Survival of patients with heart failure

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Source of data: UCI machine learning repository:https://archive.ics.uci.edu/dataset/519/heart+failure+clinical+records

Upload date of data: 2nd April 2020

Owner of the data: The original dataset version was collected by Tanvir Ahmad, Assia Munir, Sajjad Haider Bhatti, Muhammad Aftab, and Muhammad Ali Raza in 2015. Current dataset has slight difference from original dataset in terms of names of columns which was done by D. Chicco and Giuseppe Jurman in their article published on BMC Medical Informatics and Decision Making. Link:https://bmcmedinformdecismak.biomedcentral.com/articles/10.1186/s12911-020-1023-5#article-info

Objective of data collection: Dataset was collected for analyzing the survival pattern of heart failure patients who were admitted to Institute of Cardiology and Allied hospital Faisalabad-Pakistan during April-December (2015). And also rank the features which corresponds to the most important risk factors.

Analysis of data by other researchers: D.Chicco and Giuseppe Jurman found in their analysis that serum creatinine and ejection fraction plays a very important role in understanding if a patient will survive after heart failure or not. This discovery has the potential to impact on clinical practice, becoming a new supporting tool for physicians when predicting if a heart failure patient will survive or not.

Description of data set: This dataset was collected during April–December 2015. It contains the medical records of 299 heart failure patients collected at the Faisalabad Institute of Cardiology and at the Allied Hospital in Faisalabad (Punjab, Pakistan). The patients consisted of 105 women and 194 men, and their ages range between 40 and 95 years old. All 299 patients had left ventricular systolic dysfunction and had previous heart failures that put them in classes III or IV of New York Heart Association (NYHA) classification of the stages of heart failure. The dataset contains 13 features, which report clinical, body, and lifestyle information.

Description of columns:

```
column_des<-read.csv("C:/Users/rajsh/OneDrive/Documents/Rproject/Book1.csv")
print(column_des)</pre>
```

Feature		##
1 Age	1	##
2 Anaemia	2	##
3 High blood pressure	3	##
4 Creatinine phosphokinase(CPK)	4	##
5 Diabetes	5	##
6 Ejection fraction	6	##
7 Sex	7	##
8 Platelets	8	##
9 Serum creatinine	9	##
10 Serum sodium	10	##

```
## 11
                              Smoking
## 12
                                 Time
## 13
                (target) death event
##
                                                                           Measurement
                                                        Explanation
## 1
                                                 Age of the patient
                                                                                 Years
## 2
                                                                               Boolean
                       Decrease of red blood cells or hemoglobin
## 3
                                                                               Boolean
                                     If a patient has hypertension
## 4
                             Level of the CPK enzyme in the blood
                                                                                 mcg/L
## 5
                                       If the patient has diabetes
                                                                               Boolean
##
  6
      Percentage of blood leaving the heart at each contraction
                                                                            Percentage
## 7
                                                       Woman or man
                                                                                Binary
                                            Platelets in the blood kiloplatelets/mL
## 8
## 9
                                 Level of creatinine in the blood
                                                                                 mg/dL
## 10
                                      Level of sodium in the blood
                                                                                 mEq/L
## 11
                                                                               Boolean
                                              If the patient smokes
## 12
                                                   Follow-up period
                                                                                  Days
## 13
                 If the patient died during the follow-up period
                                                                               Boolean
##
                     Range
## 1
              [40, \ldots, 95]
##
  2
                      0, 1
## 3
                      0, 1
## 4
            [23, \ldots, 7861]
## 5
                       0, 1
              [14,..., 80]
## 6
## 7
                       0, 1
## 8
      [25.01,..., 850.00]
## 9
          [0.50, \ldots, 9.40]
## 10
            [114, \ldots, 148]
## 11
                       0, 1
## 12
               [4, \ldots, 285]
## 13
                       0, 1
```

The hospital physician considered a patient having **anaemia** if haematocrit levels were lower than 36%. The **creatinine phosphokinase (CPK)** states the level of the CPK enzyme in blood. When a muscle tissue gets damaged, CPK flows into the blood. Therefore, high levels of CPK in the blood of a patient might indicate a heart failure or injury. It is measured in microgram per litre (mcg/L). The **ejection fraction** states the percentage of how much blood the left ventricle pumps out with each contraction. The **serum creatinine** is a waste product generated by creatine, when a muscle breaks down. Doctors focus on serum creatinine in blood to check kidney function. If a patient has high levels of serum creatinine, it may indicate renal dysfunction. A normal range for serum creatinine is approximately 0.6 to 1.3 milligrams per deciliter (mg/dL) for adult males and 0.5 to 1.2 mg/dL for adult females. The **serum sodium test** is a routine blood exam that indicates if a patient has normal levels of sodium in the blood. An abnormally low level of sodium(<125 mEq/L) in the blood might be caused by heart failure.mEq/L represents milliequivalents per liter. The **death event** feature, that we use as the target in our binary classification study, states if the patient died or survived before the end of the follow-up period, that was 130 days on average.

Purpose of my project: Carrying out EDA to look for important factors for predicting survival of patients from heart failure.

```
#loading necessary libraries
library(ggplot2)
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
##
library(moments)
#Loading dataset
df<-read.csv("C:/Users/rajsh/OneDrive/Documents/Rproject/heart_failure_clinical_records_dataset.csv")
     age anaemia creatinine_phosphokinase diabetes ejection_fraction
## 1 75
                                     582
                                               0
## 2 55
              0
                                    7861
                                                                38
                                               0
## 3 65
              0
                                     146
                                               0
                                                                20
## 4 50
              1
                                     111
                                                                20
## 5 65
                                     160
                                                                20
              1
                                               1
## 6 90
                                      47
                                               0
                                                                40
    high_blood_pressure platelets serum_creatinine serum_sodium sex smoking time
## 1
                          265.000
                                             1.9
                      1
                                                           130 1
                          263.358
## 2
                                              1.1
                                                           136
                                                                         0
                                                                              6
                                                                 1
## 3
                         162.000
                                              1.3
                                                           129
                                                                         1
                                                                              7
                                                                1
                                                                              7
## 4
                      0 210.000
                                              1.9
                                                           137 1
                                                                        0
## 5
                          327.000
                                              2.7
                                                           116 0
                                                                              8
                          204.000
                                              2.1
                                                                        1
## 6
                                                           132 1
                                                                              8
##
   DEATH_EVENT
## 1
## 2
              1
## 3
              1
## 4
              1
## 5
              1
## 6
#Structure of data
cat("Shape of the dataset is:",dim(df),"\n")
## Shape of the dataset is: 299 13
str(df)
## 'data.frame':
                   299 obs. of 13 variables:
## $ age
                             : num 75 55 65 50 65 90 75 60 65 80 ...
## $ anaemia
                             : int 0001111101...
\#\# $ creatinine_phosphokinase: int ~582~7861~146~111~160~47~246~315~157~123~\dots
## $ diabetes
                             : int 0000100100...
```

```
## $ ejection_fraction
                                   20 38 20 20 20 40 15 60 65 35 ...
                            : int
## $ high_blood_pressure
                                   1000010001...
                            : int
## $ platelets
                                   265 263 162 210 327 ...
                            : num
                                  1.9 1.1 1.3 1.9 2.7 2.1 1.2 1.1 1.5 9.4 ...
## $ serum_creatinine
                            : num
## $ serum sodium
                            : int
                                   130 136 129 137 116 132 137 131 138 133 ...
## $ sex
                                  1 1 1 1 0 1 1 1 0 1 ...
                            : int
                                  0 0 1 0 0 1 0 1 0 1 ...
## $ smoking
                            : int
                            : int
                                  4 6 7 7 8 8 10 10 10 10 ...
## $ time
   $ DEATH EVENT
                            : int 111111111...
```

Since age is measured in years and platelets are count so they are integer columns, thus converting them into integer. By description of dataset we know that anaemia, high blood pressure, diabetes, sex and smoking are categorical columns which are encoded with 0 and 1.

```
df$age <- as.integer(df$age)</pre>
str(df)
## 'data.frame':
                    299 obs. of 13 variables:
## $ age
                              : int 75 55 65 50 65 90 75 60 65 80 ...
## $ anaemia
                              : int
                                    0 0 0 1 1 1 1 1 0 1 ...
## $ creatinine_phosphokinase: int
                                     582 7861 146 111 160 47 246 315 157 123 ...
                                    0 0 0 0 1 0 0 1 0 0 ...
## $ diabetes
                              : int
## $ ejection_fraction
                              : int
                                     20 38 20 20 20 40 15 60 65 35 ...
                                    1 0 0 0 0 1 0 0 0 1 ...
## $ high_blood_pressure
                              : int
## $ platelets
                              : num
                                    265 263 162 210 327 ...
## $ serum_creatinine
                                    1.9 1.1 1.3 1.9 2.7 2.1 1.2 1.1 1.5 9.4 ...
                              : num
                                    130 136 129 137 116 132 137 131 138 133 ...
## $ serum_sodium
                              : int
## $ sex
                                    1 1 1 1 0 1 1 1 0 1 ...
                              : int
                                    0 0 1 0 0 1 0 1 0 1 ...
## $ smoking
                              : int
                                    4 6 7 7 8 8 10 10 10 10 ...
## $ time
                              : int
   $ DEATH EVENT
                              : int 111111111...
#Getting basic descriptive statistic summary of categorical columns
#For Anaemia
Anaemia df <- data.frame(
  Anaemia = c("Present", "Absent"),
  Full_sampleNumber = c(sum(df$anaemia), 299 - sum(df$anaemia)),
  Full_samplePercentage = c(sum(df$anaemia) * 100 / nrow(df), (299 - sum(df$anaemia)) * 100 / nrow(df))
 Dead patientsNumber = c(sum(df[df$DEATH EVENT == 1, ]$anaemia),sum(df$DEATH EVENT)-sum(df[df$DEATH EVENT)
  Dead_patientsPercentage = c(sum(df[df$DEATH_EVENT == 1, ]$anaemia) * 100 /sum(df$DEATH_EVENT), (sum(df
  Survived_patientsNumber = c(sum(df[df$DEATH_EVENT == 0, ]$anaemia), ((299-sum(df$DEATH_EVENT)) - sum(
  Survived_patientsPercentage = c(sum(df[df$DEATH_EVENT == 0, ]$anaemia) * 100 /(299-sum(df$DEATH_EVENT
print(Anaemia_df)
     Anaemia Full_sampleNumber Full_samplePercentage Dead_patientsNumber
                           129
                                            43.14381
## 1 Present
                                                                      46
## 2 Absent
                           170
                                            56.85619
                                                                      50
     Dead_patientsPercentage Survived_patientsNumber Survived_patientsPercentage
                    47.91667
                                                  83
## 1
```

120

59.1133

2

52.08333

```
#For High blood pressure
High_blood_pressure_df <- data.frame(</pre>
  High blood pressure = c("Present", "Absent"),
  Full_sampleNumber = c(sum(df$high_blood_pressure), 299 - sum(df$high_blood_pressure)),
  Full_samplePercentage = c(sum(df$high_blood_pressure) * 100 / nrow(df), (299 - sum(df$high_blood_pres
  Dead_patientsNumber = c(sum(df[df$DEATH_EVENT == 1, ]$high_blood_pressure), sum(df$DEATH_EVENT) - sum
  Dead_patientsPercentage = c(sum(df[df$DEATH_EVENT == 1, ]$high_blood_pressure) * 100 / sum(df$DEATH_E
  Survived_patientsNumber = c(sum(df[df$DEATH_EVENT == 0, ]$high_blood_pressure), (299 - sum(df$DEATH_E
  Survived_patientsPercentage = c(sum(df[df$DEATH_EVENT == 0, ]$high_blood_pressure) * 100 / (299 - sum
print(High_blood_pressure_df)
    High_blood_pressure Full_sampleNumber Full_samplePercentage
## 1
                 Present
                                        105
## 2
                  Absent
                                        194
                                                         64.88294
    Dead_patientsNumber Dead_patientsPercentage Survived_patientsNumber
##
## 1
                                          40.625
                                           59.375
                      57
                                                                      137
## 2
    Survived_patientsPercentage
## 1
                        32.51232
## 2
                        67.48768
# For Diabetes
Diabetes_df <- data.frame(</pre>
  Diabetes = c("Present", "Absent"),
  Full_sampleNumber = c(sum(df$diabetes), 299 - sum(df$diabetes)),
  Full_samplePercentage = c(sum(df$diabetes) * 100 / nrow(df), (299 - sum(df$diabetes)) * 100 / nrow(df
  Dead patientsNumber = c(sum(df[df$DEATH EVENT == 1, ]$diabetes), sum(df$DEATH EVENT) - sum(df[df$DEATH
  Dead_patientsPercentage = c(sum(df[df$DEATH_EVENT == 1, ]$diabetes) * 100 / sum(df$DEATH_EVENT), (sum
  Survived_patientsNumber = c(sum(df[df$DEATH_EVENT == 0, ]$diabetes), ((299 - sum(df$DEATH_EVENT)) - s
  Survived_patientsPercentage = c(sum(df[df$DEATH_EVENT == 0, ]$diabetes) * 100 / (299 - sum(df$DEATH_E
print(Diabetes df)
##
    Diabetes Full_sampleNumber Full_samplePercentage Dead_patientsNumber
## 1 Present
                            125
                                              41.80602
## 2
      Absent
                            174
                                              58.19398
                                                                        56
   Dead_patientsPercentage Survived_patientsNumber Survived_patientsPercentage
## 1
                    41.66667
                                                   85
                                                                         41.87192
## 2
                    58.33333
                                                                         58.12808
                                                  118
# For Sex
Sex_df <- data.frame(</pre>
  Sex = c("Man", "Female"),
  Full_sampleNumber = c(sum(df$sex), 299 - sum(df$sex)),
  Full_samplePercentage = c(sum(df$sex) * 100 / nrow(df), (299 - sum(df$sex)) * 100 / nrow(df)),
  Dead_patientsNumber = c(sum(df[df$DEATH_EVENT == 1, ]$sex), sum(df$DEATH_EVENT) - sum(df[df$DEATH_EVE
  Dead_patientsPercentage = c(sum(df[df$DEATH_EVENT == 1, ]$sex) * 100 / sum(df$DEATH_EVENT), (sum(df$DEATH_EVENT)
  Survived_patientsNumber = c(sum(df[df$DEATH_EVENT == 0, ]$sex), ((299 - sum(df$DEATH_EVENT)) - sum(df
  Survived_patientsPercentage = c(sum(df[df$DEATH_EVENT == 0, ]$sex) * 100 / (299 - sum(df$DEATH_EVENT)
print(Sex_df)
```

```
##
                      Sex Full_sampleNumber Full_samplePercentage Dead_patientsNumber
## 1
                      Man
                                                                         194
                                                                                                                                                                                                 62
                                                                                                                        64.88294
## 2 Female
                                                                        105
                                                                                                                        35.11706
                                                                                                                                                                                                 34
             Dead_patientsPercentage Survived_patientsNumber Survived_patientsPercentage
## 1
                                                       64.58333
                                                                                                                                          132
                                                                                                                                                                                                          65.02463
## 2
                                                       35.41667
                                                                                                                                            71
                                                                                                                                                                                                          34.97537
# For Smoking
Smoking_df <- data.frame(</pre>
     Smoking = c("Present", "Absent"),
     Full_sampleNumber = c(sum(df\$smoking), 299 - sum(df\$smoking)),
     Full_samplePercentage = c(sum(df$smoking) * 100 / nrow(df), (299 - sum(df$smoking)) * 100 / nrow(df))
     Dead_patientsNumber = c(sum(df[df$DEATH_EVENT == 1, ]$smoking), sum(df$DEATH_EVENT) - sum(df[df$DEATH
     Dead_patientsPercentage = c(sum(df[df$DEATH_EVENT == 1, ]$smoking) * 100 / sum(df$DEATH_EVENT), (sum(
     Survived_patientsNumber = c(sum(df[df$DEATH_EVENT == 0, ]$smoking), ((299 - sum(df$DEATH_EVENT)) - su
     Survived_patientsPercentage = c(sum(df[df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 0, ]$smoking) * 100 / (299 - sum(df$DEATH_EVENT == 
print(Smoking_df)
              Smoking Full_sampleNumber Full_samplePercentage Dead_patientsNumber
##
## 1 Present
                                                                              96
                                                                                                                           32.10702
                                                                           203
                                                                                                                           67.89298
                                                                                                                                                                                                    66
## 2 Absent
##
             Dead_patientsPercentage Survived_patientsNumber Survived_patientsPercentage
```

66

137

32.51232

67.48768

We can clearly observe that: 1) Out of 299 patients 129 were found to have anaemia which is about 43% of total population. Only 64.34% patient who had anaemia could survive till follow-up period. 2) Out of 299 patients 105 were found to have high blood pressure which is about 35% of total population. Only 62.85% patient who had high blood pressure could survive till follow-up period. 3) Out of 299 patients 125 were found to have diabetes which is about 42% of total population. Only 68% patient who had diabetes could survive till follow-up period. 4) Male constituted 64.88% and Female constituted 35.12% of total population. 5) Out of 299 patients 96 were found to have diabetes which is about 32% of total population. Only 68.75% patient who had diabetes could survive till follow-up period.

Univariate analysis

31.25

68.75

1

2

```
#Getting basic descriptive statistic summary of age column
cat("For age","\n")
## For age
summary(df$age)
##
      Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                               Max.
     40.00
             51.00
                     60.00
                             60.83
                                      70.00
                                              95.00
##
cat("Standard deviation:",sd(df$age),"\n")
## Standard deviation: 11.895
```

```
cat("Skewness:",skewness(df$age),"\n")

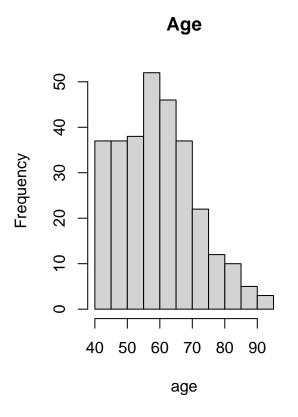
## Skewness: 0.4220415

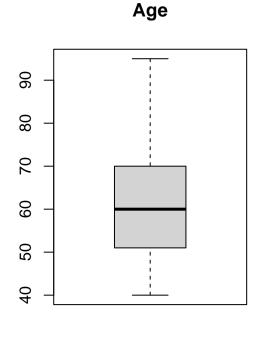
cat("Kurtosis:",kurtosis(df$age),"\n","\n")

## Kurtosis: 2.798664

##

#For age
par(mfrow=c(1,2))
hist(df$age,main="Age",xlab="age")
boxplot(df$age,main="Age")
```





```
par(mfrow=c(1,1))
```

Age of patients lie between range 40-95 years with average age of 60.83 years and standard deviation of 11.895. Most of the patients have age 55-65 years.

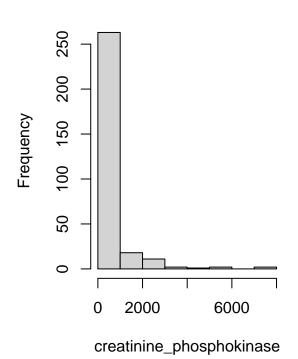
```
\begin{tabular}{ll} \#Getting basic descriptive statistic summary of Creatinine phosphokinase \\ \tt cat("For creatinine_phosphokinase","\n") \\ \end{tabular}
```

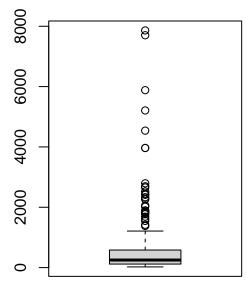
For creatinine_phosphokinase

```
summary(df$creatinine_phosphokinase)
##
     Min. 1st Qu.
                    Median
                              Mean 3rd Qu.
                                              Max.
##
             116.5
                     250.0
                             581.8
                                     582.0 7861.0
cat("Standard deviation:",sd(df$creatinine_phosphokinase),"\n")
## Standard deviation: 970.2879
cat("Skewness:",skewness(df$creatinine_phosphokinase),"\n")
## Skewness: 4.440689
cat("Kurtosis:",kurtosis(df$creatinine_phosphokinase),"\n","\n")
## Kurtosis: 27.71046
##
#For Creatinine phosphokinase
par(mfrow=c(1,2))
hist(df$creatinine_phosphokinase,main="Creatinine phosphokinase",xlab="creatinine_phosphokinase")
boxplot(df$creatinine_phosphokinase,main="Creatinine phosphokinase")
```

Creatinine phosphokinase

Creatinine phosphokinase



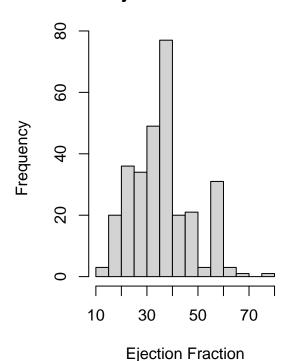


```
par(mfrow=c(1,1))
```

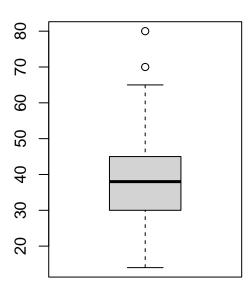
Creatinine phosphokinase column is highly right skewed & peaked with range of 23-7861, standard deviation of 970.28, Skewness of 4.44, kurtosis of 27.7. From the box-plot we can see that it has many outliers.

```
#Getting basic descriptive statistic summary of Ejection fraction
cat("For ejection_fraction","\n")
## For ejection_fraction
summary(df$ejection_fraction)
##
                              Mean 3rd Qu.
     Min. 1st Qu. Median
                                              Max.
##
     14.00
            30.00
                    38.00
                             38.08
                                     45.00
                                             80.00
cat("Standard deviation:",sd(df$ejection_fraction),"\n")
## Standard deviation: 11.83484
cat("Skewness:",skewness(df$ejection_fraction),"\n")
## Skewness: 0.5525927
cat("Kurtosis:",kurtosis(df$ejection_fraction),"\n","\n")
## Kurtosis: 3.02072
##
#For Ejection fraction
par(mfrow=c(1,2))
hist(df$ejection_fraction, main="Ejection Fraction", xlab="Ejection Fraction")
boxplot(df$ejection_fraction, main="Ejection Fraction")
```

Ejection Fraction



Ejection Fraction



```
par(mfrow=c(1,1))
```

Ejection fraction column has range of 14-80 , mean of 38.08 , standard deviation of 11.83, Skewness of 0.55 , kurtosis of 3.02.

```
#Getting basic descriptive statistic summary of Platelets
cat("For plateles","\n")
```

For plateles

```
summary(df$platelets)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 25.1 211.5 262.0 262.7 303.5 850.0
```

```
cat("Standard deviation:",sd(df$platelets),"\n")
```

Standard deviation: 98.39959

```
cat("Skewness:",skewness(df$platelets),"\n")
```

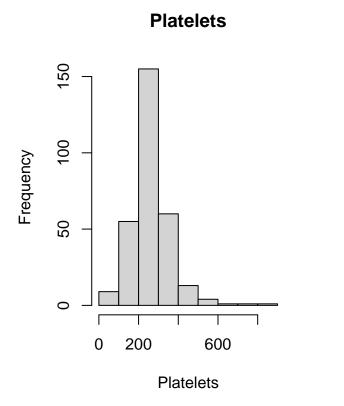
Skewness: 1.425942

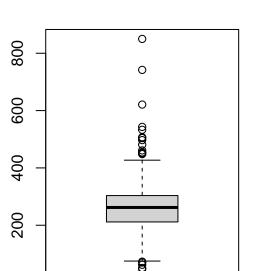
```
cat("Kurtosis:",kurtosis(df$platelets),"\n","\n")

## Kurtosis: 8.950815

##

#For Platelets
par(mfrow=c(1,2))
hist(df$platelets, main="Platelets", xlab="Platelets")
boxplot(df$platelets, main="Platelets")
```





Platelets

par(mfrow=c(1,1))

0

Platelets column is high peaked column with range of 25100-850000, mean of 263358, standard deviation of 97804.24, Skewness of 1.45, kurtosis of 9.08. From the box-plot we can see that it has many outliers.

```
#Getting basic descriptive statistic summary of Serum creatinine
cat("For serum_creatinine","\n")
```

For serum_creatinine

```
summary(df$serum_creatinine)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 0.500 0.900 1.100 1.394 1.400 9.400
```

```
cat("Standard deviation:",sd(df$serum_creatinine),"\n")

## Standard deviation: 1.03451

cat("Skewness:",skewness(df$serum_creatinine),"\n")

## Skewness: 4.43361

cat("Kurtosis:",kurtosis(df$serum_creatinine),"\n","\n")

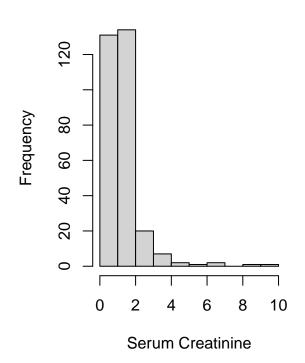
## Kurtosis: 28.37835

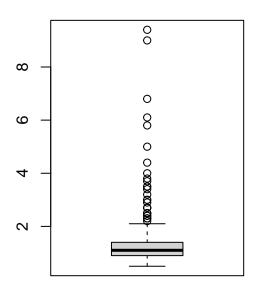
##

#For Serum creatinine
par(mfrow=c(1,2))
hist(df$serum_creatinine, main="Serum Creatinine", xlab="Serum Creatinine")
boxplot(df$serum_creatinine, main="Serum Creatinine")
```

Serum Creatinine

Serum Creatinine





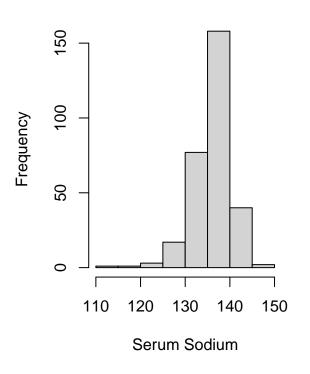
par(mfrow=c(1,1))

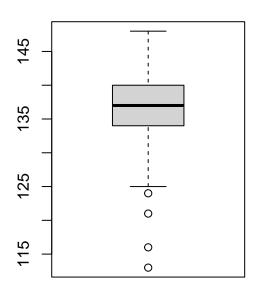
Serum creatinine is highly right skewed & peaked with range of 0.5-9.4, standard deviation of 1.03, Skewness of 4.43, kurtosis of 28.37. From the box-plot we can see that it has many outliers.

```
{\it \#Getting~basic~descriptive~statistic~summary~of~Serum~sodium}
cat("For serum_sodium","\n")
## For serum_sodium
summary(df$serum_sodium)
##
     Min. 1st Qu. Median
                             Mean 3rd Qu.
                                              Max.
##
     113.0 134.0 137.0 136.6 140.0
                                             148.0
cat("Standard deviation:",sd(df$serum_sodium),"\n")
## Standard deviation: 4.412477
cat("Skewness:",skewness(df$serum_sodium),"\n")
## Skewness: -1.04287
cat("Kurtosis:",kurtosis(df$serum_sodium),"\n","\n")
## Kurtosis: 7.031142
##
#For Serum sodium
par(mfrow=c(1,2))
hist(df$serum_sodium, main="Serum Sodium", xlab="Serum Sodium")
boxplot(df$serum_sodium, main="Serum Sodium")
```

Serum Sodium

Serum Sodium





par(mfrow=c(1,1))

Serum sodium has range of 113-148, mean of 136.6, standard deviation of 4.41, Skewness of -1.04, kurtosis of 7.03. It is left skewed.

```
#Getting basic descriptive statistic summary of Death event
cat("For time","\n")
```

For time

```
summary(df$time)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 4.0 73.0 115.0 130.3 203.0 285.0
```

```
cat("Standard deviation:",sd(df$time),"\n")
```

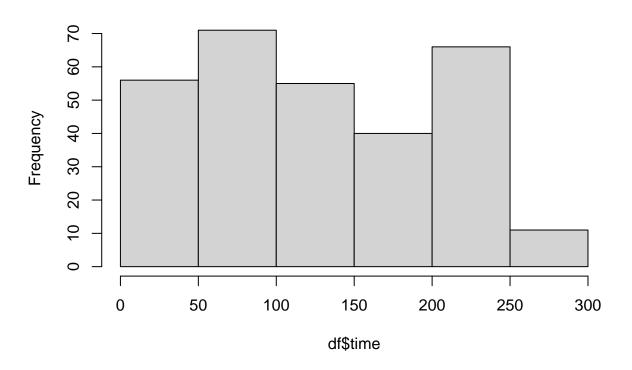
Standard deviation: 77.61421

```
cat("Skewness:",skewness(df$time),"\n")
```

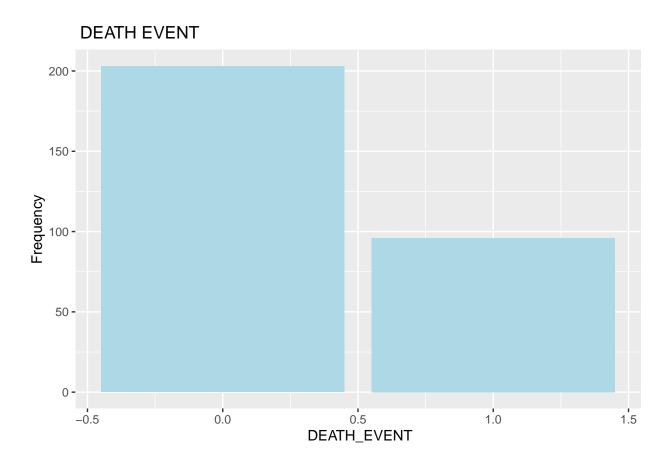
Skewness: 0.1271606

```
cat("Kurtosis:",kurtosis(df$time),"\n")
## Kurtosis: 1.788126
hist(df$time)
```

Histogram of df\$time

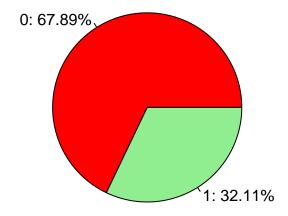


```
#For Death event
ggplot(df, aes(x = DEATH_EVENT)) +geom_bar(fill = "lightblue") +labs(title = " DEATH EVENT", x = " DEATH
```



pie(table(df\$ DEATH_EVENT), labels = paste0(names(table(df\$ DEATH_EVENT)), ": ", round(table(df\$ DEATH_EVENT))

DEATH_EVENT Distribution

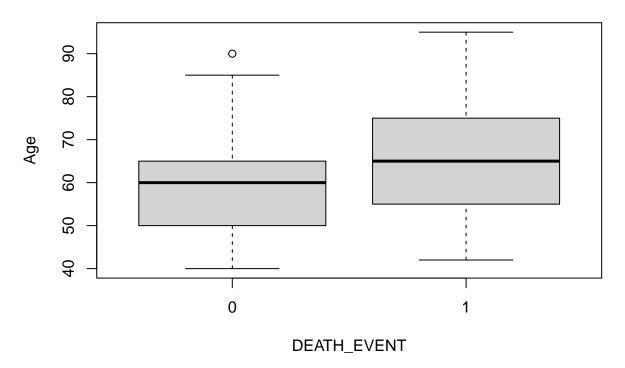


Among 299 patients 67.89% died before follow-up period.

Bivariate analysis

```
#Death vs Age
boxplot(df$age ~ df$DEATH_EVENT, main = "Death vs Age", xlab = "DEATH_EVENT", ylab = "Age")
```

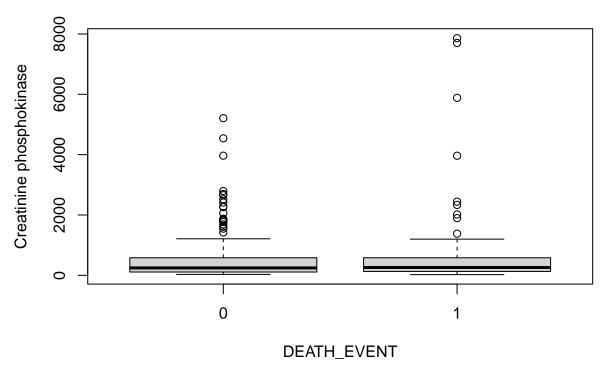
Death vs Age

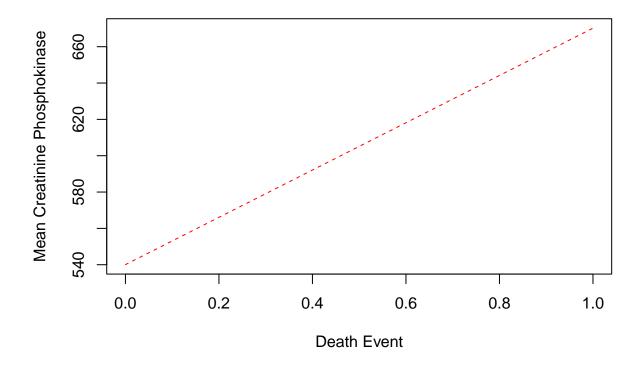


From death vs age plot we see that young age group people have more chances of survival.

```
#Death vs Creatinine phosphokinase
boxplot(df$creatinine_phosphokinase ~ df$DEATH_EVENT, main = "Death vs Creatinine phosphokinase", xlab
```

Death vs Creatinine phosphokinase

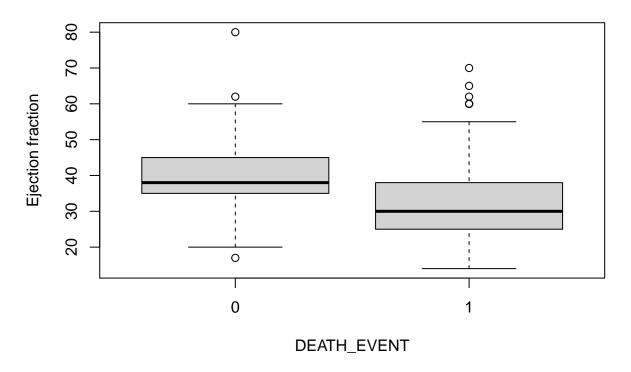




From the line plot we can see that on an average the patients who died had high value of CPK. We might consider it as potential factor to predict survival of patients. But since the range of CPK was too high than its mean so we cannot say with high certainty that it is a good factor to predict survival.

```
#Death vs Ejection fraction
boxplot(df$ejection_fraction ~ df$DEATH_EVENT, main = "Death vs Ejection fraction", xlab = "DEATH_EVENT
```

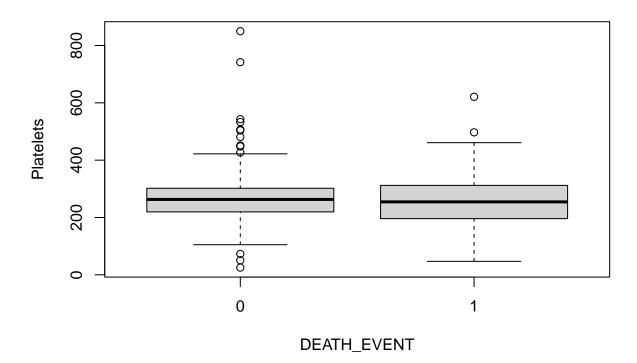
Death vs Ejection fraction

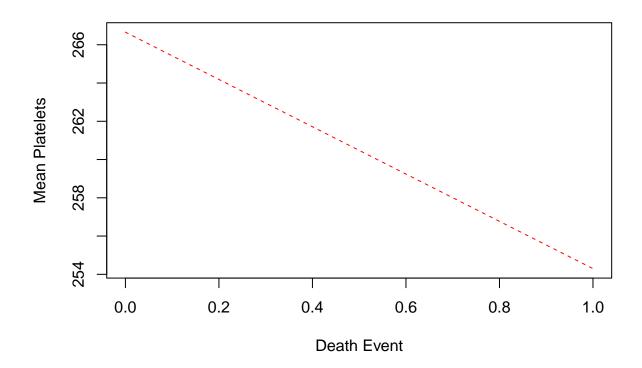


Ejection fraction vs death plot reveals that the people who died had low value of ejection fraction i.e. left ventricle does not pumps effectively with each contraction.

```
#Death vs Platelets
boxplot(df$platelets ~ df$DEATH_EVENT, main = "Death vs Platelets", xlab = "DEATH_EVENT", ylab = "Plate"
```

Death vs Platelets

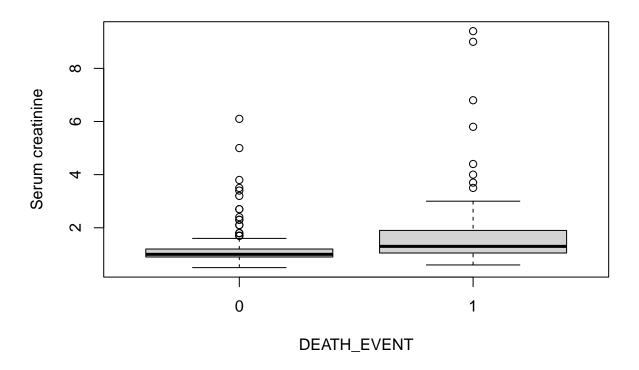




On Average plateletes count remained same for patients who survived and who died. Since there is very slight cange in average of platelets count.

```
#Death vs Serum creatinine
boxplot(df$serum_creatinine ~ df$DEATH_EVENT, main = "Death vs Serum creatinine", xlab = "DEATH_EVENT",
```

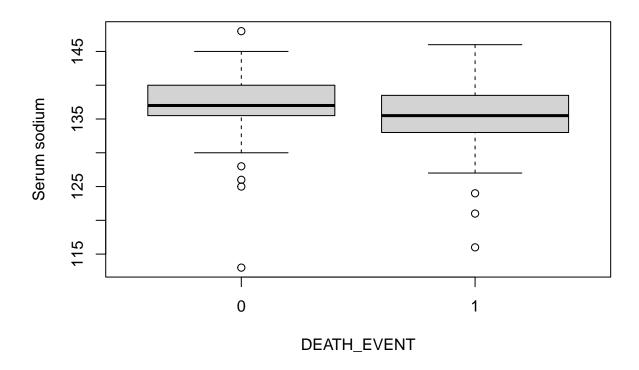
Death vs Serum creatinine



High value for Serum creatinine was observed for patients who died telling that it can be potential factor for predicting survival of patients.

```
#Death vs Serum sodium
boxplot(df$serum_sodium ~ df$DEATH_EVENT, main = "Death vs Serum sodium", xlab = "DEATH_EVENT", ylab =
```

Death vs Serum sodium



Low value for Serum sodium was observed for patients who died telling that it can be potential factor for predicting survival of patients.

#Analysis of survival and sex of patients taken together

```
#Death&Sex vs Age
boxplot(df$age ~ interaction(df$DEATH_EVENT,df$sex), main = "Death&Sex vs Age", xlab = "DEATH_EVENT & S.
```

Death&Sex vs Age



Among all patients male patients who were younger had high chances of survival since among those males who died older ones were majority. And same is observed for females.

#Death&Sex vs Serum Creatinine
boxplot(df\$serum_creatinine ~ interaction(df\$DEATH_EVENT,df\$sex), main = "Death&Sex vs Serum Creatinine")

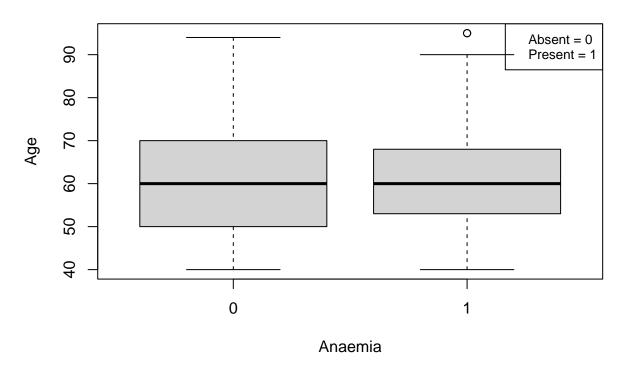
Death&Sex vs Serum Creatinine



In both males and females the patients who had high level of Serum Creatinine died before follow up period. #Relationship between different feature columns

```
#Anaemia vs Age
boxplot(df$age ~ df$anaemia, main = "Anaemia vs Age", xlab = "Anaemia", ylab = "Age")
legend("topright", legend = c("Absent = 0", "Present = 1"),cex=0.8)
```

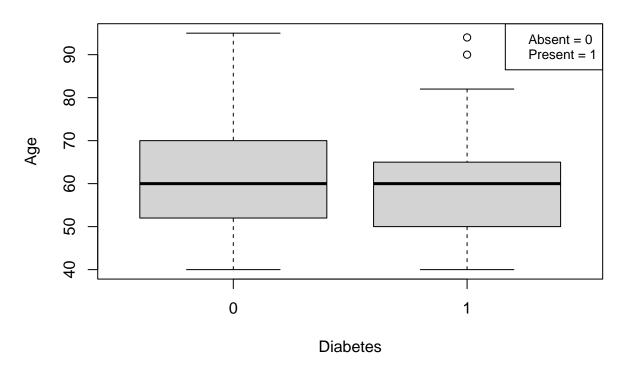
Anaemia vs Age



On average age remained same for patients who had anaemia and who didn't have anaemia.

```
#Diabetes vs Age
boxplot(df$age ~ df$diabetes, main = "Diabetes vs Age", xlab = "Diabetes", ylab = "Age")
legend("topright", legend = c("Absent = 0", "Present = 1"),cex=0.8)
```

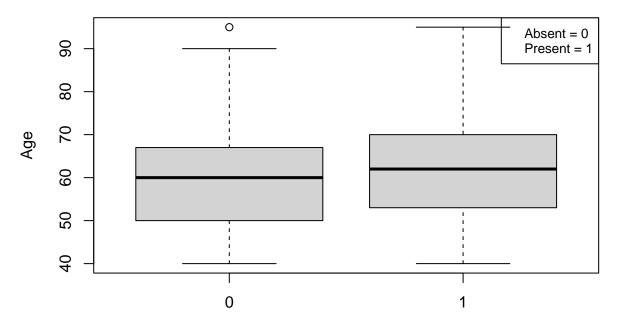
Diabetes vs Age



On average age remained same for patients who had diabetes and who didn't have diabetes.

```
#High blood pressure vs Age
boxplot(df$age ~ df$high_blood_pressure, main = "High blood pressures vs Age", xlab = "High blood pressure
legend("topright", legend = c("Absent = 0", "Present = 1"),cex=0.8)
```

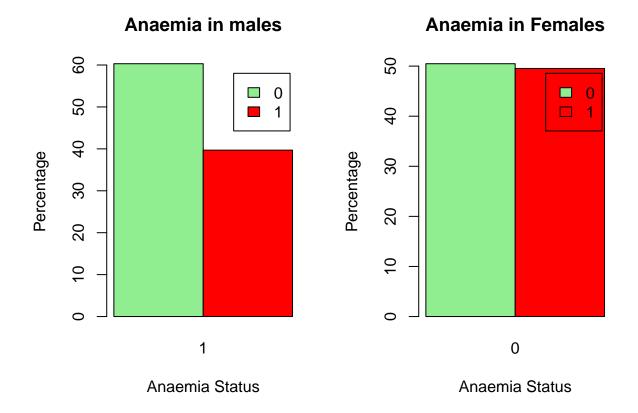
High blood pressures vs Age



High blood pressure

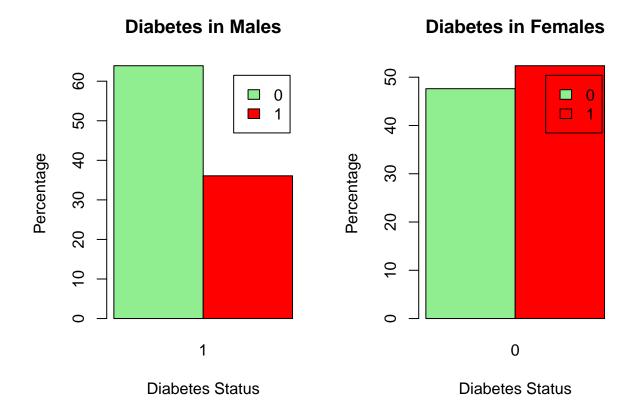
Patients who had high blood pressure were slightly aged than those who didn't had high blood pressure.

```
#Anaemia & Sex
df_males <- subset(df, sex == 1)</pre>
df_females <- subset(df, sex == 0)</pre>
# Create a table for anaemia and sex
x <- table(df_males$anaemia, df_males$sex)</pre>
y <- table(df_females$anaemia, df_females$sex)
x_percentage <- prop.table(x, margin = 2) * 100</pre>
y_percentage <- prop.table(y, margin = 2) * 100 # Calculate percentages across columns (sex)
par(mfrow=c(1,2))
# Barplot for males with percentages
barplot(x_percentage, beside = TRUE, col = c("lightgreen", "red"),
        legend = rownames(y_percentage), main = "Anaemia in males",
        xlab = "Anaemia Status", ylab = "Percentage")
# Barplot for females with percentages
barplot(y_percentage, beside = TRUE, col = c("lightgreen", "red"),
        legend = rownames(y_percentage), main = "Anaemia in Females",
        xlab = "Anaemia Status", ylab = "Percentage")
```



Among male patients around 40% had an aemia and among female around 50% had an aemia. That may mean that females are more prone to an aemia.

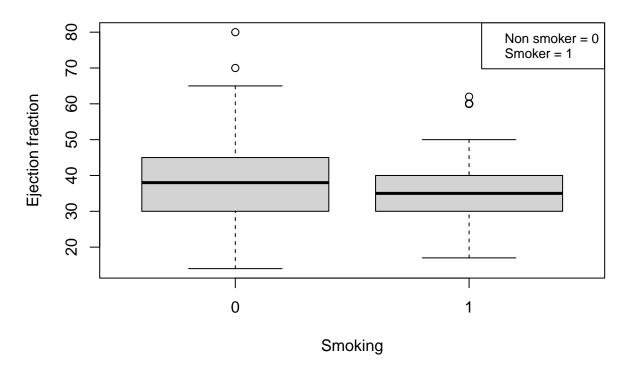
```
df_males <- subset(df, sex == 1)</pre>
df_females <- subset(df, sex == 0)</pre>
# Create a table for diabetes and sex
x <- table(df_males$diabetes, df_males$sex)</pre>
y <- table(df_females$diabetes, df_females$sex)</pre>
x_percentage <- prop.table(x, margin = 2) * 100</pre>
y_percentage <- prop.table(y, margin = 2) * 100 # Calculate percentages across columns (sex)
par(mfrow=c(1,2))
# Barplot for males with percentages
barplot(x_percentage, beside = TRUE, col = c("lightgreen", "red"),
        legend = rownames(y_percentage), main = "Diabetes in Males",
        xlab = "Diabetes Status", ylab = "Percentage")
# Barplot for females with percentages
barplot(y_percentage, beside = TRUE, col = c("lightgreen", "red"),
        legend = rownames(y_percentage), main = "Diabetes in Females",
        xlab = "Diabetes Status", ylab = "Percentage")
```



Among male patients around 35% had diabetes and among female around 52% had diabetes. That may mean that females are more prone to diabetes.

```
#Smoking vs Ejection fraction
boxplot(df$ejection_fraction ~ df$smoking, main = "Smoking vs Ejection fraction", xlab = "Smoking", ylab
legend("topright", legend = c("Non smoker = 0", "Smoker = 1"),cex=0.8)
```

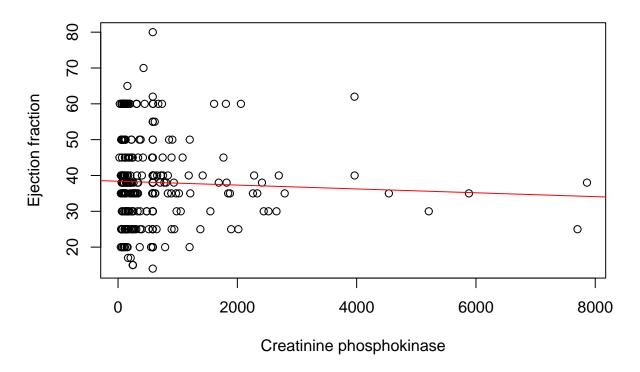
Smoking vs Ejection fraction



Smoker patients had low value of ejection fraction, meaning that smoking may cause decrease in the pumping capacity of heart.

#Creatinine phosphokinase vs Ejection fraction
plot(x=df\$creatinine_phosphokinase,y=df\$ejection_fraction,main="Creatinine phosphokinase vs Ejection fr
abline(lm(df\$ejection_fraction~df\$creatinine_phosphokinase),col="red")

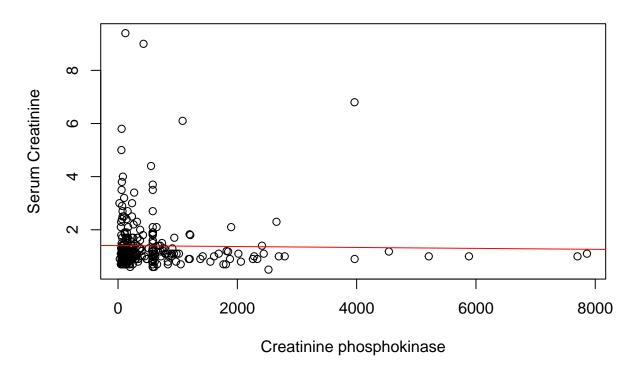
Creatinine phosphokinase vs Ejection fraction



We can see from the graph that as the CPK level increases ejection fraction decreases representing that blood pumping capacity of heart decrease whenever heart injury occurs.

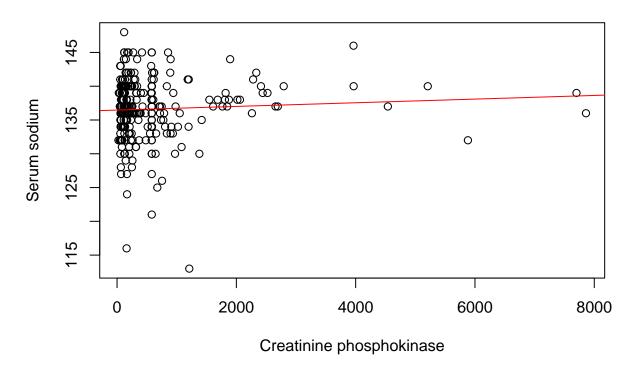
#Creatinine phosphokinase vs Serum Creatinine
plot(x=df\$creatinine_phosphokinase,y=df\$serum_creatinine,main="Creatinine phosphokinase vs Serum Creatinine(lm(df\$serum_creatinine~df\$creatinine_phosphokinase),col="red")

Creatinine phosphokinase vs Serum Creatinine



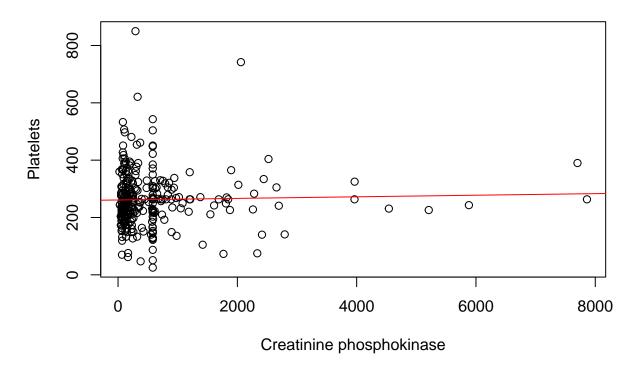
#Creatinine phosphokinase vs Serum sodium
plot(x=df\$creatinine_phosphokinase,y=df\$serum_sodium,main="Creatinine phosphokinase vs Serum sodium",xl
abline(lm(df\$serum_sodium~df\$creatinine_phosphokinase),col="red")

Creatinine phosphokinase vs Serum sodium



#Creatinine phosphokinase vs Platelets
plot(x=df\$creatinine_phosphokinase,y=df\$platelets,main="Creatinine phosphokinase vs Platelets",xlab="Cr
abline(lm(df\$platelets~df\$creatinine_phosphokinase),col="red")

Creatinine phosphokinase vs Platelets



Key findings: 1)Ejection fraction, Serum sodium and Serum creatinine can be considered as important factors to predict survival of patients. 2)Females are more prone to anaemia and diabetes than males. 3)Smoking can cause decrease in pumping capacity of heart. 4)Presence of CPK enzyme may be considered for detection of heart injury.

Splitting Dataset into training and testing data

```
train_indices <- sample(1:nrow(df), 0.75 * nrow(df),replace = F)
train_data <- df[train_indices, ]
test_data <- df[-train_indices, ]
train_data</pre>
```

```
##
        age anaemia creatinine_phosphokinase diabetes ejection_fraction
## 111
                   0
                                                           0
        85
                                              129
                                                                               60
##
   48
         60
                   0
                                              582
                                                           1
                                                                               38
   229
         65
                   0
                                               56
                                                           0
                                                                               25
##
##
  259
         45
                                               66
                                                           1
                                                                               25
                   1
## 78
                   0
         42
                                              102
                                                           1
                                                                               40
   170
         70
                   0
                                              835
                                                           0
                                                                               35
##
                                                           0
## 138
        68
                   1
                                              646
                                                                              25
   226
        75
                   0
                                              675
                                                           1
                                                                              60
   236
        77
                   1
                                              109
                                                           0
                                                                              50
##
   230
        72
                   0
                                                           0
                                                                               25
##
                                              211
## 206
                   1
                                              167
                                                           1
                                                                               45
         50
## 285
         50
                   1
                                               54
                                                           0
                                                                               40
                                              145
                                                           0
                                                                               25
## 185
        58
```

## 19	97 45	0	582	1	38
## 40		0	235	1	38
## 98	3 70	1	59	0	60
## 20)5 78	1	64	0	40
## 28	30 55	0	84	1	38
## 27	70 40	0	582	1	35
## 9	65	0	157	0	65
## 21	10 49	0	972	1	35
## 26	65 61	0	582	1	38
## 10	3 80	0	898	0	25
## 80) 55	0	336	0	45
## 36	69	0	582	1	35
## 26	51 55	0	66	0	40
## 11	L 75	1	81	0	38
## 21		0	582	0	62
## 15		0	115	0	45
## 28		0	582	1	55
## 41		0	582	0	20
## 18		0	1896	1	25
## 17		1	115	0	20
## 14		1	168	0	38
## 23		1	720	1	40
## 15		1	231	1	25
## 15		0	30	1	60
## 44		0	127	1	50
## 86		0	78	0	50
## 21		0	212	1	17
## 22		0	582	1	35
## 28		0	582	0	40
## 10		1	1876	1	35
## 17		1	149	0	38
## 70 ## 26		0	113	1	25
## 25		1 0	135 582	1	38 30
## 10		1	61	1	40
## 29		0	2060	1	60
## 20		1	101	0	40
## 29		0	2413	0	38
## 74		0	224	1	50
## 15		1	910	0	50
## 42		0	124	1	30
## 29		0	1820	0	38
## 53		0	3964	1	62
## 1	75	0	582	0	20
## 22		0	582	0	20
## 18	37 50	0	582	0	50
## 99	60	1	156	1	25
## 96	5 58	1	133	0	60
## 73		0	5882	0	35
## 45	60	1	588	1	60
## 72	2 58	0	582	1	35
## 17	78 49	1	69	0	50
## 27		0	64	0	40
## 28	84 65	0	1688	0	38

## 233	40	1	129	0	35
## 271	44	0	582	1	30
## 64	45	0	582	0	35
## 281	70	0	2695	1	40
## 235	53	1	582	0	45
## 147	52	0	132	0	30
## 124	60	1	582	0	30
## 33	50	1	249	1	35
## 276	45	0	582	0	38
## 175	65	0	198	1	35
## 149	75	1	582	0	30
## 47	51	0	1380	0	25
## 266	50	1	298	0	35
## 137	65	1	59	1	60
## 267	55	0	1199	0	20
## 57	70	1	75	0	35
## 102	75	0	582	0	45
## 262	62	1	655	0	40
## 208	85	0	212	0	38
## 49	80	1	553	0	20
## 141	80	0	805	0	38
## 232	70	0	93	0	35
## 37 ## 68	90 72	1 1	60 110	1 0	50
## 00 ## 77	70	0	92	0	25 60
## 143	50	0	482	1	30
## 51	68	1	577	0	25
## 290	90	1	337	0	38
## 162	45	1	130	0	35
## 81	70	0	69	0	40
## 157	52	1	58	0	35
## 249	40	0	624	0	35
## 58	60	1	607	0	40
## 248	64	0	143	0	25
## 244	73	1	1185	0	40
## 5	65	1	160	1	20
## 139	62	0	281	1	35
## 2	55	0	7861	0	38
## 129	61	0	248	0	30
## 105	60	0	53	0	50
## 134	63	0	193	0	60
## 200	60	0	1211	1	35
## 264	68	1	157	1	60
## 287	60	0	253	0	35
## 255	53	1	446	0	60
## 84	79 50	1	55	0	50
## 299 ## 112	50 55	0	196 60	0 0	45 35
## 112 ## 18	55 45		582	0	
## 18 ## 136	45 75	0	582 582	0	14 40
## 158	75 50	0	250	0	25
## 195	45	0	582	0	20
## 28	70	0	122	1	45
## 250	53	0	207	1	40
200	50	•	201	-	40

## 180	55	0	835	0	40
## 12	62	0	231	0	25
## 75	69	0	582	0	20
## 161	66	1	72	0	40
## 272	51	0	582	1	40
## 140	50	0	1548	0	30
## 198	65	0	167	0	30
## 31	94	0	582	1	38
## 62	50	0	318	0	40
## 173	70	1	171	0	60
## 181	40	0	478	1	30
## 107	55	0	748	0	45
## 209	60	1	2281	1	40
## 121	60	1	737	0	60
## 21	65	1	52	0	25
## 55	60	1	260	1	38
## 128	58	1	200	1	60
## 204	60	0	59	0	25
## 16	82	1	379	0	50
## 3	65	0	146	0	20
## 177	69	0	1419	0	40
## 85 ## 6	59	1	280	1	25
## 6 ## 63	90	1 0	47	0 0	40
## 03 ## 241	55 70	0	109 81	1	35 35
## 241	46	0	168	1	17
## 258	58	0	132	1	38
## 247	55	0	2017	0	25
## 256	52	1	191	1	30
## 172	52	0	3966	0	40
## 123	60	0	96	1	38
## 283	42	0	64	0	30
## 231	60	0	166	0	30
## 148	64	0	1610	0	60
## 186	60	1	104	1	30
## 183	65	0	395	1	25
## 67	42	1	250	1	15
## 279	50	1	1051	1	30
## 234	53	1	707	0	38
## 66	60	0	68	0	20
## 93	42	0	582	0	60
## 39	60	0	2656	1	30
## 146	50	0	185	0	30
## 192	64	1	62	0	60
## 245	54	0	582	1	38
## 46	50	0	582	1	38
## 27	95	1	112	0	40
## 20	48	1	582	1	55
## 110	45	0	292	1	35
## 109	63	0	936	0	38
## 60 ## 131	72 53	0 1	364 1808	1 0	20 60
## 131 ## 165	53 45	0	1808 2442	1	30
## 165	45 59	1	2442 129	0	
## 100	59	1	129	U	45

	400					070	4		٥.	
	130	53	1			270	1		35	
	257	65	0			326	0		38	
	224	47	0			582	0		25	
##	168	59	0			66	1		20	
##	218	54	1			427	0		70	
##	71	41	0			148	0		40	
	89	44	0			84	1		40	
	227	58	1			57			25	
	125	60	0			582	0		40	
	194	73	1			231	1		30	
	213					224			50	
		78	0							
	166	80	0			776			38	
	199	50	1			582			20	
	294	63	1			103	1		35	
	69	70	0			161	0		25	
##	29	58	1			60	0		38	
##	133	46	0			719	0		40	
##	118	85	1			102	0		60	
##	24	53	0			63	1		60	
##	13	45	1			981	0		30	
##	94	60	1			154	0		25	
##	26	80	0			148	1		38	
	50	57	1			129	0		30	
	176	60	1			95	0		60	
	151	72	0			233	0		45	
	115	60	1			754			40	
	76					47			20	
		60 E0	1				0			
	182	59	1			176	1		25	
	286	55	1			170	1		40	
	92	60	0			897	1		45	
	32	85	0			23	0		45	
	38	82	1			855	1		50	
	120	86	0			582	0		38	
	116	58	1			400	0		40	
##	242	65	0			582	1		30	
##	190	40	0			244	0		45	
##	223	42	1			86	0		35	
##	201	63	1			1767	0		45	
##	10	80	1			123	0		35	
##	167	53	0			196	0		60	
	228	55	1			2794			35	
	88	65	1			68			60	
	135	81	0			4540	0		35	
##	19	70	1			125	0		25	
	15	49	1			80	0		30	
	101	65	1			305	0		25	
	292	60	0			320			35 45	
	132	60	1			1082			45	
	196	77	1			418			45	
##		high	_blood_pressu			serum_		serum_sodium		
	111			0	306.000		1.20	132	1	1
	48			1	451.000		0.60	138	1	1
	229			0	237.000		5.00	130	0	0
##	259			0	233.000		0.80	135	1	0

##	78	0	237.000	1.20	140	1	0
	170	1	305.000	0.80	133	0	0
	138	0	305.000	2.10	130	1	0
	226	0	265.000	1.40	125	0	0
	236	1	406.000	1.10	137	1	0
	230	0	274.000	1.20	134	0	0
	206	0	362.000	1.00	136	0	0
	285	0	279.000	0.80	141	1	0
	185	0	219.000	1.20	137	1	
	197	1	263.358	1.18	137	0	1
##		0	329.000	3.00	142	0	0
			255.000	1.10			
##		0	277.000		136	0	0
	205	0		0.70	137	1	1
	280	0	451.000	1.30	136	0	0
	270	0	222.000	1.00	132	1	0
##		0	263.358	1.50	138	0	0
	210	1	268.000	0.80	130	0	0
	265	0	147.000	1.20	141	1	0
	103	0	149.000	1.10	144	1	1
##		1	324.000	0.90	140	0	0
##		0	228.000	3.50	134	1	0
	261	0	203.000	1.00	138	1	0
##		1	368.000	4.00	131	1	1
	212	1	147.000	0.80	140	1	1
	153	1	184.000	0.90	134	1	1
	288	0	543.000	1.00	132	0	0
##		1	263.358	1.83	134	1	1
	188	0	365.000	2.10	144	0	0
	174	0	189.000	0.80	139	1	0
##		1	76.000	1.10	137	1	0
	239	0	257.000	1.00	136	0	0
	156	0	194.000	1.70	140	1	0
	152	1	244.000	0.90	139	1	0
##		1	218.000	1.00	134	1	0
##		0	406.000	0.70	140	1	0
	211	1	389.000	1.00	136	1	1
	220	1	371.000	0.70	140	0	0
	282	0	51.000	2.70	136	1	1
	108	0	226.000	0.90	138	1	0
	17	0	262.000	0.90	140	1	0
##		0	497.000	1.83	135	1	0
	268	0	133.000	1.70	140	1	0
##		1	263.358	1.83	134	0	0
	100	0	221.000	1.10	140	0	0
	297	0	742.000	0.80	138	0	0
	207	0	226.000	0.80	141	0	0
	298	0	140.000	1.40	140	1	1
##		0	149.000	1.30	137	1	1
	159	0	235.000	1.30	134	1	0
##		1	153.000	1.20	136	0	1
	296	0	270.000	1.20	139	0	0
##		0	263.358	6.80	146	0	0
##		1	265.000	1.90	130	1	0
##	221	0	263.358	1.83	134	1	0

## 1	87	0	153.000	0.60	134	0	0
## 9		1	318.000	1.20	137		0
## 9		1	219.000	1.00	141		0
## 7		0	243.000	1.00	132	1	1
## 4		0	194.000	1.10	142		0
## 7		0	122.000	0.90	139	1	1
## 1		0	132.000	1.00	140		0
## 2		0	189.000	0.70	140		0
## 2		0	263.358	1.10	138	1	1
## 2		0	255.000	0.90	137	1	0
## 2		1	263.358	1.60	130	1	1
## 6	34	0	385.000	1.00	145	1	0
## 2	281	0	241.000	1.00	137	1	0
## 2	235	0	305.000	1.10	137	1	1
## 1	.47	0	218.000	0.70	136	1	1
## 1	.24	1	127.000	0.90	145	0	0
## 3	33	1	319.000	1.00	128	0	0
## 2	276	1	422.000	0.80	137	0	0
## 1	.75	1	281.000	0.90	137	1	1
## 1	.49	0	225.000	1.83	134	1	0
## 4	17	1	271.000	0.90	130	1	0
## 2	266	0	362.000	0.90	140	1	1
## 1	.37	0	172.000	0.90	137	0	0
## 2	267	0	263.358	1.83	134	1	1
## 5	57	0	223.000	2.70	138	1	1
## 1	.02	1	263.358	1.18	137	1	0
## 2		0	283.000	0.70	133	0	0
## 2	208	0	186.000	0.90	136	1	0
## 4		1	140.000	4.40	133	1	0
## 1		0	263.358	1.10	134	1	0
## 2		0	185.000	1.10	134	1	1
## 3		0	226.000	1.00	134		0
## 6		0	274.000	1.00	140		1
## 7		1	317.000	0.80	140		1
## 1		0	329.000	0.90	132		0
## 5		1	166.000	1.00	138		0
## 2		0	390.000	0.90	144		0
		0	174.000	0.80	139	1	1
## 8		0	293.000	1.70	136		0
		0	277.000	1.40	136		0
		0	301.000	1.00	142	1	1
## 5		0	216.000	0.60	138	1	1
		0	246.000	2.40	135		0
	244	1	220.000	0.90	141		0
## 5		0	327.000	2.70	116		0
		0	221.000	1.00	136		0
## 2		0	263.358	1.10	136		0
	.29	1	267.000	0.70	136	1	1
	.05	1	286.000	2.30	143		0
	.34	1	295.000	1.30	145		1
			263.358			1	
		0		1.80	113	1 0	1
## 2		0	208.000	1.00	140		0
## 2		0	279.000	1.70	140		0
## 2	35	1	263.358	1.00	139	1	0

##	84	1	172.000	1.80	133	1	0
##	299	0	395.000	1.60	136	1	1
##	112	0	228.000	1.20	135	1	1
##	18	0	166.000	0.80	127	1	0
##	136	0	263.358	1.18	137	1	0
##	158	0	262.000	1.00	136	1	1
##	195	1	126.000	1.60	135	1	0
##	28	1	284.000	1.30	136	1	1
##	250	0	223.000	1.20	130	0	0
##	180	0	279.000	0.70	140	1	1
##	12	1	253.000	0.90	140	1	1
##	75	0	266.000	1.20	134	1	1
	161	1	242.000	1.20	134	1	0
	272	0	221.000	0.90	134	0	0
##	140	1	211.000	0.80	138	1	0
##	198	0	259.000	0.80	138	0	0
##	31	1	263.358	1.83	134	1	0
##		1	216.000	2.30	131	0	0
	173	1	176.000	1.10	145	1	1
	181	0	303.000	0.90	136	1	0
	107	0	263.000	1.30	137	1	0
	209	0	283.000	1.00	141	0	0
	121	1	210.000	1.50	135	1	1
##		1	276.000	1.30	137	0	0
##		0	255.000	2.20	132	0	1
	128	0	300.000	0.80	137	0	0
	204	1	212.000	3.50	136	1	1
##		0	47.000	1.30	136	1	0
##		0	162.000	1.30	129	1	1
	177	0	105.000	1.00	135	1	1
##		1	302.000	1.00	141	0	0
##		1	204.000	2.10	132	1	1
##		0	254.000	1.10	139	1	1
	241	1	533.000	1.30	139	0	0
	127	1	271.000	2.10	124	0	0
	258	1	253.000	1.00	139	1	0
	247	0	314.000	1.10	138	1	0
	256	1	334.000	1.00	142	1	1
	172	0	325.000	0.90	140	1	1
	123	0	228.000	0.75	140	0	0
	283	0	215.000	3.80	128	1	1
	231	0	62.000	1.70	127	0	0
	148	0	242.000	1.00	137	1	0
	186	0	389.000	1.50	136	1	0
	183	0	265.000	1.20	136	1	1
##		0	213.000	1.30	136	0	0
	279	0	232.000	0.70	136	0	0
	234	0	330.000	1.40	137	1	1
	66	0	119.000	2.90	127	1	1
	93	0	263.358	1.18	137	0	0
##		0	305.000	2.30	137	1	0
	146	0	266.000	0.70	141	1	1
	192	0	309.000	1.50	135	0	0
	245	0	264.000	1.80	134	1	0
		-					-

##	46	0	310.000	1.90	135	1	1
##	27	1	196.000	1.00	138	0	0
##	20	0	87.000	1.90	121	0	0
	110	0	850.000	1.30	142	1	1
	109	0	304.000	1.10	133	1	1
##		1	254.000	1.30	136	1	1
	131	1	249.000	0.70	138	1	1
	165	0	334.000				
				1.10	139	1	0
	160	1	362.000	1.10	139	1	1
	130	0	227.000	3.40	145	1	0
	257	0	294.000	1.70	139	0	0
	224	0	130.000	0.80	134	1	0
	168	0	70.000	2.40	134	1	0
	218	1	151.000	9.00	137	0	0
##	71	0	374.000	0.80	140	1	1
##	89	1	235.000	0.70	139	1	0
##	227	0	189.000	1.30	132	1	1
##	125	0	217.000	3.70	134	1	0
##	194	0	160.000	1.18	142	1	1
##	213	0	481.000	1.40	138	1	1
	166	1	192.000	1.30	135	0	0
	199	1	279.000	1.00	134	0	0
	294	0	179.000	0.90	136	1	1
##		0	244.000	1.20	142	0	0
##		0	153.000	5.80	134	1	0
	133	1	263.358	1.18	137	0	0
			507.000				
	118	0		3.20	138	0	0
##		0	368.000	0.80	135	1	0
##		0	136.000	1.10	137	1	0
##		0	210.000	1.70	135	1	0
##		0	149.000	1.90	144	1	1
##		0	395.000	1.00	140	0	0
	176	0	337.000	1.00	138	1	1
##	151	1	235.000	2.50	135	0	0
	115	1	328.000	1.20	126	1	0
##	76	0	204.000	0.70	139	1	1
##	182	0	221.000	1.00	136	1	1
##	286	0	336.000	1.20	135	1	0
##	92	0	297.000	1.00	133	1	0
##	32	0	360.000	3.00	132	1	0
##	38	1	321.000	1.00	145	0	0
##	120	0	263.358	1.83	134	0	0
##	116	0	164.000	1.00	139	0	0
	242	0	249.000	1.30	136	1	1
	190	1	275.000	0.90	140	0	0
	223	0	365.000	1.10	139	1	1
	201	0	73.000	0.70	137	1	0
	10	1	388.000	9.40	133	1	1
	167	0	220.000	0.70	133	1	
							1
	228	1	141.000	1.00	140	1	
##		1	304.000	0.80	140	1	0
	135	0	231.000	1.18	137	1	1
##		1	237.000	1.00	140	0	0
##	15	1	427.000	1.00	138	0	0

##	101			0	298.000	1.10	141	1	0
##	292			0	133.000	1.40	139	1	0
##	132			0	250.000	6.10	131	1	0
	196			0	223.000	1.80	145	1	0
##			DEATH_EVENT						
	111	90	1						
##		40	1						
	229	207	0						
	259	230	0						
##		74	0						
	170	145	0						
	138 226	108 205	0						
	236	209	0						
	230	209	0						
	206	187	0						
	285	250	0						
	185	170	1						
	197	185	0						
##		30	1						
##		85	0						
##	205	187	0						
##	280	246	0						
##	270	244	0						
##	9	10	1						
	210	187	0						
	265	237	0						
	103	87	0						
##		74	0						
##		30	1						
	261	233	0						
##		10	1						
	212	192	0						
	153 288	118	0						
##		250 31	1						
	188	172	1						
	174	146	0						
	14	11	1						
	239	210	0						
	156	120	0						
##	152	117	0						
##	44	33	0						
##	86	79	0						
##	211	188	0						
	220	197	0						
	282	250	0						
	108	88	0						
##		14	1						
##		67	1						
	268	244	0						
##		23	1						
	100 297	86 278	0						
##	291	210	U						

##	207	187	0
##	298	280	0
##	74	72	0
##	159	121	0
##	42	32	1
##	296	271	0
##	53	43	1
##	1	4	1
##	221	198	1
##	187	172	1
##	99	85	0
##	96	83	0
##	73	72	1
##	45	33	1
##	72	71	0
##	178	147	0
##	274	245	0
##	284	250	0
##	233	209	0
##	271	244	0
##	64	61	1
##	281	247	0
##	235	209	0
##	147	112	0
##	124 33	95 28	0
## ##	276	245	0
##	175	146	0
##	149	113	1
##	47	38	1
##	266	240	0
##	137	107	0
##	267	241	1
##	57	54	0
##	102	87	0
##	262	233	0
##	208	187	0
##	49	41	1
##	141	109	1
##	232	208	0
##	37	30	1
##	68	65	1
##	77	74	0
##	143	109	0
##	51	43	1
##	290	256	0
##	162	121	0
##	81	75	0
##	157	120	0
##	249	214	0
##	58 248	54	0
## ##	248244	214 213	0
##	5	213 8	1
##	J	0	1

##	139	108	0
##	2	6	1
##	129	104	0
##	105	87	0
##	134	107	0
##	200	186	0
##	264	237	0
##	287	250	0
##	255	215	0
##	84	78	0
##	299	285	0
##	112	90	0
##	18	14	1
##	136	107	0
##	158	120	0
##	195	180	1
##	28	26	1
##	250	214	0
##	180	147	0
##	12 75	10 73	1
## ##	75 161	121	0
##	272	244	0
##	140	108	0
##	198	186	0
##	31	27	1
##	62	60	1
##	173	146	0
##	181	148	0
##	107	88	0
##	209	187	0
##	121	95	0
##	21	16	0
##	55	45	1
##	128	104	0
##	204	187	0
##	16	13	1
##	3	7	1
##	177	147	0
##	85	78	1
##	6	8	1
##	63	60	0
##	241	212	0
##	127	100	1
##	258	230	0
##	247	214	1
##	256	216	0
##	172	146	0
##	123	95	0
##	283	250	0
##	231	207	1
##	148	113	0
##	186	171	1
##	183	154	1

##	67	65	1
##	279	246	C
##	234	209	C
##	66	64	1
##	93	82	C
##	39	30	C
##	146	112	C
##	192	174	C
##	245	213	0
##	46	35	1
##	27 20	24 15	1 1
##	110	88	C
##	109	88	C
##	60	59	1
##	131	106	C
##	165	129	1
##	160	121	C
##	130	105	C
##	257	220	C
##	224	201	C
##	168	135	1
##	218	196	1
##	71	68	C
##	89	79	C
##	227	205	C
##	125	96	1
##	194	180	C
##	213	192	C
##	166	130	1
##	199	186	C
##	294	270	C
##	69	66	1
##	29	26	1
## ##	133	107	C
##	118 24	94 22	C
##	13	11	1
##	94	82	1
##	26	23	1
##	50	42	1
##	176	146	C
##	151	115	1
##	115	91	C
##	76	73	1
##	182	150	1
##	286	250	C
##	92	80	C
##	32	28	1
##	38	30	1
##	120	95	1
##	116	91	C
##	242	212	C
##	190	174	C

##	223	201	0
##	201	186	0
##	10	10	1
##	167	134	0
##	228	206	0
##	88	79	0
##	135	107	0
##	19	15	1
##	15	12	0
##	101	87	0
##	292	258	0
##	132	107	0
##	196	180	1

test_data

##			${\tt anaemia}$	${\tt creatinine_phosphokinase}$	${\tt diabetes}$	${\tt ejection_fraction}$
##	4	50	1	111	0	20
##	7	75	1	246	0	15
##	8	60	1	315	1	60
##	22	65	1	128	1	30
##	23	68	1	220	0	35
	30	82	0	70	1	30
##	34	50	1	159	1	30
	35	65	0	94	1	50
##	43	70	0	571	1	45
	52	53	1	91	0	20
	54	70	1	69	1	50
	56	95	1	371	0	30
	59	49	0	789	0	20
	61	45	0	7702	1	25
	65	45	0	582	0	80
	79	75	1	203	1	38
	82	67	0	582	0	50
	83	60	1	76	1	25
	87	55	0	47	0	35
	90	57	1	115	0	25
	91	70	0	66	1	45
	95	58	0	144	1	38
	97	63	1	514	1	25
	104	42	0	5209	0	30
	106	72	1	328	0	30
	113	50	0	369	1	25
	114	70	1	143	0	60
##	117	60	1	96	1	60
##	119	65	1	113	1	60
	122	66	1	68	1	38
	126	43	1	358	0	50
	142	46	1	291	0	35
	144	61	1	84	0	40
##	145	72	1	943	0	25
	150	60	0	2261	0	35
	154	50	0	1846	1	35
##	155	65	1	335	0	35

##	163	63	1		582	0		40	
##	164	50	1		2334	1		35	
##	169	65	0		582	1		40	
##	171	51	1		582	1		35	
##	179	63	1		122	1		60	
##	184	75	0		99	0		38	
##	189	60	1		151	1		40	
##	191	80	0		582	1		35	
##	193	50	1		121	1		40	
##	202	45	0		308	1		60	
##	203	70	0		97	0		60	
##	214	48	1		131	1		30	
##	215	65	1		135	0		35	
##	216	73	0		582	0		35	
##	217	70	0		1202	0		50	
##	219	68	1		1021	1		35	
##	222	65	0		118	0		50	
##	225	58	0		582	1		25	
##	237	75	0		119	0		50	
##	238	70	0		232	0		30	
##	240	55	1		180	0		45	
	243	40	0		90	0		35	
	246	61	1		80	1		38	
	251	50	0		2522	0		30	
	252	55	0		572	1		35	
	253	50	0		245	0		45	
	254	70	0		88	1		35	
	260	53	0		56	0		50	
	263	65	1		258	1		25	
	269	45	0		582	1		38	
	273	67	0		213	0		38	
	275	60	1		257	1		30	
	277	70	0		618	0		35	
	278	70	0		582	1		38	
	289	65 45	0		892	1		35	
	291293	45 52	0		615 190	1		55 38	
	295	62	0		61	1 1		38	
##	290		blood_pressure	nlatelets			serum sodium		amokina
##	Δ	111811_	0	210.000	ser um_c	1.90	137	1	0
##			0	127.000		1.20	137	1	0
##			0	454.000		1.10	131	1	1
##			1	297.000		1.60	136	0	0
##			1	289.000		0.90	140	1	1
##			0	200.000		1.20	132	1	1
##			0	302.000		1.20	138	0	0
##			1	188.000		1.00	140	1	0
##	43		1	185.000		1.20	139	1	1
##	52		1	418.000		1.40	139	0	0
##	54		1	351.000		1.00	134	0	0
##	56		0	461.000		2.00	132	1	0
##			1	319.000		1.10	136	1	1
##			1	390.000		1.00	139	1	0
##	65		0	263.358		1.18	137	0	0

##	79	1	283.000	0.60	131	1	1
	82	0	263.358	1.18	137	1	1
	83	0	196.000	2.50	132		0
	87	1	173.000	1.10	137	1	0
	90	1	181.000	1.10	144	1	0
	91	0	249.000	0.80	136	1	1
	95	1	327.000	0.70	142		0
	97	1	254.000	1.30	134	1	0
	104	0	226.000	1.00	140	1	1
	106	1	621.000	1.70	138	0	1
	113	0	252.000	1.60	136	1	0
			351.000				
	114 117	0	271.000	1.30	137	0	0
		1		0.70	136	0	0
	119	1	203.000	0.90	140	0	0
	122	1	162.000	1.00	136	0	0
	126	0	237.000	1.30	135	0	0
	142	0	348.000	0.90	140	0	0
	144	1	229.000	0.90	141	0	0
	145	1	338.000	1.70	139	1	1
	150	1	228.000	0.90	136		0
	154	0	263.358	1.18	137	1	1
	155	1	235.000	0.80	136	0	0
	163	0	448.000	0.90	137	1	1
	164	0	75.000	0.90	142	0	0
	169	0	270.000	1.00	138	0	0
	171	0	263.358	1.50	136	1	1
	179	0	267.000	1.20	145	1	0
	184	1	224.000	2.50	134	1	0
	189	1	201.000	1.00	136	0	0
	191	0	350.000	2.10	134	1	0
	193	0	260.000	0.70	130	1	0
	202	1	377.000	1.00	136	1	0
	203	1	220.000	0.90	138	1	0
	214	1	244.000	1.60	130	0	0
	215	1	290.000	0.80	134	1	0
	216	1	203.000	1.30	134	1	0
	217	1	358.000	0.90	141	0	0
	219	0	271.000	1.10	134	1	0
	222	0	194.000	1.10	145	1	1
	225	0	504.000	1.00	138	1	0
##	237	1	248.000	1.10	148	1	0
##	238	0	173.000	1.20	132	1	0
##	240	0	263.358	1.18	137	1	1
##	243	0	255.000	1.10	136	1	1
##	246	0	282.000	1.40	137	1	0
##	251	1	404.000	0.50	139	0	0
##	252	0	231.000	0.80	143	0	0
##	253	1	274.000	1.00	133	1	0
##	254	1	236.000	1.20	132	0	0
##	260	0	308.000	0.70	135	1	1
##	263	0	198.000	1.40	129	1	0
##	269	0	302.000	0.90	140		0
	273	0	215.000	1.20	133	0	0
##	275	0	150.000	1.00	137	1	1

##	277			0	327.000	1.10	142	0	0
##	278			0	25.100	1.10	140	1	0
##	289			0	263.358	1.10	142	0	0
	291			0	222.000	0.80	141	0	0
	293			0	382.000	1.00	140	1	1
	295			1	155.000	1.10	143	1	1
##			DEATH_EVENT						
##		7	1						
##		10	1						
##		10	1						
	22	20	1						
	23	20	1						
	30	26	1						
	34	29	0						
	35 43	29 33	1						
	43 52	43	1 1						
	52 54	43	1						
	56	50	1						
	59	55	1						
	61	60	1						
	65	63	0						
	79	74	0						
	82	76	0						
	83	77	1						
	87	79	0						
	90	79	0						
	91	80	0						
	95	83	0						
##	97	83	0						
##	104	87	0						
	106	88	1						
##	113	90	0						
	114	90	1						
	117	94	0						
	119	94	0						
	122	95	0						
	126	97	0						
	142	109	0						
	144	110	0						
	145	111	1						
	150	115	0						
	154	119	0						
	155 163	120 123	0						
	164	126	1						
	169	140	0						
	171	145	0						
	179	147	0						
	184	162	1						
	189	172	0						
	191	174	0						
	193	175	0						
	202	186	0						

```
## 203
        186
## 214
        193
                       1
## 215
        194
                       0
## 216
                       0
        195
## 217
        196
                       0
## 219
                       0
        197
## 222
        200
## 225
                       0
        205
## 237
        209
                       0
## 238
                       0
        210
## 240
        211
                       0
                       0
## 243
        212
## 246
        213
                       0
## 251
                       0
        214
## 252
        215
                       0
## 253
        215
                       0
## 254
        215
                       0
## 260
        231
                       0
## 263
        235
                       1
## 269
        244
                       0
## 273
        245
                       0
## 275
        245
                       0
## 277
        245
                       0
## 278
        246
                       0
## 289
                       0
        256
## 291
        257
                       0
## 293
        258
                       0
## 295
        270
                       0
```

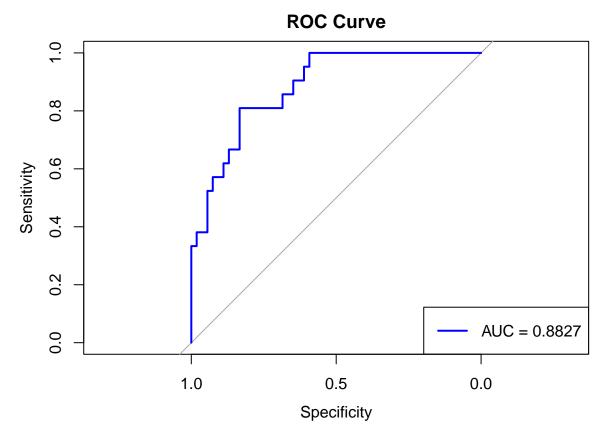
Model development (logistic)

```
logistic_model = glm(DEATH_EVENT~., data=train_data, family="binomial")
summary(logistic_model)
```

```
##
## Call:
## glm(formula = DEATH_EVENT ~ ., family = "binomial", data = train_data)
##
## Coefficients:
##
                            Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                           7.9632799 6.0737111
                                                 1.311 0.1898
                                                 2.516
## age
                           0.0434821 0.0172850
                                                        0.0119 *
                          -0.3808572
                                     0.4272709
                                               -0.891
## anaemia
                                                        0.3727
## creatinine_phosphokinase 0.0001882 0.0002380
                                                 0.791
                                                        0.4292
## diabetes
                           0.2101722 0.4112369
                                                 0.511
                                                        0.6093
## ejection_fraction
                          -0.0825491 0.0190516 -4.333 1.47e-05 ***
## high_blood_pressure
                          -0.2561613   0.4318960   -0.593
                                                        0.5531
                          -0.0031861 0.0023648
                                               -1.347
## platelets
                                                        0.1779
## serum_creatinine
                           0.6122125 0.1903677
                                                 3.216
                                                        0.0013 **
## serum sodium
                          -0.0416713 0.0432340 -0.964
                                                        0.3351
                          -0.4533968 0.4677745 -0.969
                                                        0.3324
## sex
## smoking
                          -0.2713557
                                     0.4774305 - 0.568
                                                        0.5698
## time
```

```
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 285.62 on 223 degrees of freedom
## Residual deviance: 166.87 on 211 degrees of freedom
## AIC: 192.87
##
## Number of Fisher Scoring iterations: 6
Applying this model to test data
pred_prob <- predict(logistic_model, newdata = test_data, type = "response")</pre>
#Setting threshold probability to 0.5
threshold <- 0.5
test_predictions <- ifelse(pred_prob > threshold, 1, 0)
test_predictions
##
               22 23 30 34
                                35
                                    43
                                        52
                                            54
                                                56
                                                    59
                                                         61
                                                             65
                                                                 79
                                                                     82
                                                                         83
                                                                             87
                                                                                  90
##
     1
                 1
                                 0
                                          1
                                              0
                                                              0
                                                                  0
                                                                      0
                                                                           1
                     1
                         1
                             1
                                      1
                                                  1
                                                      1
                                                          1
       95 97 104 106 113 114 117 119 122 126 142 144 145 150 154 155 163 164 169
           1
                 0
                     0
                         1
                             0
                                 0
                                      0
                                          0
                                              0
                                                  0
                                                      0
                                                          0
                                                              0
                                                                  0
                                                                      0
                                                                           0
## 171 179 184 189 191 193 202 203 214 215 216 217 219 222 225 237 238 240 243 246
                 0
                         0
                             0
                                 0
                                      0
                                          0
                                              0
                                                      0
                                                          0
                                                              0
                                                                  0
                                                                      0
                                                                          0
             0
                     0
                                                  0
## 251 252 253 254 260 263 269 273 275 277 278 289 291 293 295
                 0
                         0
                             0
                                 0
                                     0
                                        0
                                             0
                                                  0
                     0
Model evaluation
library(caret)
## Loading required package: lattice
# Create a confusion matrix
conf_matrix <- confusionMatrix(factor(test_predictions), factor(test_data$DEATH_EVENT))</pre>
# Display the confusion matrix and accuracy
conf_matrix
## Confusion Matrix and Statistics
##
             Reference
##
## Prediction 0 1
            0 51 10
##
##
            1 3 11
##
##
                  Accuracy : 0.8267
##
                    95% CI: (0.7219, 0.9043)
##
       No Information Rate: 0.72
       P-Value [Acc > NIR] : 0.02306
##
```

```
##
##
                     Kappa: 0.5214
##
   Mcnemar's Test P-Value: 0.09609
##
##
               Sensitivity: 0.9444
##
##
               Specificity: 0.5238
            Pos Pred Value: 0.8361
##
##
            Neg Pred Value: 0.7857
                Prevalence: 0.7200
##
##
            Detection Rate: 0.6800
##
      Detection Prevalence: 0.8133
##
         Balanced Accuracy: 0.7341
##
##
          'Positive' Class: 0
##
print(paste("Accuracy:", conf_matrix$overall["Accuracy"]))
## [1] "Accuracy: 0.8266666666667"
#Here we obtained an accuracy of 0.85 which is good for predicting survival of patients.
Drawing ROC curve
library(pROC)
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
## The following objects are masked from 'package:stats':
##
##
       cov, smooth, var
# Create ROC curve
roc_curve <- roc(test_data$DEATH_EVENT, pred_prob)</pre>
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
# Plot ROC curve
plot(roc_curve, main = "ROC Curve", col = "blue", lwd = 2)
legend("bottomright", legend = paste("AUC =", round(auc(roc_curve), 4)), col = "blue", lwd = 2)
```



Again AUC score is 0.8843 which represent good measure of the discriminative ability of a classification model.