

20F-0292

BCS 5E

PROBABILITY AND
STATISTICS PROJECT

- **Dataset Name: Crop Recommendation**
- **Dataset Link:** <https://www.kaggle.com/datasets/atharvaingle/crop-recommendation-dataset>
- **Tools Used: R Programming Language, RStudio, R Markdown, RShiny, flexdashboard**

CODES:

- 1) Declaration and making of simple dashboard interface

title: "CROP RECOMMENDATION"

output:

flexdashboard::flex_dashboard

runtime: shiny

- 2) Importing our excel file into RMarkdown and Displaying Table in Table section

Table

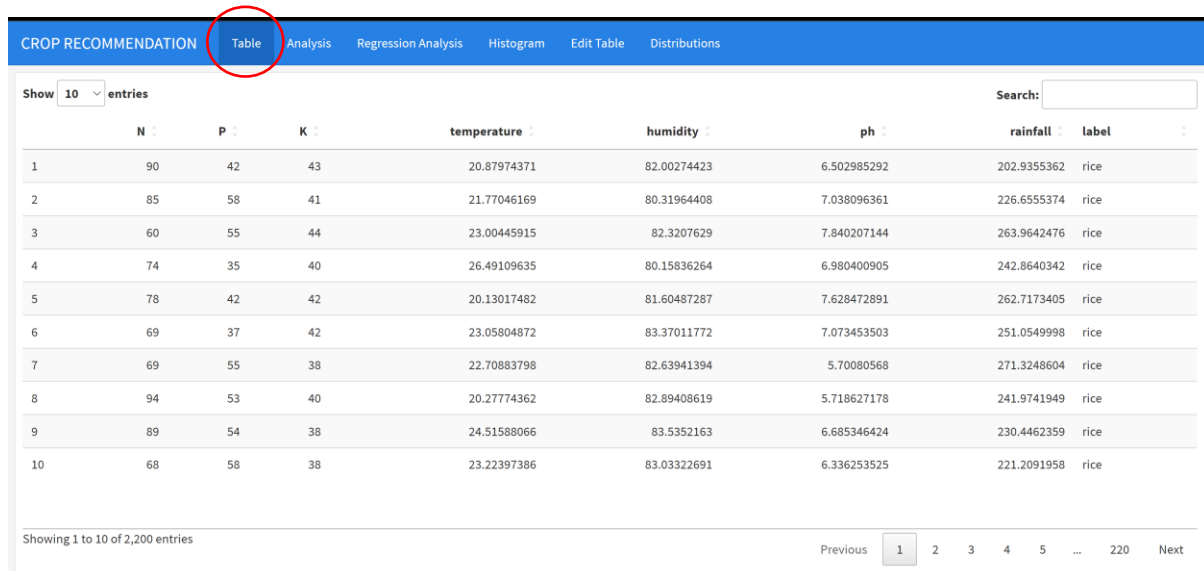
```
```\r
```

```
library(DT)
```

```
startData <- read.csv("crop.csv")
```

```
DT::datatable(startData)
```

```
```\r
```



	N	P	K	temperature	humidity	ph	rainfall	label
1	90	42	43	20.87974371	82.00274423	6.502985292	202.9355362	rice
2	85	58	41	21.77046169	80.31964408	7.038096361	226.6555374	rice
3	60	55	44	23.00445915	82.3207629	7.840207144	263.9642476	rice
4	74	35	40	26.49109635	80.15836264	6.980400905	242.8640342	rice
5	78	42	42	20.13017482	81.60487287	7.628472891	262.7173405	rice
6	69	37	42	23.05804872	83.37011772	7.073453503	251.0549998	rice
7	69	55	38	22.70883798	82.63941394	5.70080568	271.3248604	rice
8	94	53	40	20.27774362	82.89408619	5.718627178	241.9741949	rice
9	89	54	38	24.51588066	83.5352163	6.685346424	230.4462359	rice
10	68	58	38	23.22397386	83.03322691	6.336253525	221.2091958	rice

Figure 1 Table Interface

You can also filter the searches for the ones you best want

3) Doing statistical analysis on whole data in Analysis section

```
# Analysis
```

```
## Column {.sidebar}
```

```
```{r}
```

```
output$summary1 <- renderPrint({
 if(input$datalist=="All"){
 myVec2=c()
 val <- 2
 while(val <= row_num){
 myVec2 <- append(myVec2,mean(t(data_origin[val,])))
 val<-val+1
 }
 summary(myVec2)
 }
 else{
 label <- input$datalist
 rows <- length(t(data_origin[,2]))
 summary(data_origin[2:rows,label])
 }
})
```

```
output$standD <- renderText({
 if(input$datalist=="All"){
 myVec2=c()
 val <- 2
 while(val <= row_num){
 myVec2 <- append(myVec2,mean(t(data_origin[val,])))
 val<-val+1
 }
 c("Standard Deviation: ",sd(myVec2))
 }
```

```

 }
 else{
 label <- input$datalist
 rows <- length(t(data_origin[,2]))
 c("Standard Deviation: ",sd(data_origin[2:rows,label]))
 }
 })

output$IQR <- renderText({
 if(input$datalist=="All"){
 myVec2=c()
 val <- 2
 while(val <= row_num){
 myVec2 <- append(myVec2,mean(t(data_origin[val,])))
 val<-val+1
 }
 c("IQR: ",IQR(myVec2))
 }
 else{
 label <- input$datalist
 rows <- length(t(data_origin[,2]))
 c("IQR: ",IQR(data_origin[2:rows,label]))
 }
})

data <- read.csv("crop.csv")
data_origin <- subset(data, select = -c(8,8))
selectInput("datalist", "Choose a column:", c(rownames(t(data_origin))))
...

Column

...{r}
column(8, verbatimTextOutput("minval"))
column(8, verbatimTextOutput("maxval"))
column(8, verbatimTextOutput("summary1"))
column(8, verbatimTextOutput("standD"))
column(8, verbatimTextOutput("IQR"))
...

```



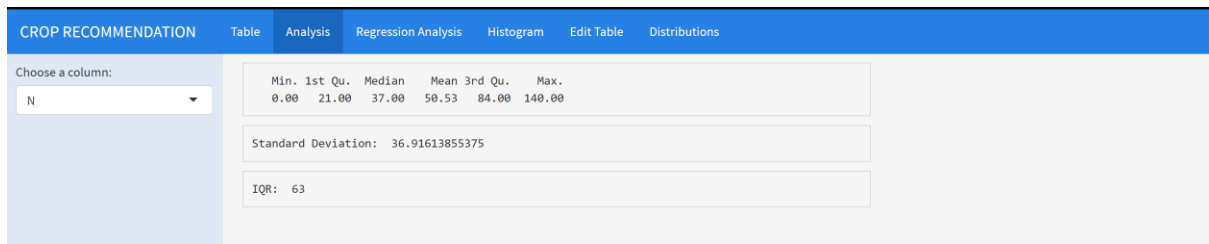


Figure 2 Analytics done on Nitrogen

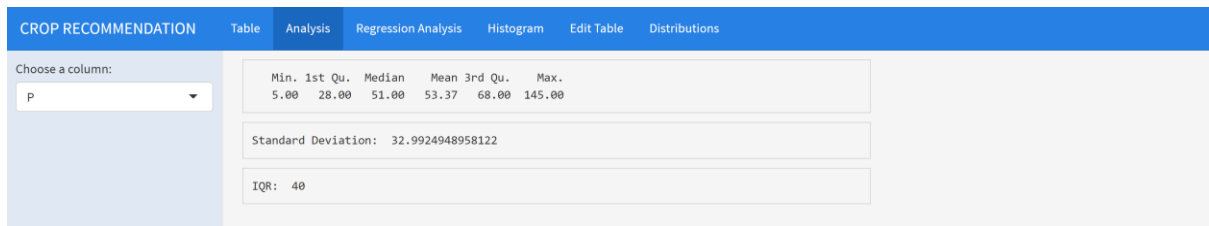


Figure 3 Analytics done for Phosphorus

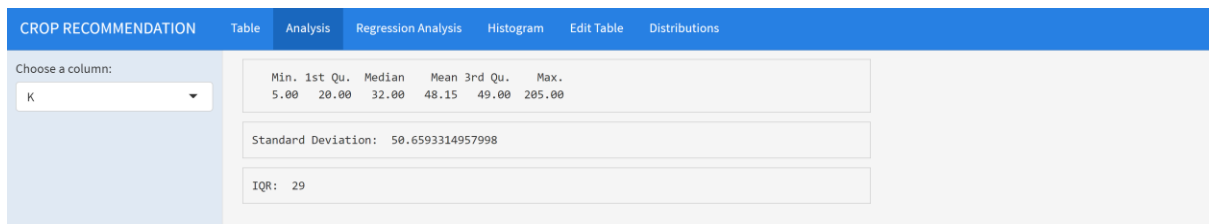


Figure 4 Analysis done for Potassium

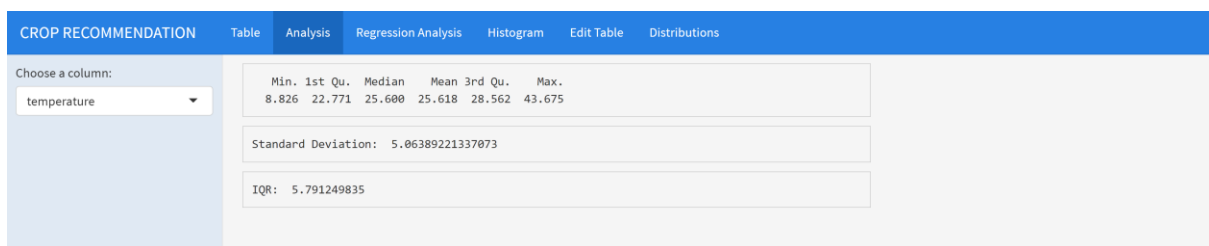


Figure 5 Analysis Done for temperature

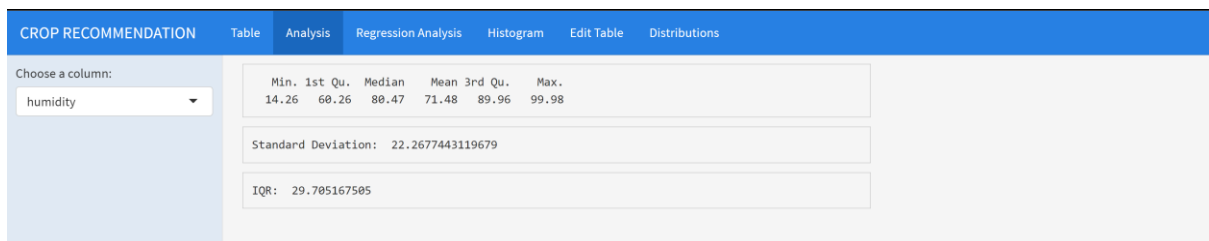


Figure 6 Analysis done for humidity

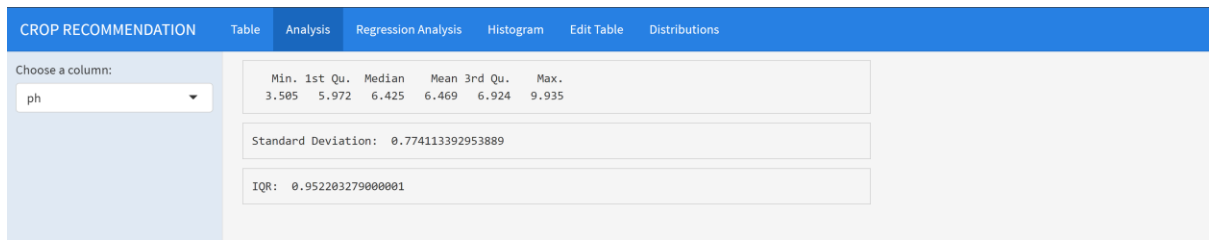


Figure 7 Analysis done for ph

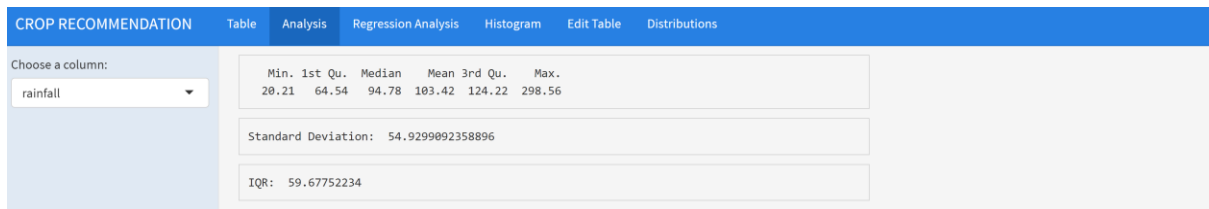


Figure 8 Analysis Done for rainfall

#### 4) Regression Analysis on Each Data

# Regression Analysis

## ph~Rainfall

Regression analysis of "rainfall" with "ph"

```
```{r}
```

```
library(datasets)
```

```
mydata=as.data.frame(read.csv("crop.csv"))
```

```
colnames(mydata)=c("N", "P", "K", "temperature", "humidity", "ph", "rainfall")
```

```
model1=lm(ph~rainfall, data=mydata)
```

```
plot(ph~rainfall, data=mydata,
      xlab="rainfall", ylab="ph")
```

```
abline(model1)
```

```
```
```

Regression analysis of "rainfall" with "ph"

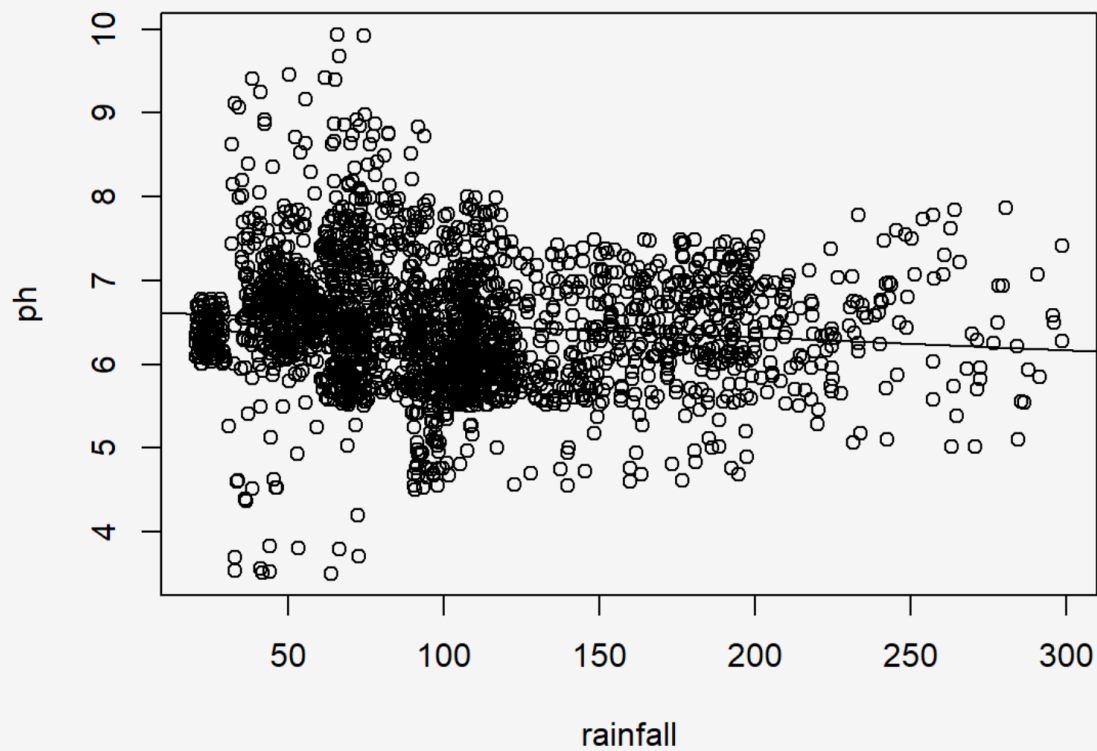


Figure 9 Rainfall ph regression analysis

```
ph~humidity
```

```
Regression analysis of "Humidity" with "ph"
```

```
```{r}
```

```
library(datasets)
```

```
mydata=as.data.frame(read.csv("crop.csv"))
```

```
colnames(mydata)=c("N", "P", "K", "temperature", "humidity", "ph", "rainfall")
```

```
model1=lm(ph~humidity, data=mydata)
```

```
plot(ph~humidity, data=mydata,
```

```
      xlab="humidity", ylab="ph")
```

```
abline(model1)
```

```
```
```

### Regression analysis of “Humidity” with “ph”

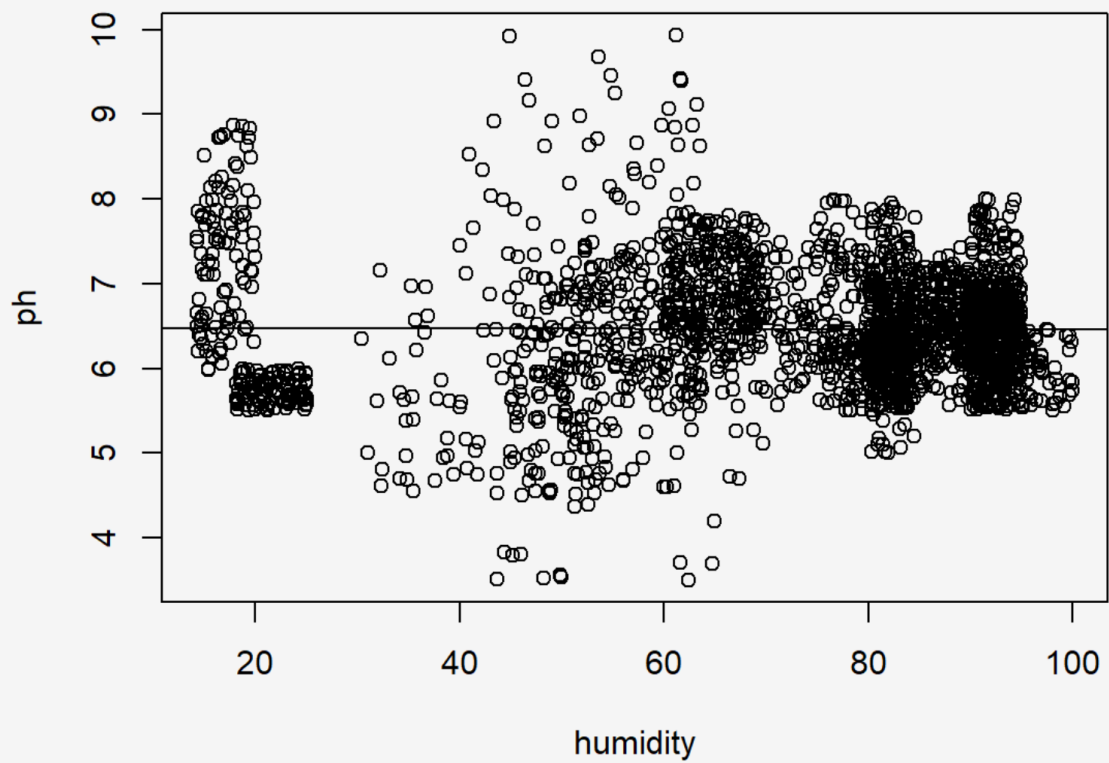


Figure 10 Regression analysis for ph and humidity

```
N~ph
```

```
Regression analysis of "ph" with "N"
```

```
```{r}
```

```
library(datasets)
```

```
mydata=as.data.frame(read.csv("crop.csv"))
```

```
colnames(mydata)=c("N", "P", "K", "temperature", "humidity", "ph", "rainfall")
```

```
model1=lm(N~ph, data=mydata)
```

```
plot(N~ph, data=mydata,
```

```
      xlab="ph", ylab="N")
```

```
abline(model1)
```

```
```
```



### Regression analysis of "ph" with "N"

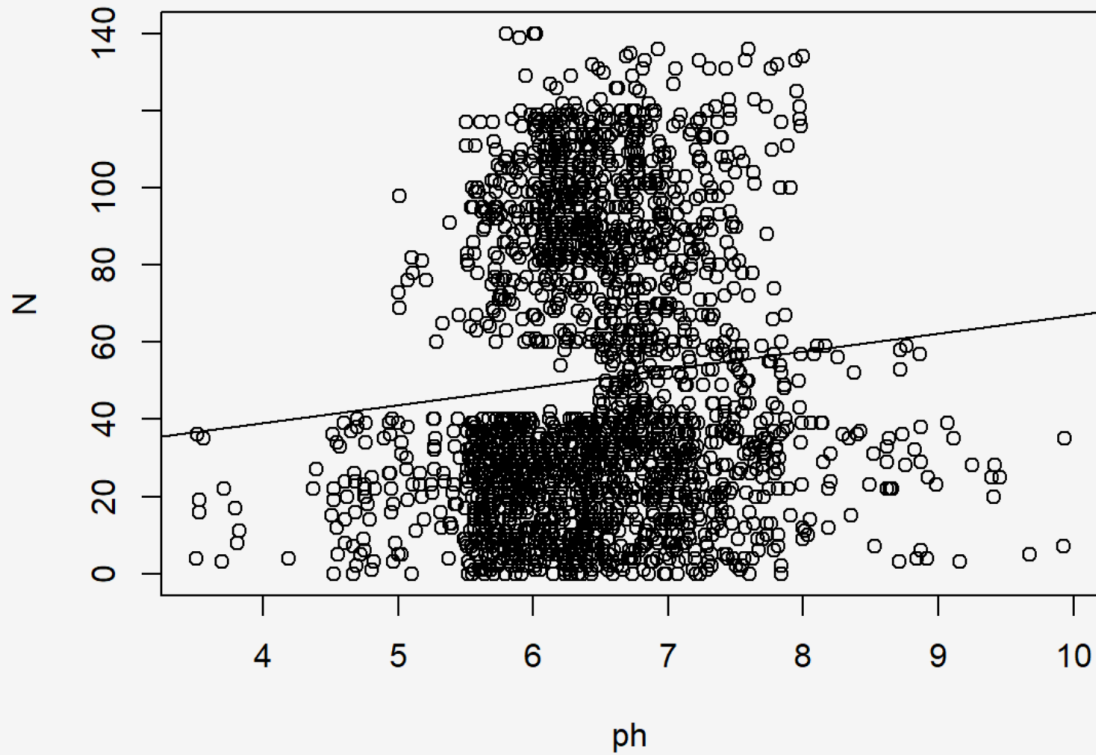


Figure 11 Regression analysis for ph and N

```
P~ph
```

```
Regression analysis of "ph" with "P"
```

```
``{r}
```

```
library(datasets)
```

```
mydata=as.data.frame(read.csv("crop.csv"))
```

```
colnames(mydata)=c("N", "P", "K", "temperature", "humidity", "ph", "rainfall")
```

```
model1=lm(P~ph, data=mydata)
```

```
plot(P~ph, data=mydata,
```

```
 xlab="ph", ylab="P")
```

```
abline(model1)
```

```
``
```

### Regression analysis of “ph” with “P”

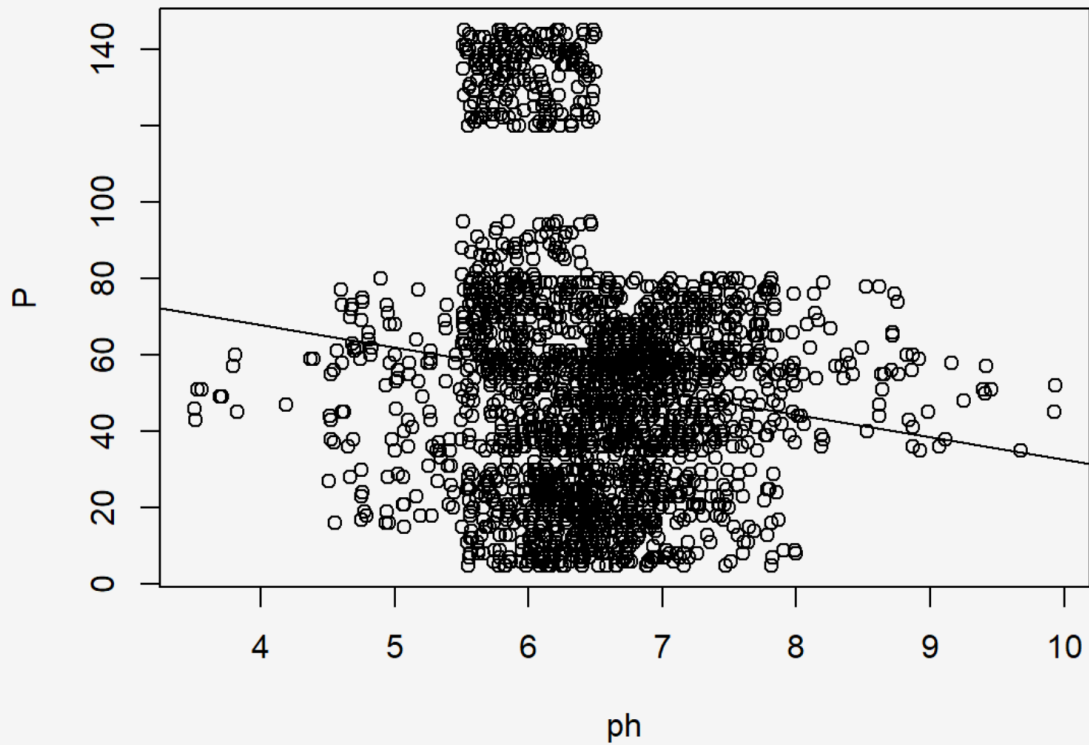


Figure 12 Regression analysis for ph and P

```
K~ph
```

```
Regression analysis of "ph" with "K"
```

```
``{r}
```

```
library(datasets)
```

```
mydata=as.data.frame(read.csv("crop.csv"))
```

```
colnames(mydata)=c("N", "P", "K", "temperature", "humidity", "ph", "rainfall")
```

```
model1=lm(K~ph, data=mydata)
```

```
plot(K~ph, data=mydata,
```

```
 xlab="ph", ylab="K")
```

```
abline(model1)
```

```
``
```

### Regression analysis of “ph” with “K”

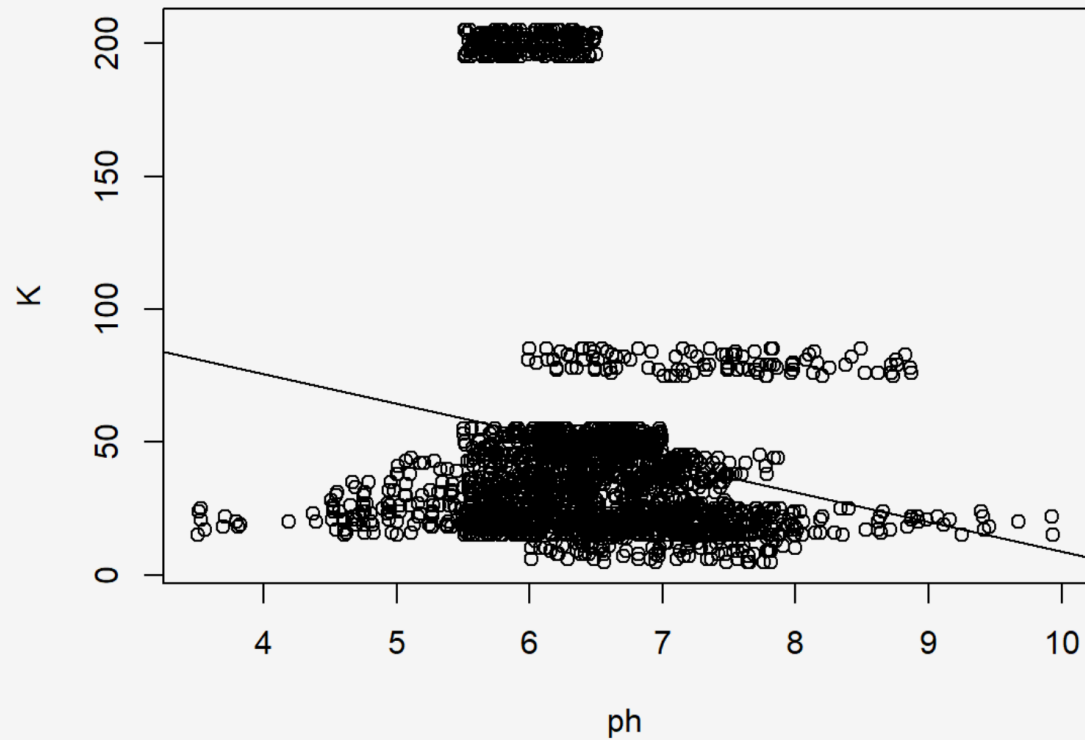


Figure 13 Regression Analysis for ph and K

#### 5) Histogram on Each Data (Most effort put on this one)

# Histogram

```
```{r setup, include=FALSE}
library(tidyverse)
library(DT)
startData <- read.csv("crop.csv")
updateData <- reactive(
  startData %>% group_by(!!! rlang::syms(input$GB)) %>%
  summarise_if(is.numeric, sum, na.rm=T))
```
```

## Column {.sidebar}

```

```{r}

selectInput(inputId = "GB",label= "Group By", choices = names(startData))

selectInput(inputId = "Metric",label= "Metric", choices =
names (select_if(startData, is.numeric)))

```

```

## Column

### Histogram

```

```{r}

renderPlot({
  updateData()%>%
    ggplot(aes(x=!! rlang::sym(input$GB),y=!! rlang::sym(input$Metric),fill=!!
rlang::sym(input$GB))) +
    geom_col()
})

```

```

### Table

```

```{r}

renderDT(
  updateData(), rownames = F, extensions = 'Buttons', filter="top", editable=T,
  options = list(
    dom = 'Blfrtip',
    buttons = c('copy', 'csv', 'excel', 'pdf', 'print'),
    lengthMenu = list(c(10,50, 100, -1), c(10,50, 100, "All"))
  )
)

```

```

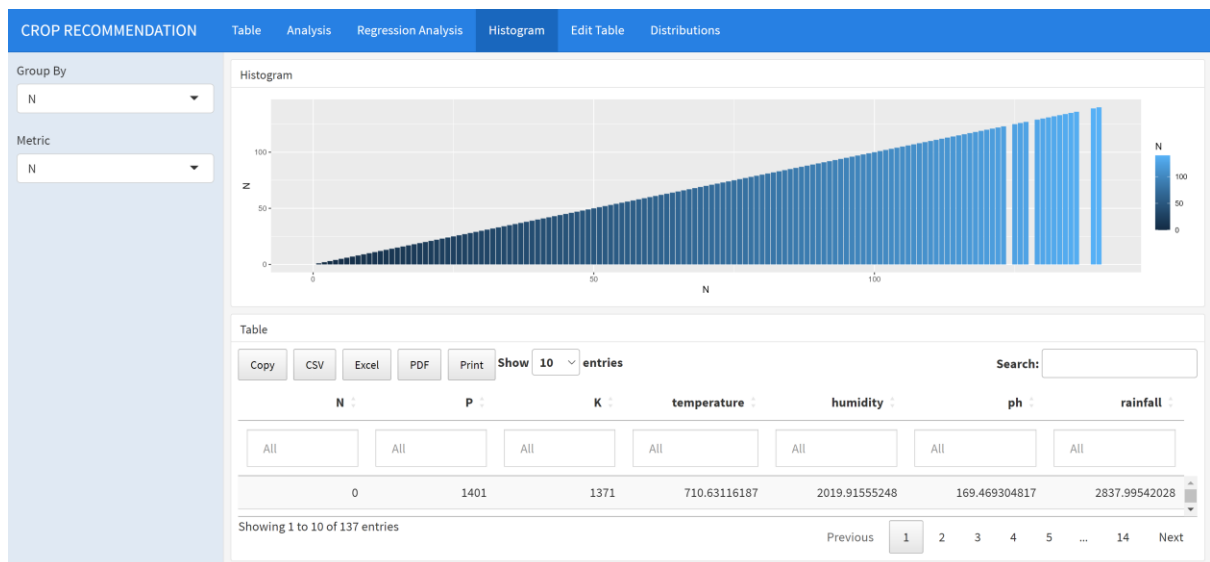


Figure 14 Histogram for N and N

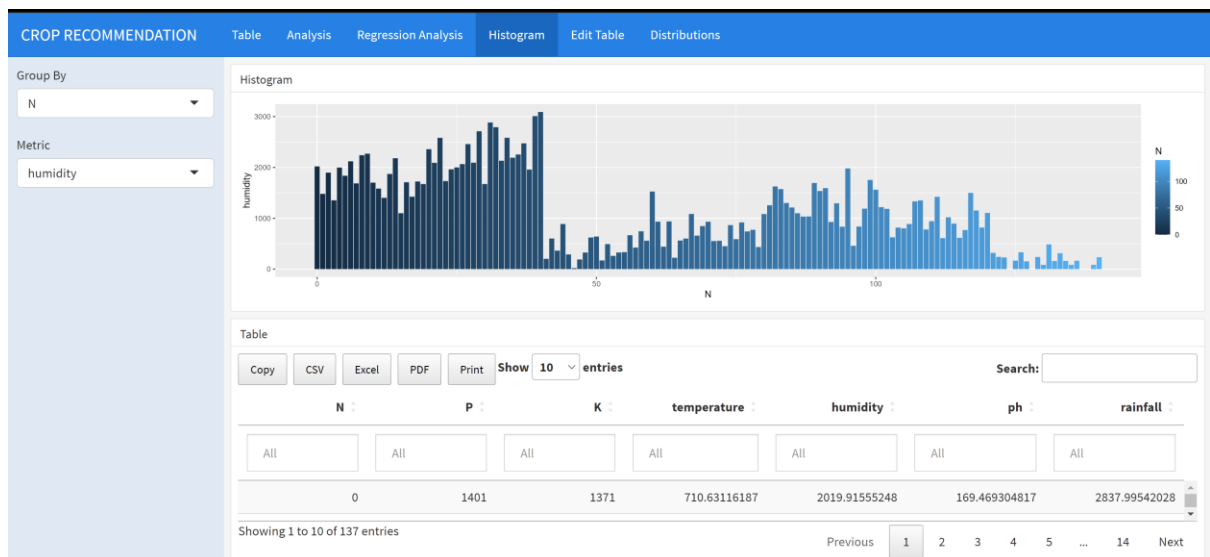


Figure 15 Histogram For N and Humidity

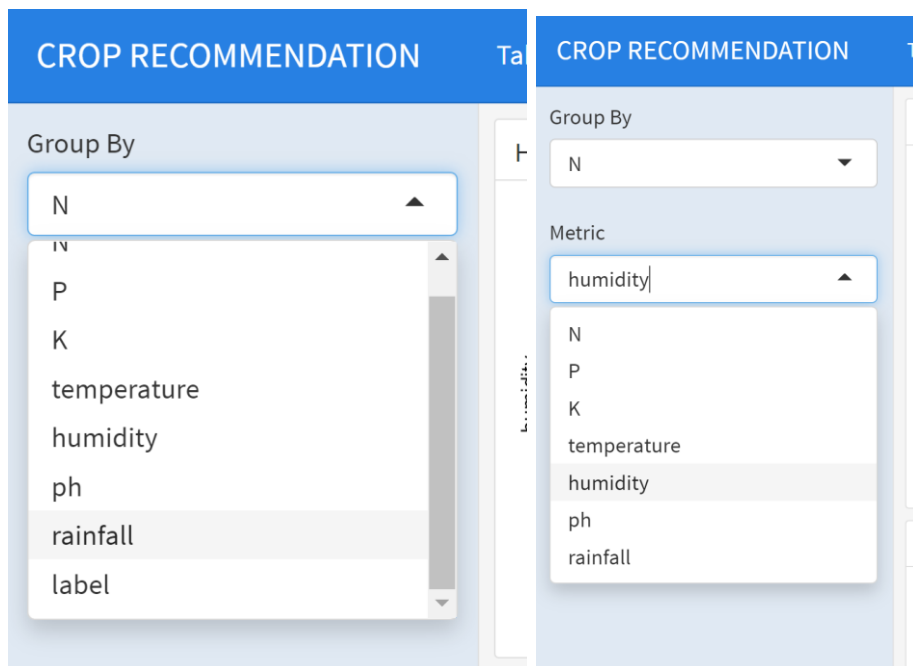


Figure 16 Special thing about this Histogram is that it is dynamic and you can choose metric and group by by your self on which you want to apply methods

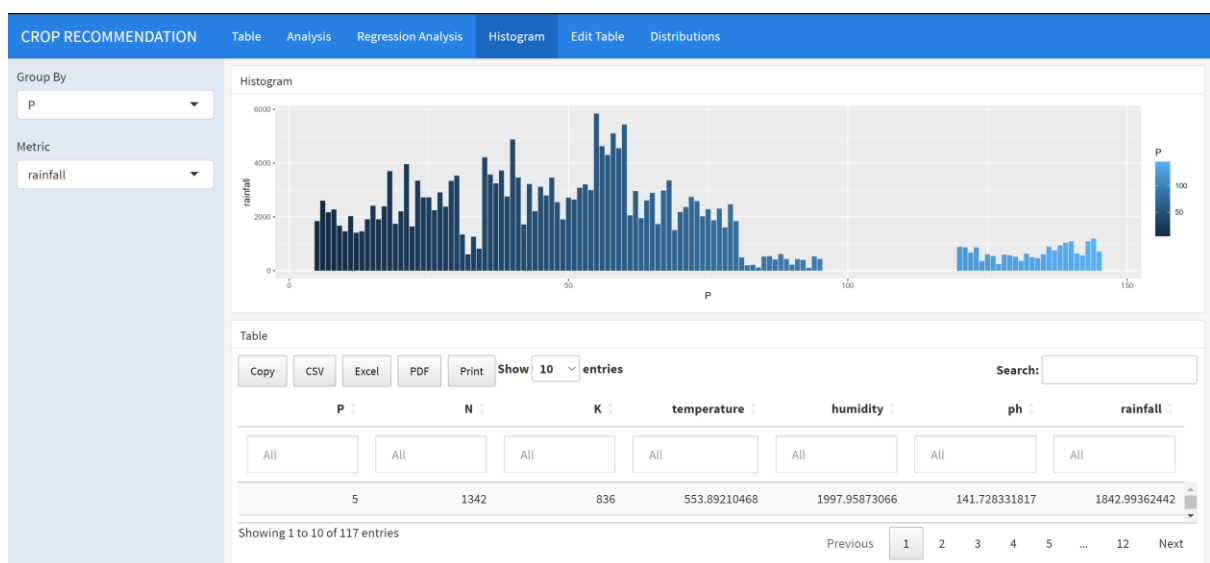


Figure 17 Histogram of P and Rainfall

|       |      |     |       |     |       |                 |         |
|-------|------|-----|-------|-----|-------|-----------------|---------|
| Table | Copy | CSV | Excel | PDF | Print | Show 10 entries | Search: |
|-------|------|-----|-------|-----|-------|-----------------|---------|

Figure 18 You can also print data to your favourite form

CROP RECOMMENDATION.pdf

1 / 1 100%

CROP RECOMMENDATION

| P  | N    | K   | temperature  | humidity      | ph            | rainfall      |
|----|------|-----|--------------|---------------|---------------|---------------|
| 5  | 1342 | 836 | 553.89210468 | 1997.95873066 | 141.728331817 | 1842.99362442 |
| 6  | 1113 | 847 | 642.75437685 | 2195.73253787 | 152.750300468 | 2601.2710867  |
| 7  | 1377 | 831 | 648.19627282 | 2241.38981848 | 162.899706487 | 2170.35098161 |
| 8  | 784  | 564 | 490.87856078 | 1850.87084754 | 133.085856241 | 2274.68748182 |
| 9  | 728  | 626 | 428.97181879 | 1542.27162446 | 109.358972873 | 1674.58125827 |
| 10 | 794  | 542 | 374.12338002 | 1262.17735889 | 89.881886839  | 1461.42910416 |
| 11 | 1066 | 708 | 542.9399817  | 1924.03752339 | 136.70722379  | 2026.43927936 |
| 12 | 1086 | 594 | 441.2723966  | 1558.42350019 | 108.048196212 | 1411.62691496 |
| 13 | 755  | 528 | 386.58788553 | 1389.46987153 | 96.847147804  | 1460.86526596 |
| 14 | 1057 | 672 | 463.3102165  | 1728.81627888 | 123.012254527 | 1912.49521361 |

Figure 19 Pdf of 10 datas

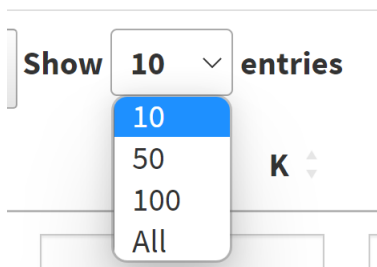


Figure 20 Also you can choose how much data you want to print in which format

## 6) Editing the data

### # Edit Table

```

```{r}

library(DT)

startData <- read.csv("crop.csv", header = TRUE, sep = ",")

DT::datatable(startData)

var1 <- renderDT(
  updateData(), rownames = F, extensions = 'Buttons', filter="top", editable=T,
  options = list(
    dom = 'Blfrrtip',
    buttons = c('Insert', 'Delete'),
    lengthMenu = list(c(10,50), c(10,50))
  )
)
...

```

	N	P	K	temperature	humidity	ph	rainfall	label
1	90	42	43	20.87974	82.00274	6.502985	202.9355	rice
2	85	58	41	21.77046	80.31964	7.038096	226.6555	rice
3	60	55	44	23.00446	82.32076	7.840207	263.9642	rice
4	74	35	40	26.4911	80.15836	6.980401	242.864	rice
5	78	42	42	20.13017	81.60487	7.628473	262.7173	rice
6	69	37	42	23.05805	83.37012	7.073454	251.055	rice
7	69	55	38	22.70884	82.63941	5.700806	271.3249	rice
8	94	53	40	20.27774		5.718627	241.9742	rice
9	89	54	38	24.51588	83.53522	6.685346	230.4462	rice
10	68	58	38	23.22397	83.03323	6.336254	221.2092	rice
11	91	53	40	26.52724	81.41754	5.386168	264.6149	rice
12	90	46	42	23.97898	81.45062	7.502834	250.0832	rice
13	78	58	44	26.8008	80.88685	5.108682	284.4365	rice
14	93	56	36	24.01498	82.05687	6.984354	185.2773	rice
15	94	50	37	25.66585	80.66385	6.94802	209.587	rice
16	60	48	39	24.28209	80.30026	7.042299	231.0863	rice
17	85	38	41	21.58712	82.78837	6.249051	276.6552	rice
18	91	35	39	23.79392	80.41818	6.97086	206.2612	rice
19	77	38	36	21.86525	80.1923	5.953933	224.555	rice

Figure 21 You can also edit and save the data

CROP RECOMMENDATION							
Table Analysis Regression Analysis Histogram Edit Table Distributions							
Show	10	entries					
	N	P	K	temperature	humidity	ph	
1	90	42	43	20.87974371	82.00274423	6.502985292	
2	85	58	41	21.77046169	80.31964408	7.038096361	
3	60	55	44	23.00445915	82.3207629	7.840207144	
4	74	35	40	26.49109635	80.15836264	6.980400905	
5	78	42	42	20.13017482	81.60487287	7.628472891	
6	69	37	42	23.05804872	83.37011772	7.073453503	
7	69	55	38	22.70883798	82.63941394	5.70080568	
8	94	53	40	20.27774362	82.89408619	5.718627178	
9	89	54	38	24.51588066	83.5352163	6.685346424	
10	68	58	38	23.22397386	83.03322691	6.336253525	

Showing 1 to 10 of 2,200 entries

Figure 22 The editing will be saved here Dynamically

7) Distributions:

```
# Distributions
```

```
```{r}

library(tidyverse)
library(DT)

sData <- read.csv("crop.csv")

uData <- reactive(

sData %>% group_by(!!! rlang::syms(input$GB)) %>%

summarise_if(is.numeric,sum,na.rm=T))

```
```

```
## Column {.sidebar}
```

```
```{r}

selectInput("Distributions", "Select a Distribution type", choices = c("Normal", "Uniform"))

sliderInput("sampleSize", "Select a sample size", min = 0, max = 300, value= 270, step= 0)

conditionalPanel(condition = "input.Distribution == 'Normal'", textInput("Mean", "Please Select Mean", 10), textInput("sd",
"Please Select Standard Deviation"))

conditionalPanel(condition = "input.Distribution == 'Uniform'", textInput("u1", "Val1: ", value = 1), textInput("u2", "Val2: ",
value = 1))

server <- function(input, output, session){

output$myPlot <- renderPlot({

distType <- input$Distribution

size <- input$sampleSize

if(distType == "Normal"){

randomVec <- rnorm(size, mean = as.numeric(input$mean), sd=as.numeric(input$sd))

}

else{

randomVec <- runif(size, min = 0, max = 300)

}

hist(randomVec, col="blue")})

```
```

```
## Column
```

Distribution Plot

```
```{r}  
plotOutput("myPlot")
```
```

Table

```
```{r}  
renderDT(
 updateData(), rownames = F, extensions = 'Buttons', filter="top", editable=T,
 options = list(
 dom = 'Blfrtip',
 buttons = c('copy', 'csv', 'excel', 'pdf', 'print'),
 lengthMenu = list(c(10,50, 100, -1), c(10,50, 100, "All"))
)
)
```
```

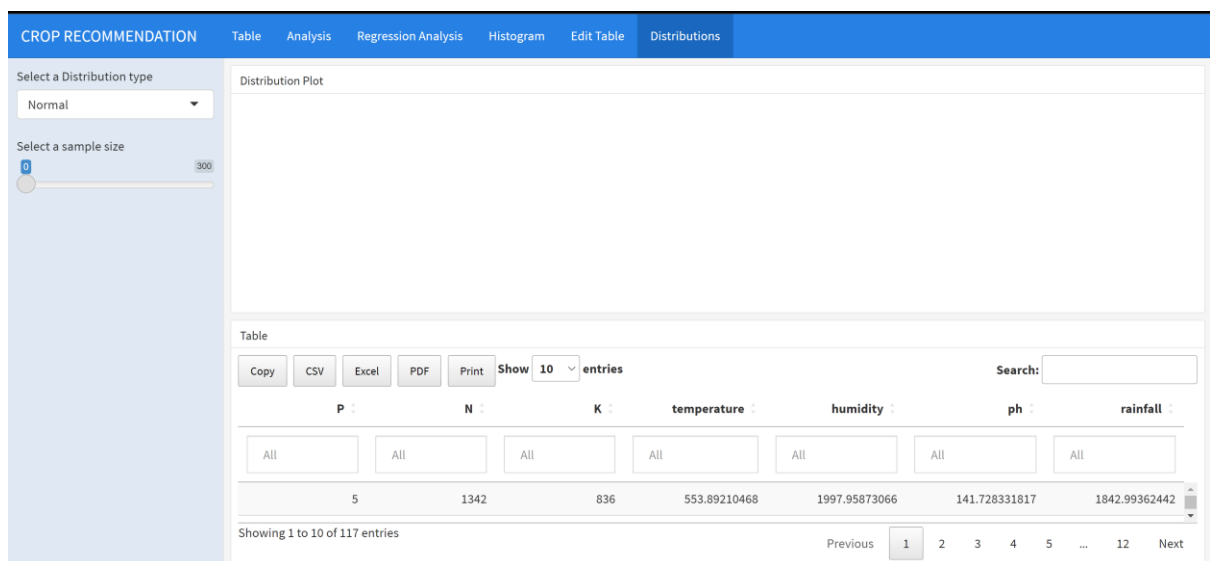


Figure 23 Distribution picture

CROP RECOMMENDATION

Select a Distribution type

Normal

Normal

Uniform

Figure 24 Applied Uniform and Normal Distribution on this data as it was continuous