

Project on water potability

Explanatory Analysis

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I am took a water potability dataset from kaggle. Data file name is **water_potability.csv**

Context

Access to safe drinking-water is essential to health, a basic human right and a component of effective policy for health protection. This is important as a health and development issue at a national, regional and local level. In some regions, it has been shown that investments in water supply and sanitation can yield a net economic benefit, since the reductions in adverse health effects and health care costs outweigh the costs of undertaking the interventions.

Content

The water_potability.csv file contains water quality metrics for 3276 different water bodies.

pH value: PH is an important parameter in evaluating the acid–base balance of water. It is also the indicator of acidic or alkaline condition of water status. WHO has recommended maximum permissible limit of pH from 6.5 to 8.5. The current investigation ranges were 6.52–6.83 which are in the range of WHO standards.

Hardness: Hardness is mainly caused by calcium and magnesium salts. These salts are dissolved from geologic deposits through which water travels. The length of time water is in contact with hardness producing material helps determine how much hardness there is in raw water. Hardness was originally defined as the capacity of water to precipitate soap caused by Calcium and Magnesium.

Solids (Total dissolved solids - TDS): Water has the ability to dissolve a wide range of inorganic and some organic minerals or salts such as potassium, calcium, sodium, bicarbonates, chlorides, magnesium, sulfates etc. These minerals produced un-wanted taste and diluted color in appearance of water. This is the important parameter for the use of water. The water with high TDS value indicates that water is highly mineralized. Desirable limit for TDS is 500 mg/l and maximum limit is 1000 mg/l which prescribed for drinking purpose.

Chloramines: Chlorine and chloramine are the major disinfectants used in public water systems. Chloramines are most commonly formed when ammonia is added to chlorine to treat drinking water. Chlorine levels up to 4 milligrams per liter (mg/L or 4 parts per million (ppm)) are considered safe in drinking water.

Sulfate: Sulfates are naturally occurring substances that are found in minerals, soil, and rocks. They are present in ambient air, groundwater, plants, and food. The principal commercial use of sulfate is in the chemical industry. Sulfate concentration in seawater is about 2,700 milligrams per liter (mg/L).

It ranges from 3 to 30 mg/L in most freshwater supplies, although much higher concentrations (1000 mg/L) are found in some geographic locations.

Conductivity: Pure water is not a good conductor of electric current rather's a good insulator. Increase in ions concentration enhances the electrical conductivity of water. Generally, the amount of dissolved solids in water determines the electrical conductivity. Electrical conductivity (EC) actually measures the ionic process of a solution that enables it to transmit current. According to WHO standards, EC value should not exceeded 400 $\mu\text{S}/\text{cm}$.

Organic_carbon: Total Organic Carbon (TOC) in source waters comes from decaying natural organic matter (NOM) as well as synthetic sources. TOC is a measure of the total amount of carbon in organic compounds in pure water. According to US EPA < 2 mg/L as TOC in treated / drinking water, and < 4 mg/Lit in source water which is use for treatment.

Trihalomethanes: THMs are chemicals which may be found in water treated with chlorine. The concentration of THMs in drinking water varies according to the level of organic material in the water, the amount of chlorine required to treat the water, and the temperature of the water that is being treated. THM levels up to 80 ppm is considered safe in drinking water.

Turbidity: The turbidity of water depends on the quantity of solid matter present in the suspended state. It is a measure of light emitting properties of water and the test is used to indicate the quality of waste discharge with respect to colloidal matter. The mean turbidity value obtained for Wondo Genet Campus (0.98 NTU) is lower than the WHO recommended value of 5.00 NTU.

Potability: Indicates if water is safe for human consumption where 1 means Potable and 0 means Not potable.

We are doing explonetry analysis on it then we will fit a model on it.

Some basic step we have to follow for analyzing a dataset.this

step 2. Cleaning the dataset

step 1. Loading the dataset

step 3. Visualized the dataset

step 4. Fit a model on it

Step 1:

I load water portability data from my local file.and show 1st nth row of the dataset using `head()`

```
options(warn=-1)
```

```
data =read.csv("C:/Users/SVMY/Downloads/R project/water_potability.csv")
head(data)
```

	ph	Hardness	Solids	Chloramines	Sulfate	Conductivity	Organic_carbon
1	NA	204.8905	20791.32	7.300212	368.5164	564.3087	10.379783
2	3.716080	129.4229	18630.06	6.635246	NA	592.8854	15.180013
3	8.099124	224.2363	19909.54	9.275884	NA	418.6062	16.868637
4	8.316766	214.3734	22018.42	8.059332	356.8861	363.2665	18.436524
5	9.092223	181.1015	17978.99	6.546600	310.1357	398.4108	11.558279
6	5.584087	188.3133	28748.69	7.544869	326.6784	280.4679	8.399735
	Trihalomethanes	Turbidity	Potability				
1	86.99097	2.963135	0				
2	56.32908	4.500656	0				

3	66.42009	3.055934	0
4	100.34167	4.628771	0
5	31.99799	4.075075	0
6	54.91786	2.559708	0

I want to see column names and dimension of our dataset.

```
colnames(data) #columns name

[1] "ph"           "Hardness"      "Solids"        "Chloramines"
[5] "Sulfate"      "Conductivity"  "Organic_carbon" "Trihalomethanes"
[9] "Turbidity"    "Potability"

dim(data) #dimension of data

[1] 3276 10
```

Our dataset contains 3276 rows and 10 columns.

Step 2:

we want to see if there are any null values in the dataset.

```
sum(is.na(data)) #sum of null values if in dataset

[1] 1434
```

It shows that there are 1434 empty cells. Now we have to remove these empty cells from our dataset otherwise it will make trouble to analyze our dataset.

```
data=na.omit(data)
```

```
sum(is.na(data))

[1] 0
```

we are interested in the type of each variable.

```
sapply(data, typeof)

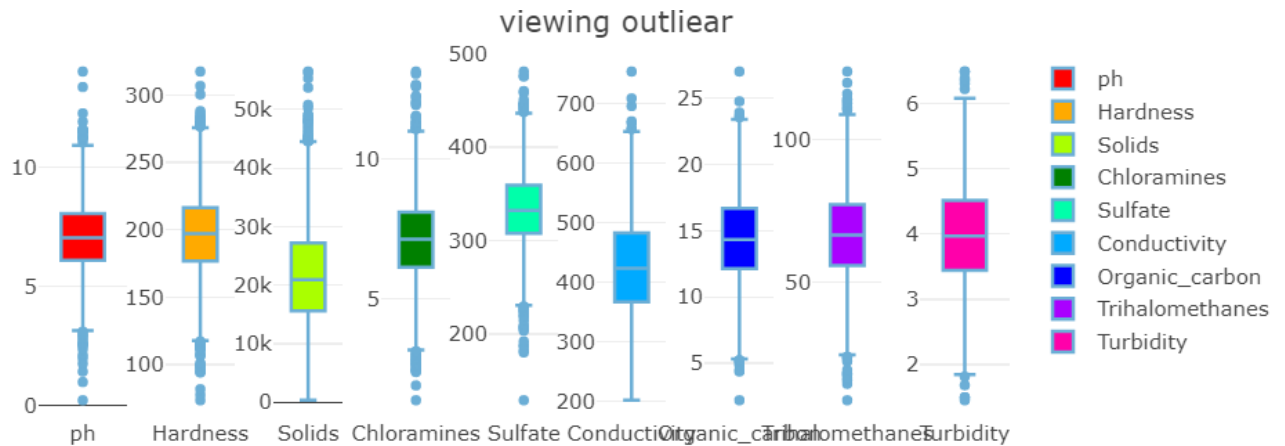
      ph      Hardness      Solids      Chloramines      Sulfate
"double" "double"    "double"    "double"    "double"
Conductivity Organic_carbon Trihalomethanes      Turbidity      Potability
"double"  "double"    "double"    "double"    "integer"
```

here data types are integer and double(float) so we do not need to change anything.

Outlier

```
options(warn=-1)
library(plotly)
library(webshot)
l <- htmltools::tagList()
for(i in 1:9){
  l[[i]]=as.widget(plot_ly(y = data[,i], boxpoints = 'outliers', type = "box", name = colnames(data)[i], margin = list(
    line = list(color = 'rgb(107,174,214)'), fillcolor = palette(rainbow(9))[i]))
```

```
}
fig <- subplot(1[[1]], 1[[2]],1[[3]],1[[4]],1[[5]],1[[6]],1[[7]],1[[8]],1[[9]]) %>% layout(title = 'viewing outleiar')
fig
```



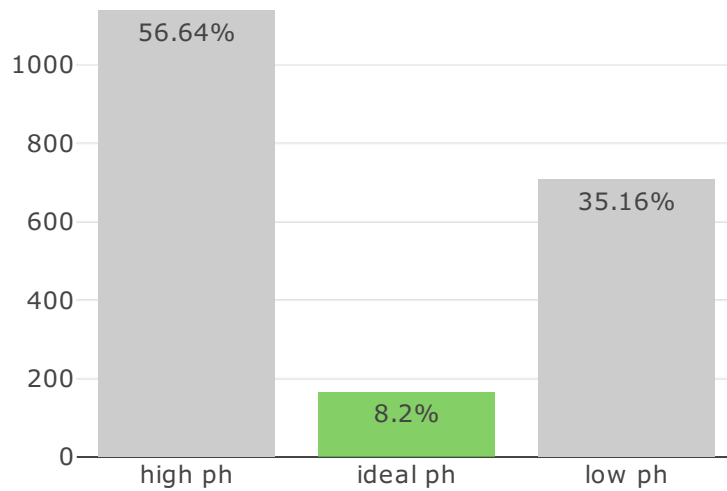
Now visualizing each variables which define water purity.

pH value:

```
library("webshot")
library(plotly)

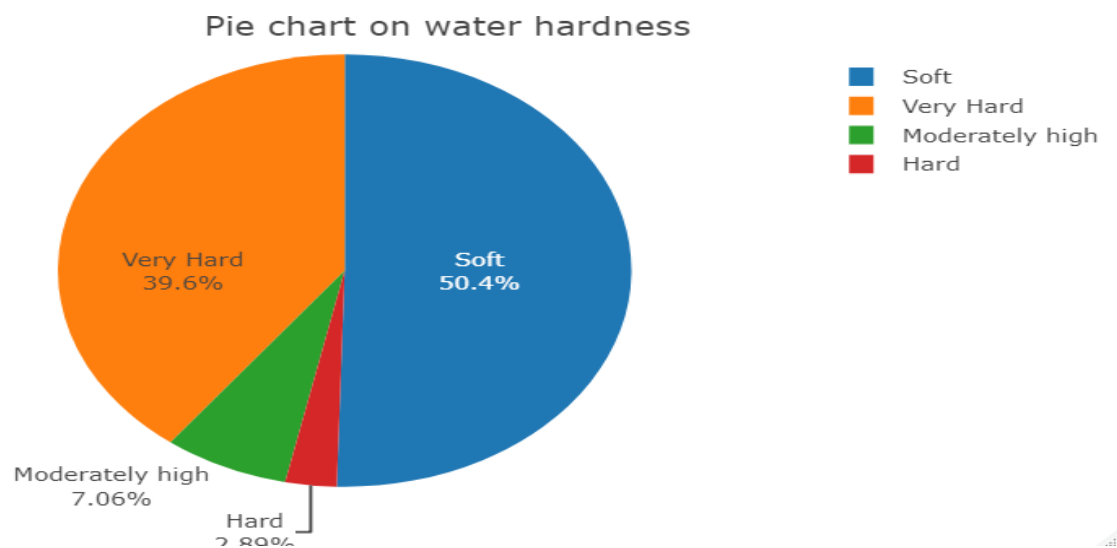
FALSE Loading required package: ggplot2
FALSE
FALSE Attaching package: 'plotly'
FALSE The following object is masked from 'package:ggplot2':
FALSE
FALSE last_plot
FALSE The following object is masked from 'package:stats':
FALSE
FALSE filter
FALSE The following object is masked from 'package:graphics':
FALSE
FALSE layout

x=c(length(which(data$ph<6.52)),length(which(data$ph>6.52 & data$ph<6.83)),length(which(data$ph>6.83)))
txt=c(paste0(round((x[1]/sum(x))*100,2), "%"),paste0(round((x[2]/sum(x))*100,2), "%"),paste0(round((x[3]/sum(x))*100,2), "%"))
fig <- plot_ly(
  x = c("low ph", "ideal ph", "high ph"),
  y = x,
  name = "SF Zoo",
  type = "bar",
  text=txt,
  marker = list(color = c('rgba(204,204,204,1)', 'rgba(103,195,66,0.8)', 'rgba(204,204,204,1)'))
)
fig
```



Hardness:

```
label=c("Soft","Moderately high","Hard","Very Hard")
values=c(length(which(data>0 & data<60)),length(which(data>61 & data<120)),length(which(data>121 & data<180)),length(which(data>181 & data<300)))
fig <- plot_ly(type='pie', labels=label, values=values,
               textinfo='label+percent',
               insidetextorientation='radial')
fig <- fig %>% layout(title = "Pie chart on water hardness")
fig
```



Solids:

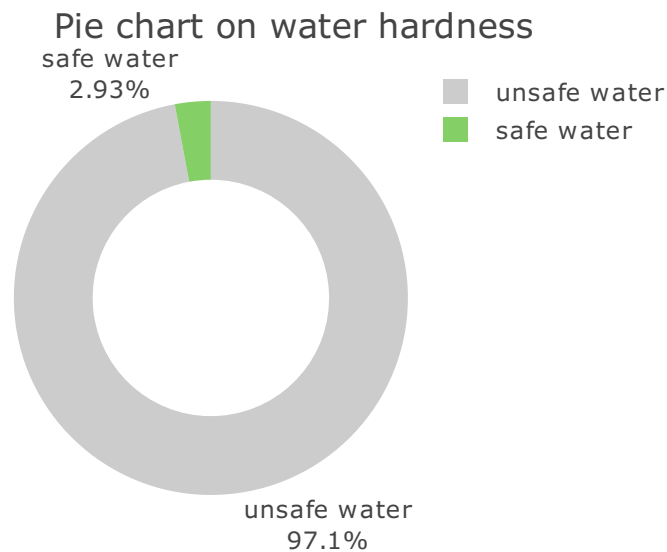
there is only one value which is below 1000 this means that all the water is safe corresponding this solids variable.

```
length(which(data$Solids<1000))
```

```
[1] 1
```

Chloramines:

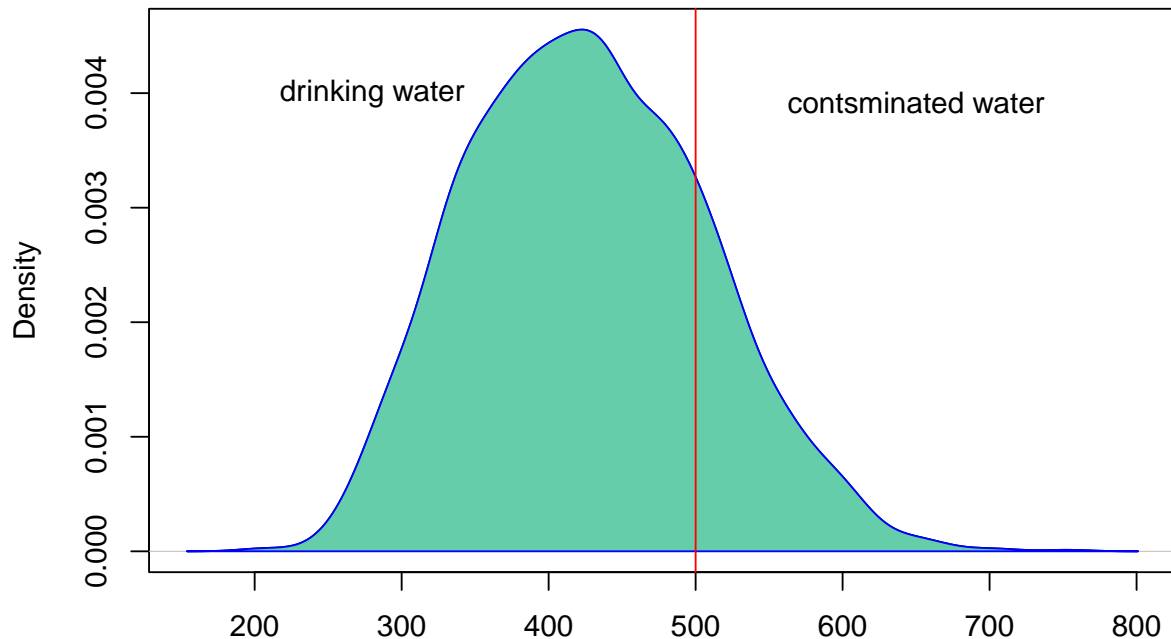
```
label=c("safe water","unsafe water")
values=c(length(which( data$Chloramines<4)),length(which(data$Chloramines>4)))
colors <- c('rgba(103,195,66,0.8)','rgba(204,204,204,1)')
fig <- plot_ly(, labels=label, values=values,
               textinfo='label+percent',marker = list(colors = colors))
fig <- fig %>% add_pie(hole = 0.6)
fig <- fig %>% layout(title = "Pie chart on water hardness")
fig
```

**Sulfate:**

```
plot(density(data$Conductivity), main="frequency density plot")
polygon(density(data$Conductivity), col="aquamarine3", border="blue")
abline(v=500,col="red")
```

```
#locator() #this function is use to locate point in the diagram
text(280, 0.003999010, expression("drinking water"))
text(650.0549, 0.00392039, expression("contsminated water"))
```

frequency density plot



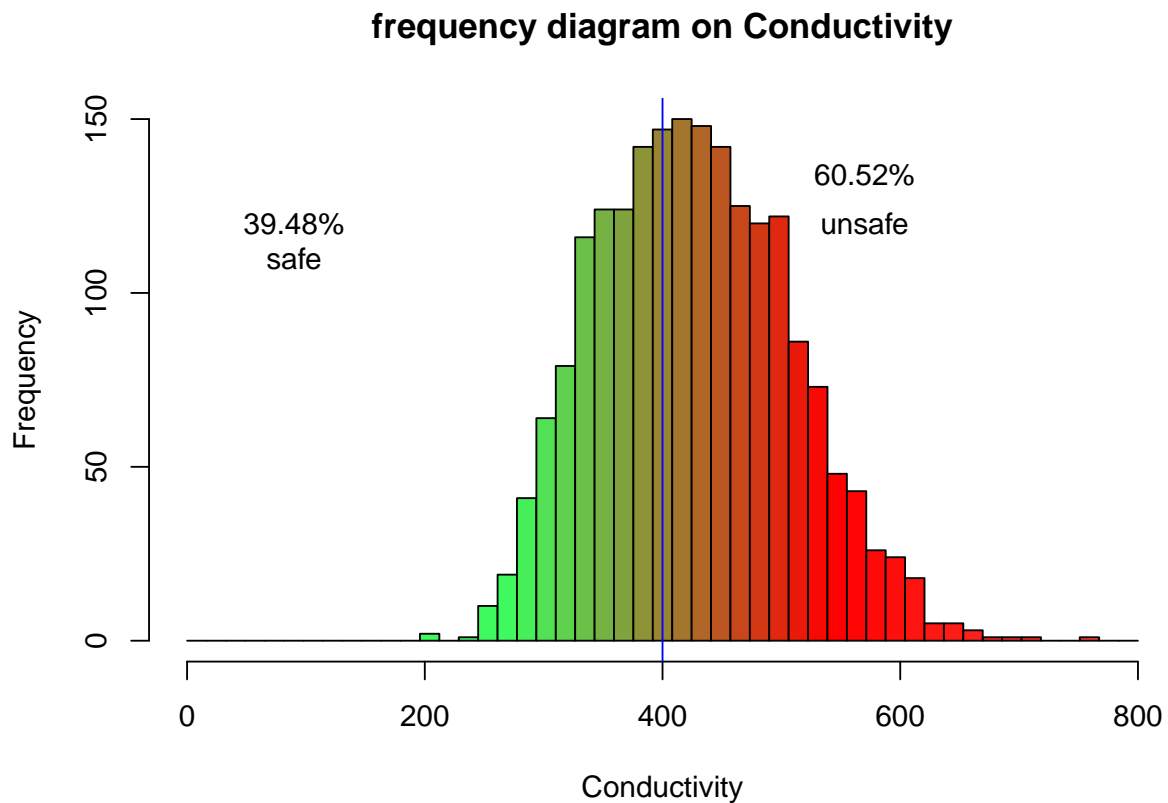
N = 2011 Bandwidth = 15.87

Area right side of red line is dirty water, we can't drink this water & area left side of red line is drinking water.

Conductivity:

```
x=c(length(which(data$Conductivity<400)),length(which(data$Conductivity>400)))
txt=c(paste0(round((x[1]/sum(x))*100,2), "%"),paste0(round((x[2]/sum(x))*100,2), "%"))

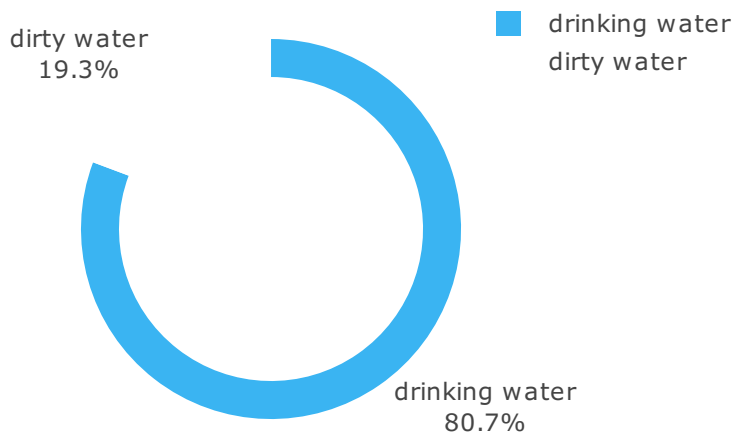
colfunc <- colorRampPalette(c("#3FFA5E", "#3FFA5E", "red", "#FA4B3F"))
hist(data$Conductivity, breaks = seq(from=0, to=800, length=50), col=colfunc(50), main = "frequency diagram of conductivity")
abline(v=400, col="blue")
#locator()
text(90, 120, txt[1])
text(570, 134, txt[2])
text(90, 110, "safe")
text(570, 120, "unsafe")
```



Trihalomethanes:

```
label1=c("drinking water","dirty water")
values1=c(length(which( data$Trihalomethanes<80)),length(which(data$Trihalomethanes>80)))
colors <- c('#3AB4F2','#FFFFFF')
fig <- plot_ly( labels=label1, values=values1,
               textinfo='label+percent',
               insidetextorientation='radial',marker = list(colors = colors))
fig <- fig %>% add_pie(hole = 0.8)
fig <- fig %>% layout(title = "Pie chart on Trihalomethanes")
fig
```


Pie chart on Trihalomethanes



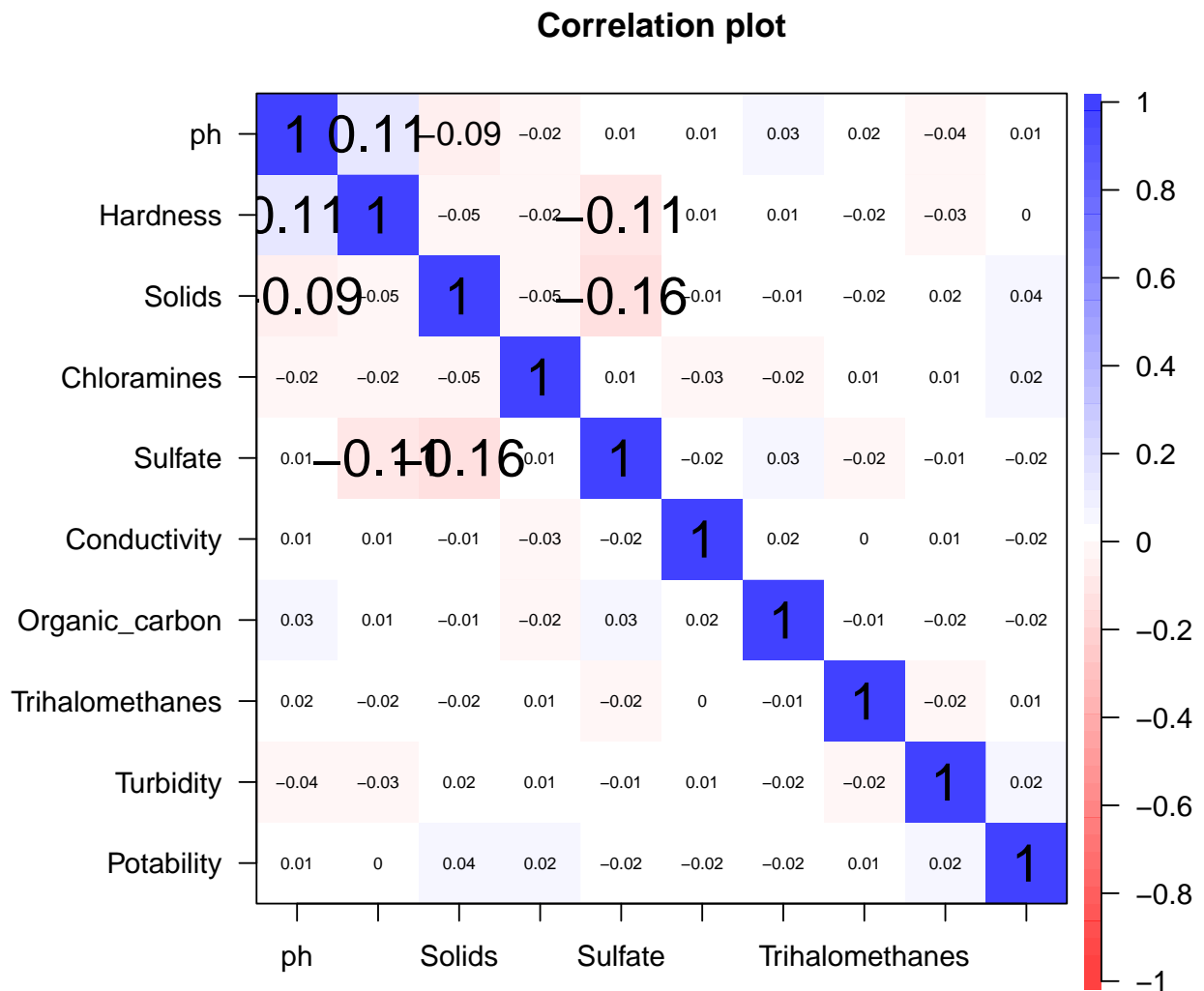
Correlation plot

```
library(psych)
```

```
Attaching package: 'psych'
```

```
The following objects are masked from 'package:ggplot2':  
  %+%, alpha
```

```
corPlot(data, cex = 1.2)
```



this correlation plot is not usefull for us.

Fitting a model:

Splitting dataset into training and test dataset and our spliting ratio 70-30,then extract indepnedent(x) and dependent variables(y)

```
#split dataset
library(caTools)
train = sample.split(data, SplitRatio = .70)
train <- subset( data, train==TRUE )[,1:10]
test <- subset( data, train==FALSE )[,1:10]
trainst=scale(train)
testst=scale(test)
```

```
x=trainst[:,1:9]  
y=trainst[:,10]
```

here we can't perform linear model because our response is in binary format ,so we will use GLM(gerenalized linear model).