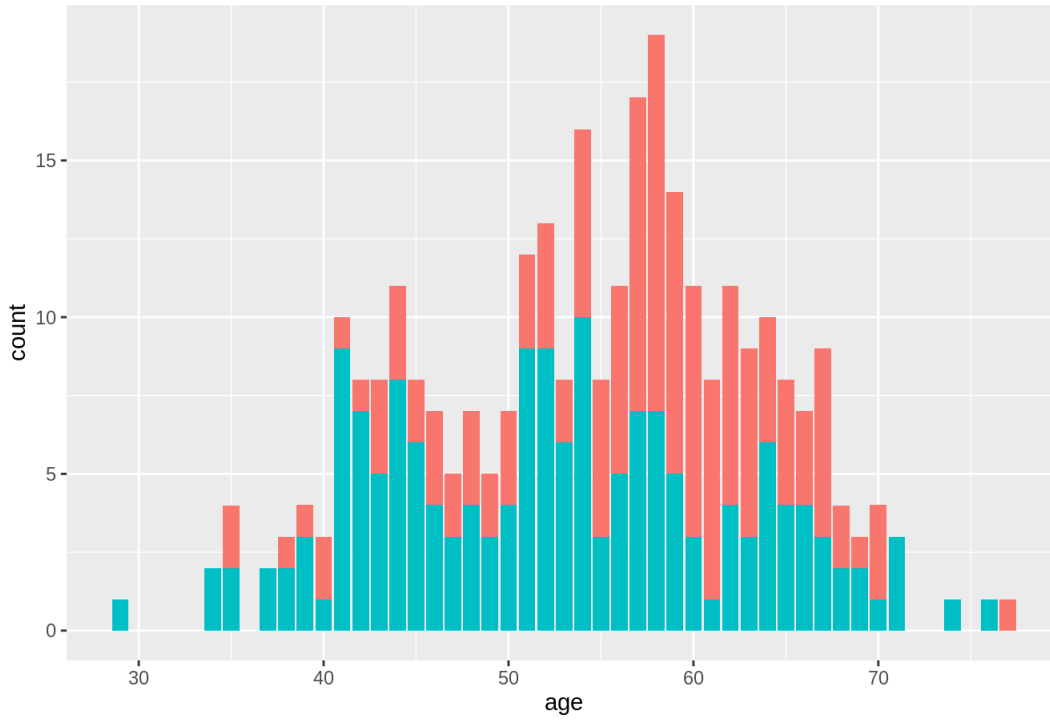


```
ggplot(df, aes(factor(sex), fill = thall)) +  
  geom_bar(position = position_dodge()) +  
  xlab("cinsiyet") +  
  ylab("Gozlenme Sikliklari")
```



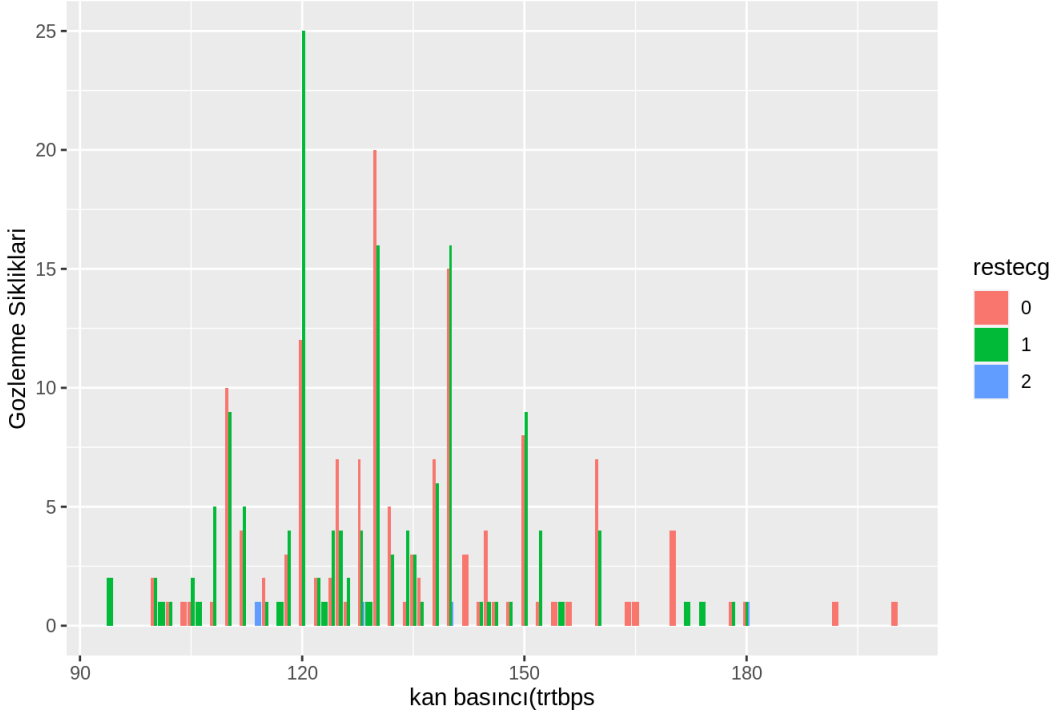
```
ggplot(df, aes(x = age, fill=factor(output)))+  
  geom_bar() +  
  theme(legend.position = "output") +  
  ggtitle("yaş ve kalp krizi geçirme olasılığı")
```

## yaş ve kalp krizi geçirme olasılığı



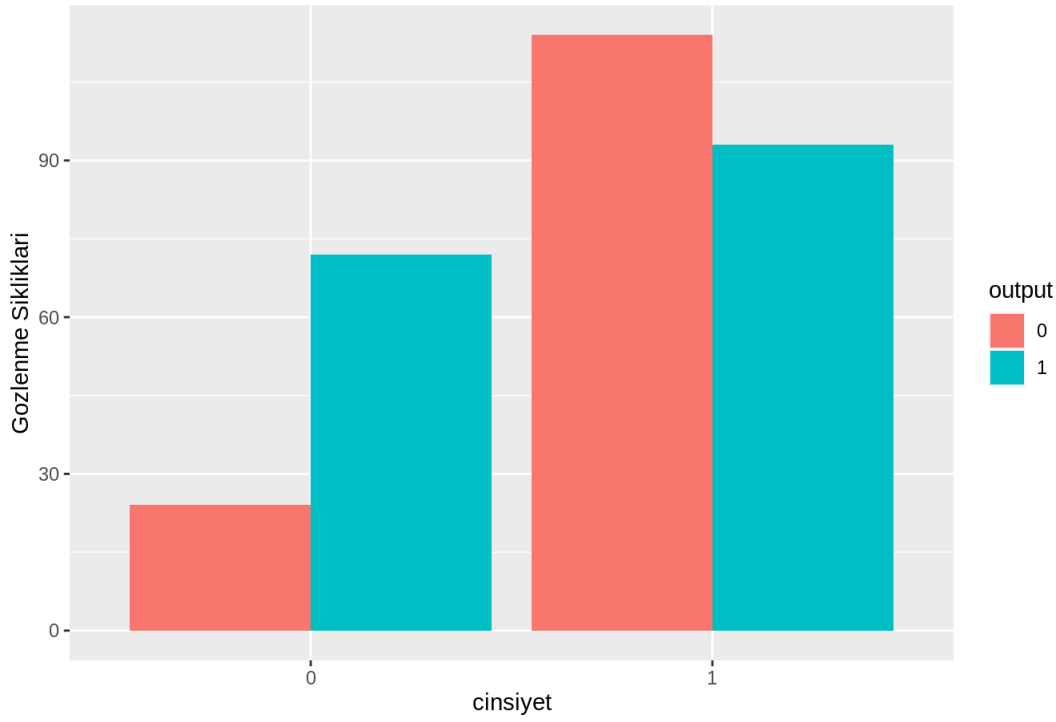
```
ggplot(df, aes(trtbps, fill = restecg)) +  
  geom_bar(position = position_dodge()) +  
  ggtitle("dinlenme kan basıncı ve dinlenme elektrokardiyografik sonuçları") +  
  xlab("kan basıncı(trtbps)") +  
  ylab("Gözlenme Sıklıkları")
```

## dinlenme kan basıncı ve dinlenme elektrokardiyografik sonuçları



```
ggplot(df, aes(factor(sex), fill = output)) +  
  geom_bar(position = position_dodge()) +  
  ggtitle("cinsiyete göre kalp krizi geçirme ") +  
  xlab("cinsiyet") +  
  ylab("Gözlenme Sıklıkları")
```

## cinsiyete göre kalp krizi geçirme



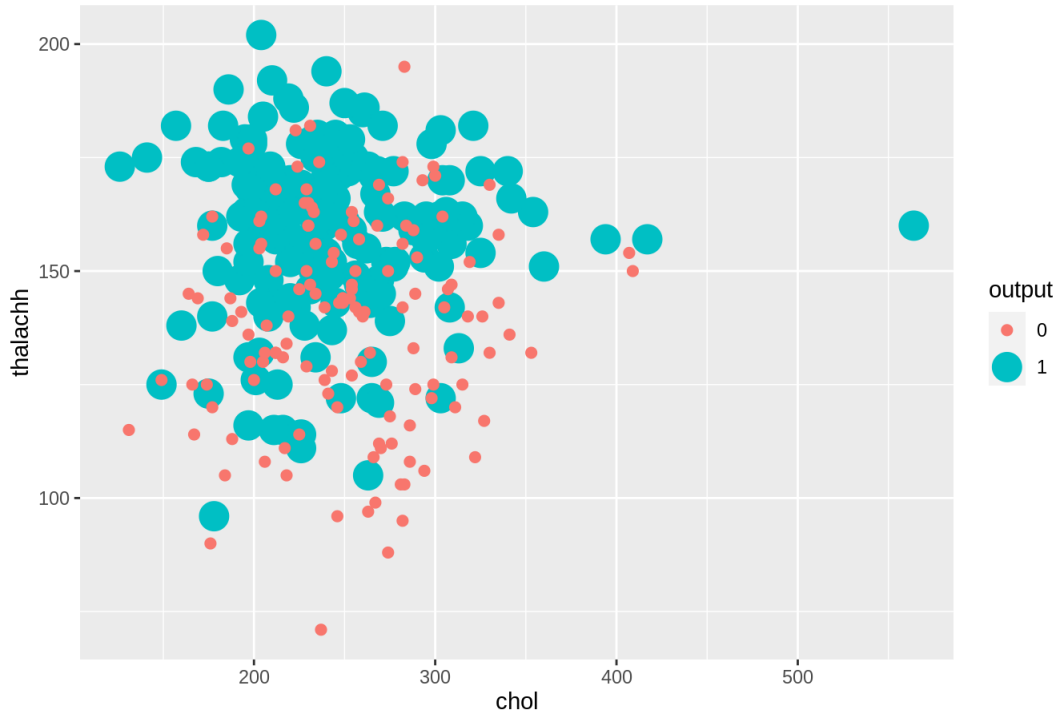
```
library(corrplot)
```

```
## corrplot 0.92 loaded
```

```
ggplot(df, aes(chol, thalach, color = output, size = output)) +  
  geom_point() +  
  ggtitle("kolesterol,maksimum kalp atışı ve kalp krizi olasılığı")
```

```
## Warning: Using size for a discrete variable is not advised.
```

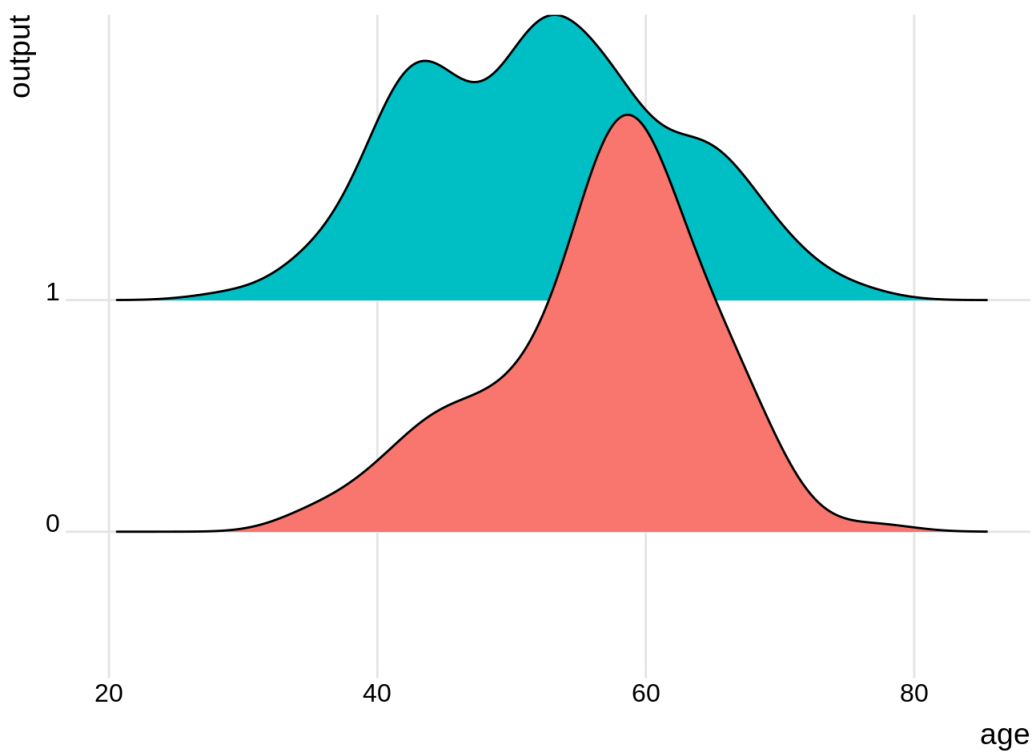
## kolesterol,maksimum kalp atışı ve kalp krizi olasılığı



```
library(ggribes)
```

```
ggplot(df, aes(x = age, y = output, fill = cp)) +  
  geom_density_ridges() +  
  theme_ridges() +  
  theme(legend.position = "none")
```

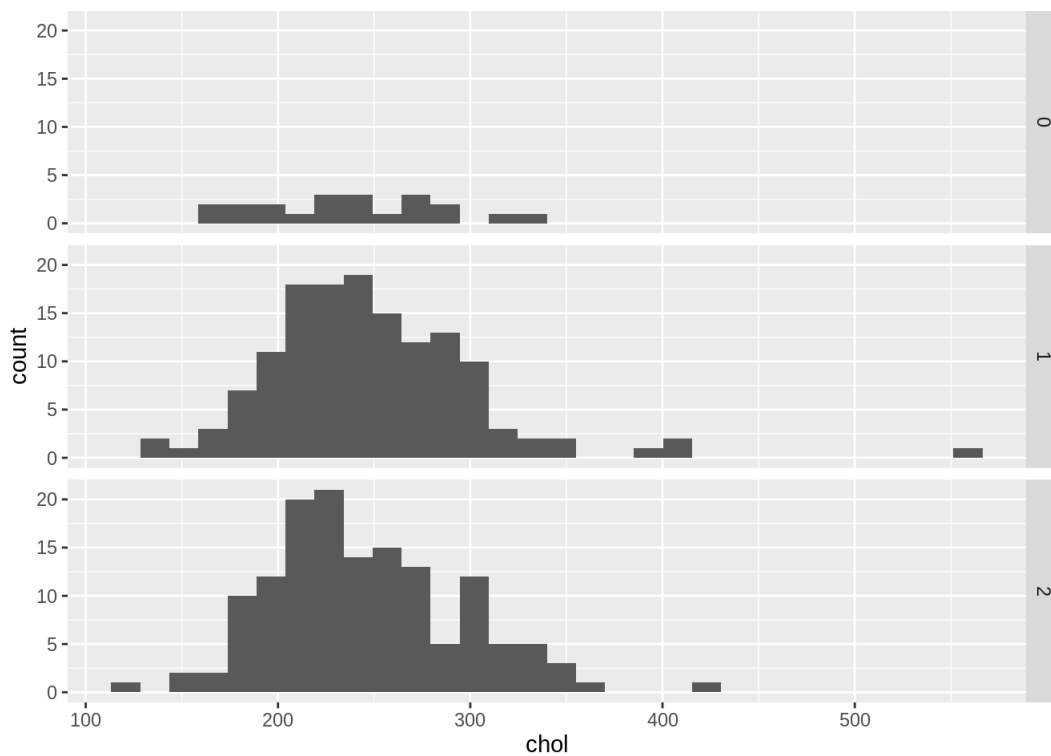
```
## Picking joint bandwidth of 2.8
```



```
ggplot(df, aes(chol)) +  
  geom_histogram() +  
  facet_grid(slp ~ .)
```

*#ayrı ayrı verir*

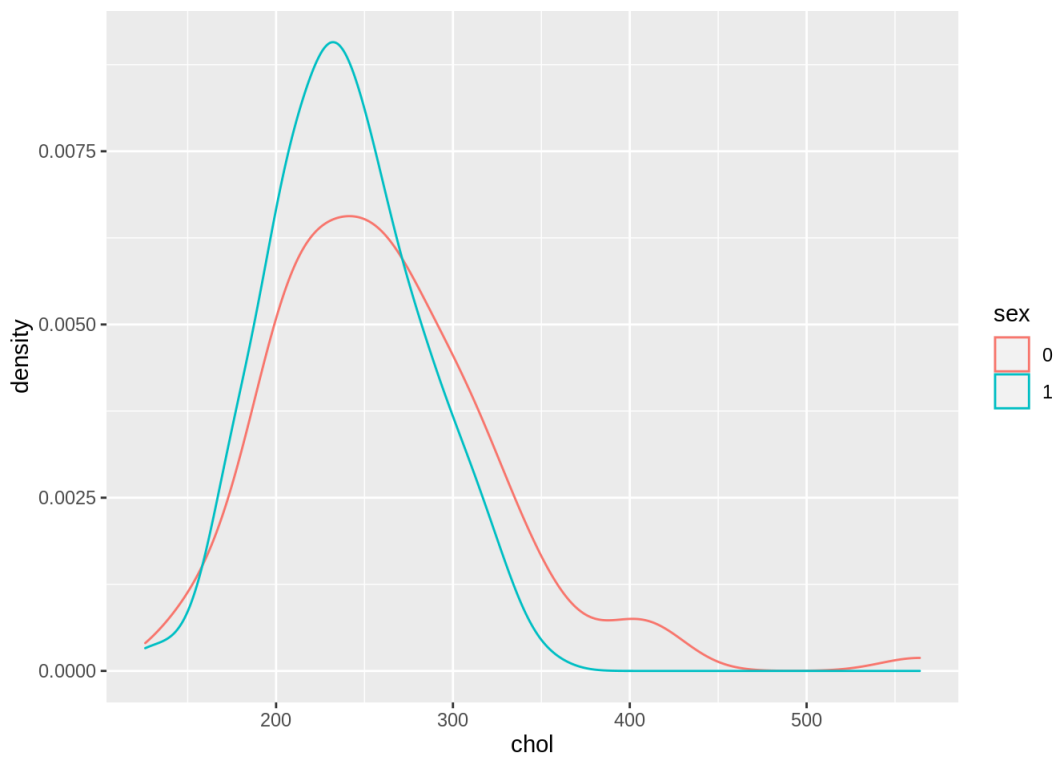
## `stat\_bin()` using `bins = 30`. Pick better value with `binwidth`.



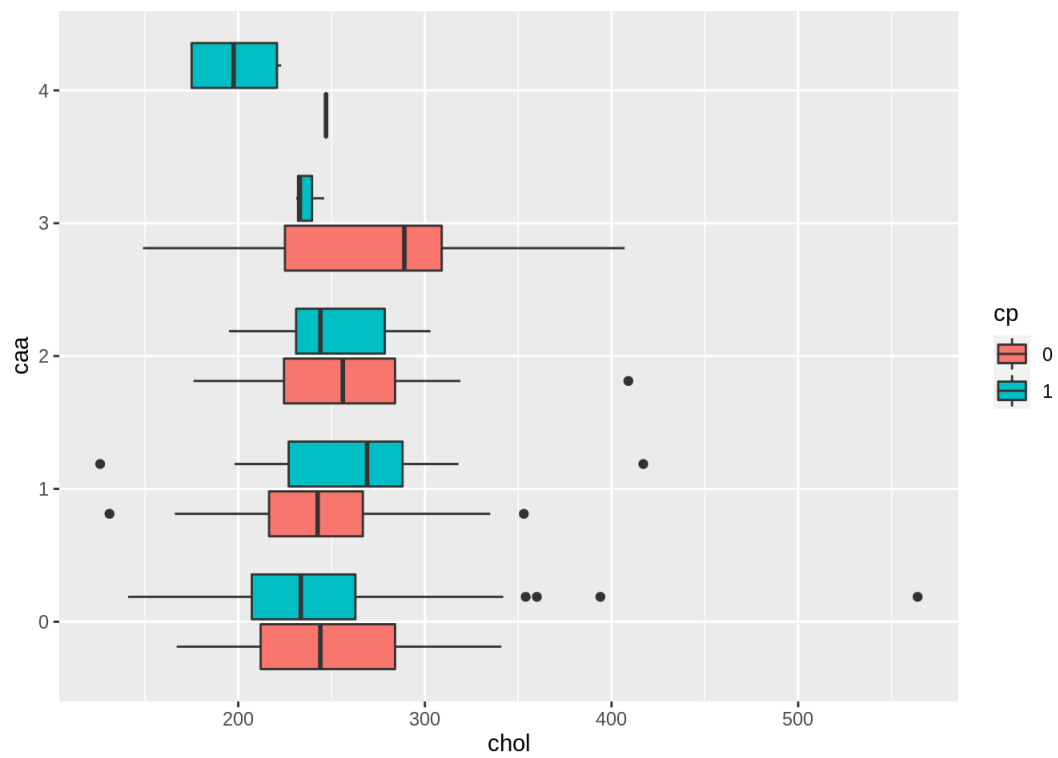
*#çoklu frekans*

```
ggplot(df, aes(chol, y = ..density..)) +  
  geom_density(aes(colour = sex), binwidth = 500)
```

## Warning: Ignoring unknown parameters: binwidth



```
ggplot(df, aes(x = chol, y = caa, fill = cp)) +  
  geom_boxplot()
```



```
ggplot(df, aes(x = chol, y = oldpeak, fill = sex)) +  
  geom_violin(alpha = 0.6)
```

## Warning: position\_dodge requires non-overlapping x intervals

