

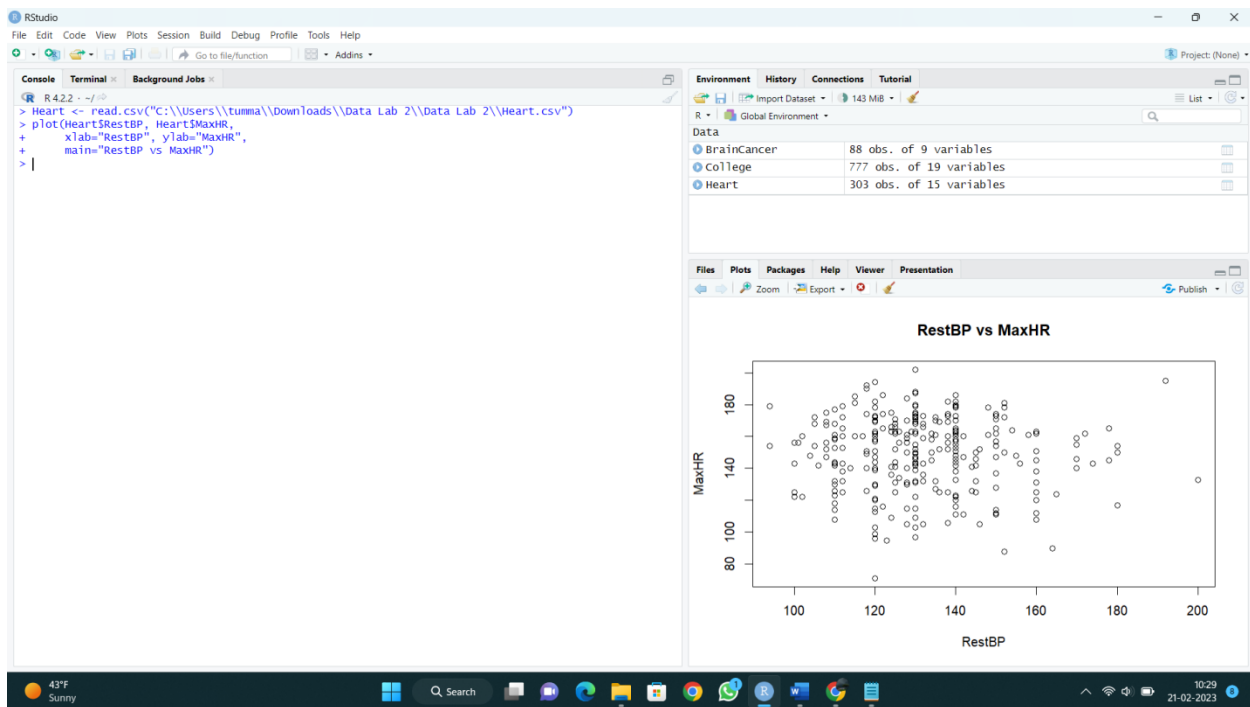
1. Visualizations from Heart

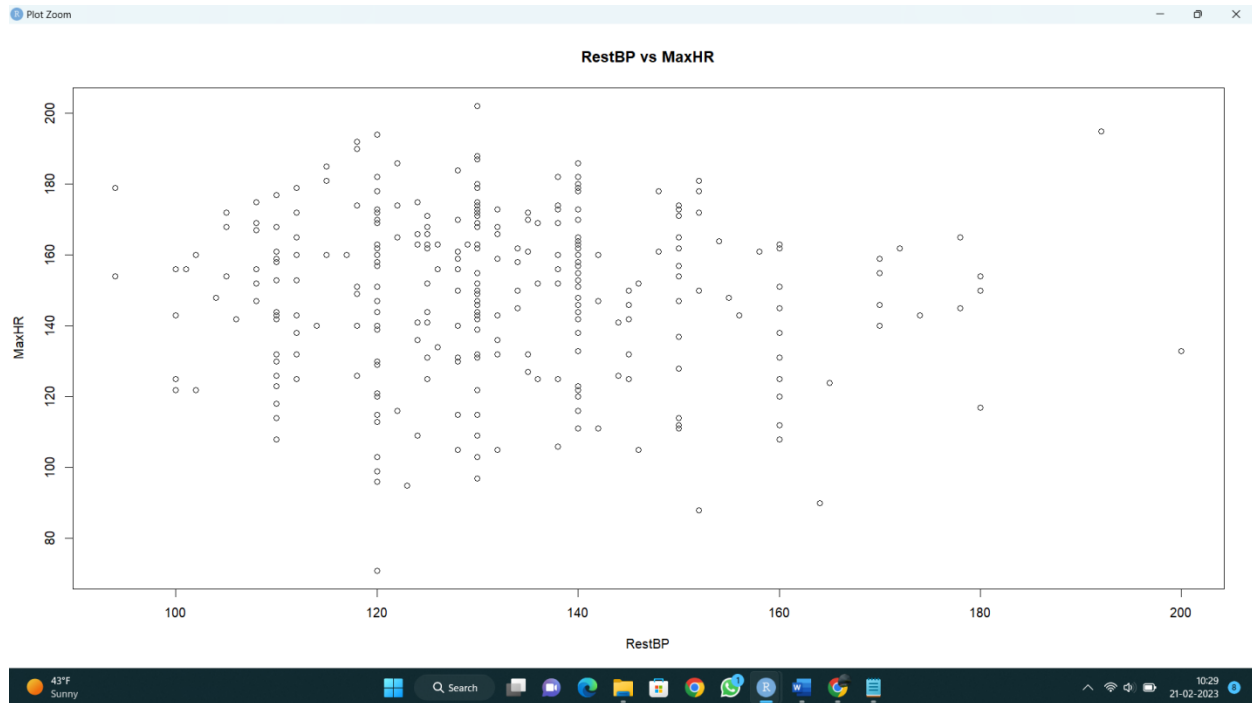
Used the read.csv function to import the Credit csv file

```
Heart <- read.csv("C:\\Users\\tumma\\Downloads\\Data Lab 2\\Data Lab 2\\Heart.csv")
```

1.1 Scatter plot of RestBP vs MaxHR

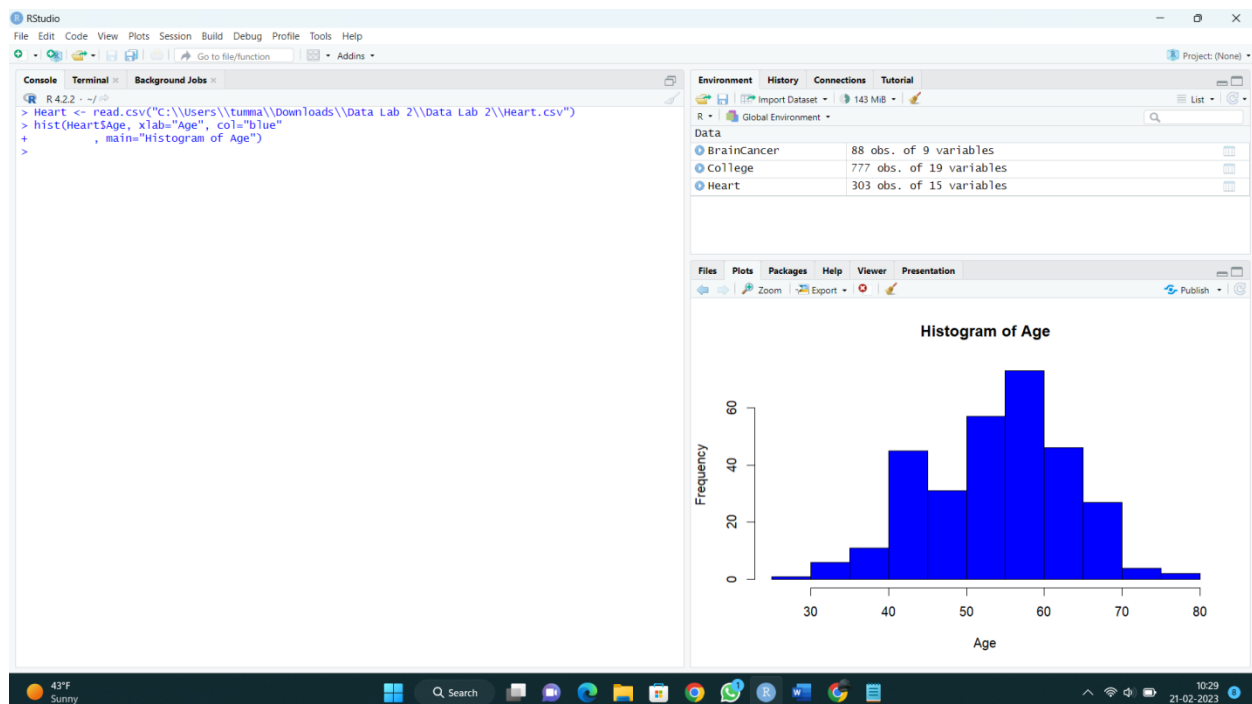
```
plot(Heart$RestBP, Heart$MaxHR,  
     xlab="RestBP", ylab="MaxHR",  
     main="RestBP vs MaxHR")
```

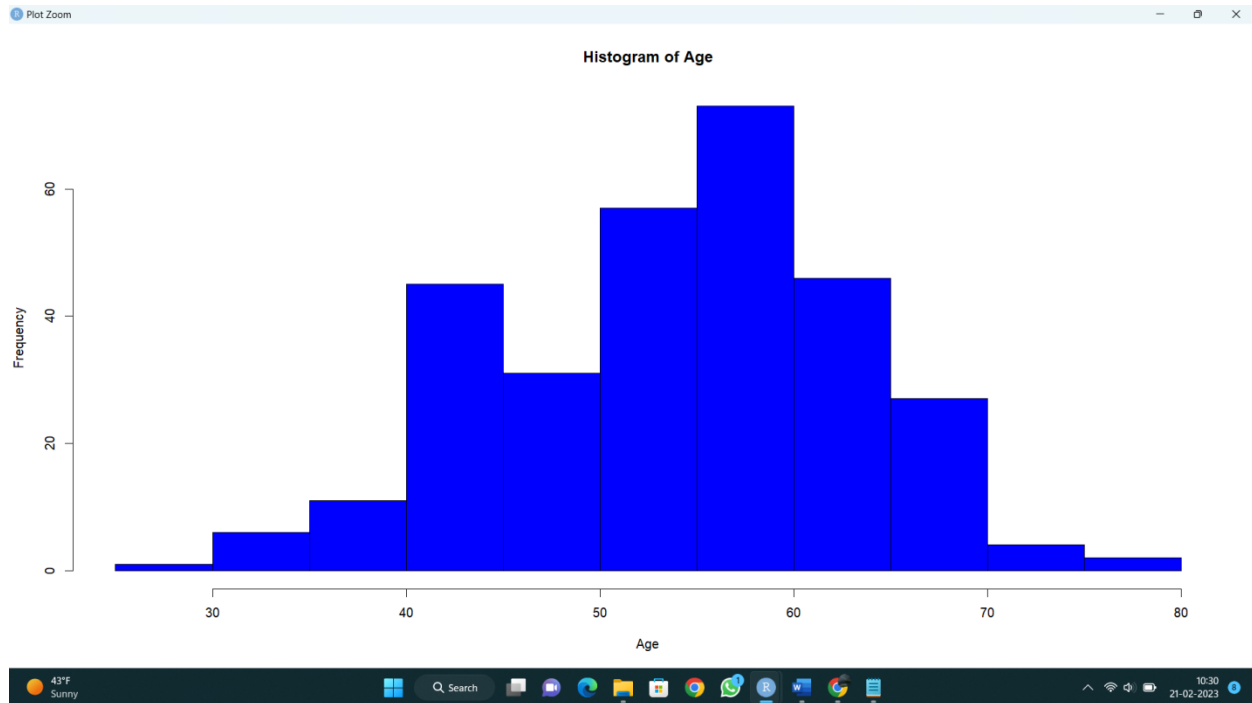




1.2 Histogram of Age

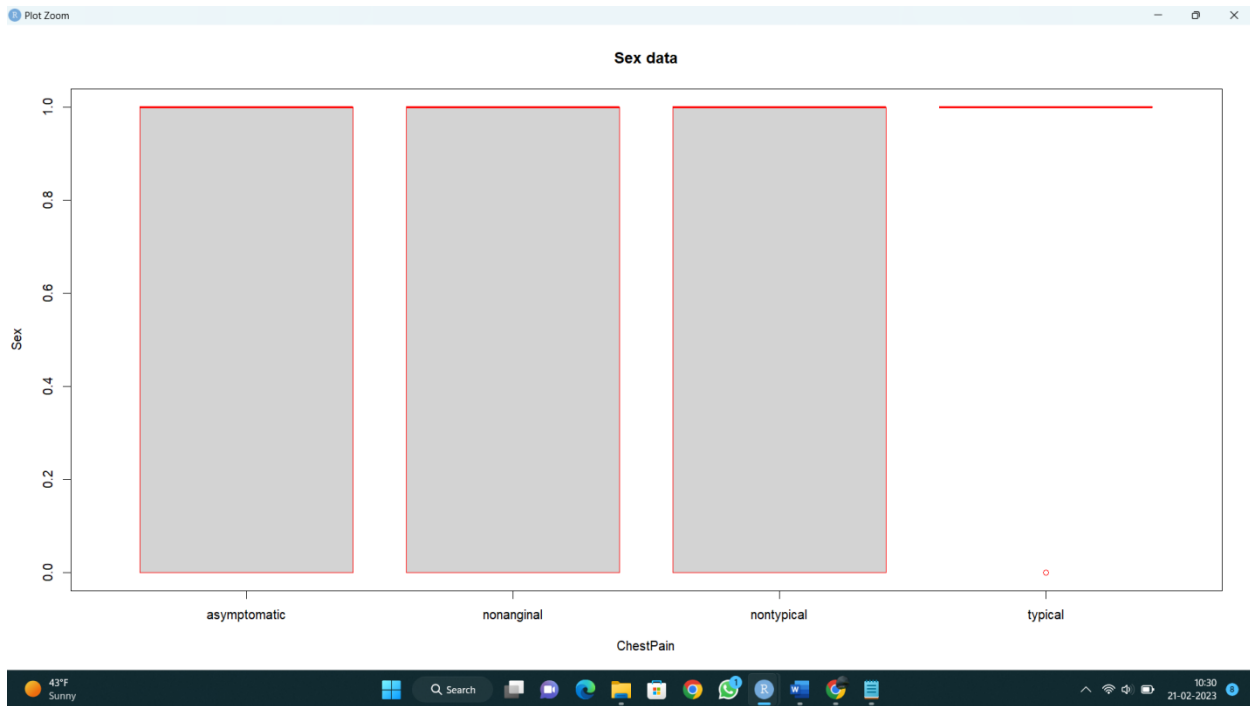
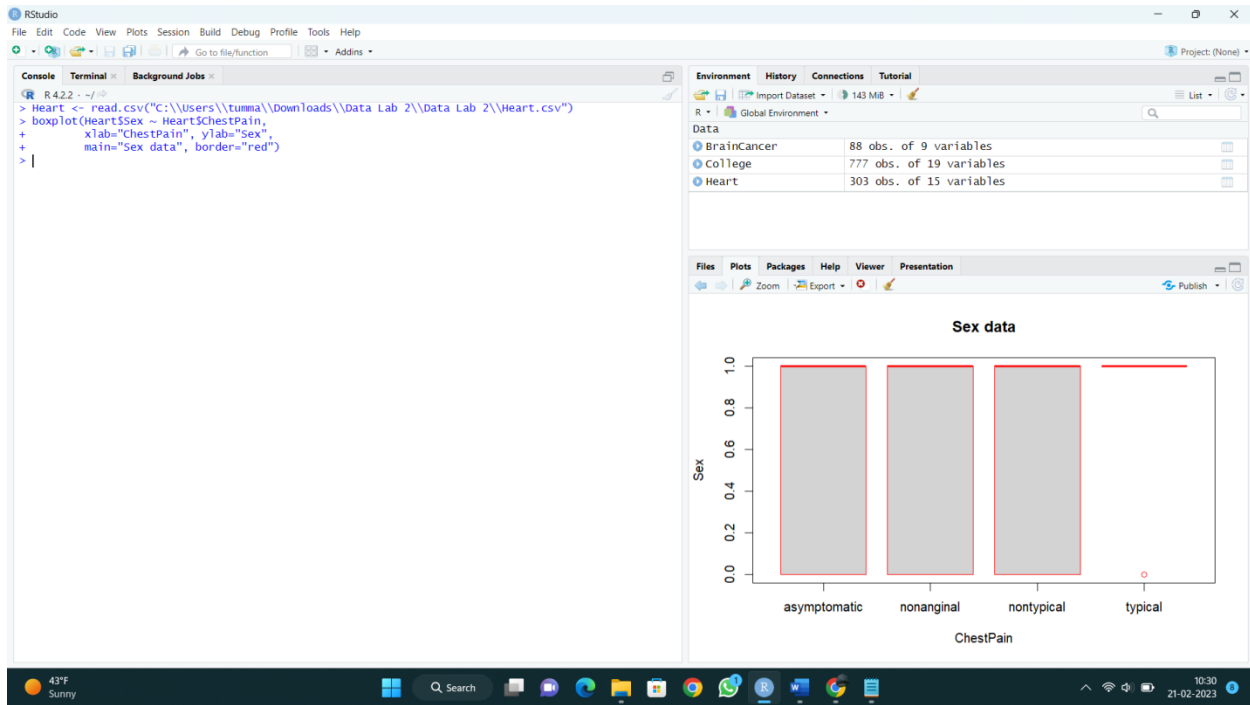
```
hist(Heart$Age, xlab="Age", col="blue",
     , main="Histogram of Age")
```





1.3 Boxplot of Sex vs ChestPain

```
boxplot(Heart$Sex ~ Heart$ChestPain,  
        xlab="ChestPain", ylab="Sex",  
        main="Sex data", border="red")
```



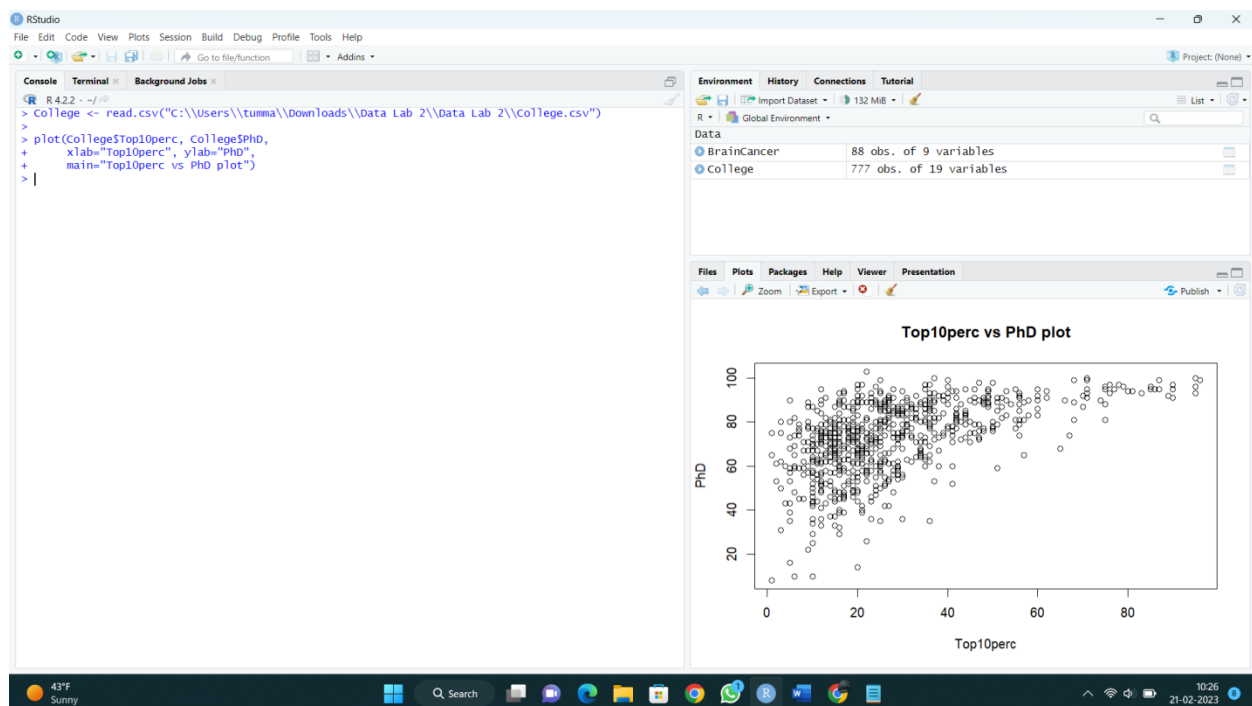
2 Visualizations from College

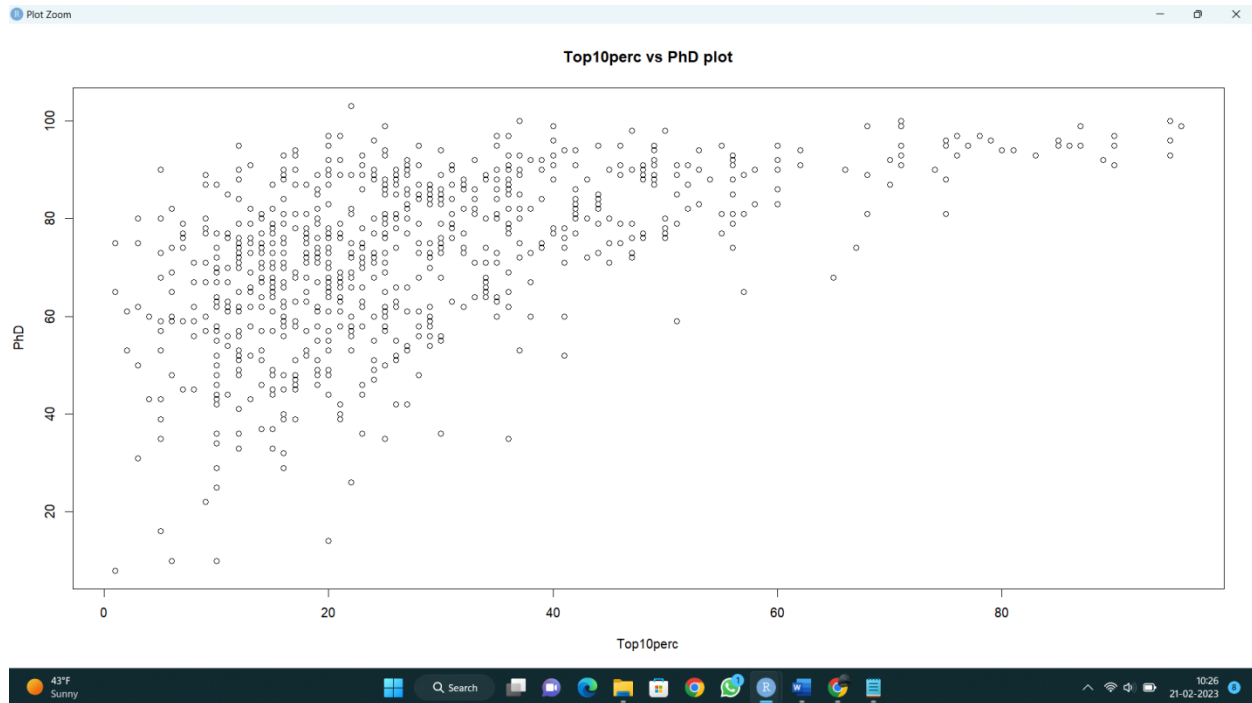
Used the read.csv function to import the College csv file

```
College <- read.csv("C:\\Users\\tumma\\Downloads\\Data Lab 2\\Data Lab 2\\College.csv")
```

2.1 Scatter plot of **Top10perc** vs **PhD**

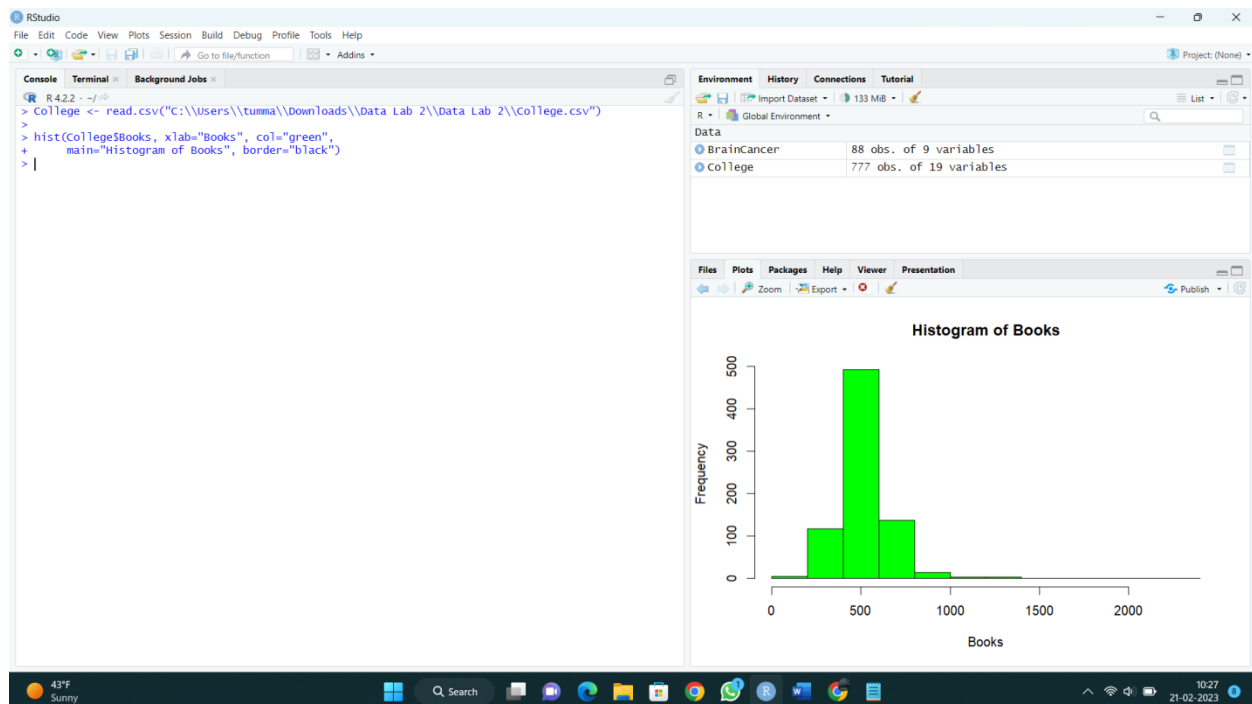
```
plot(College$Top10perc, College$PhD,  
      xlab="Top10perc", ylab="PhD",  
      main="Top10perc vs PhD plot")
```

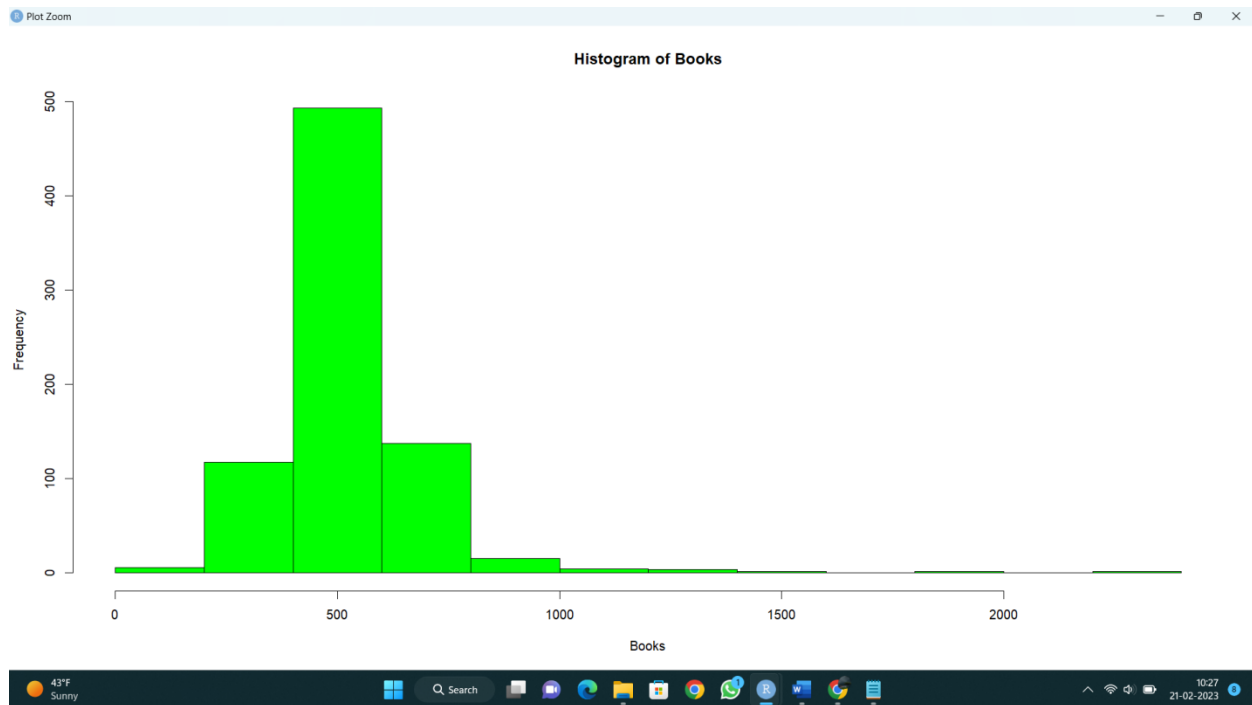




2.2 Histogram of Books

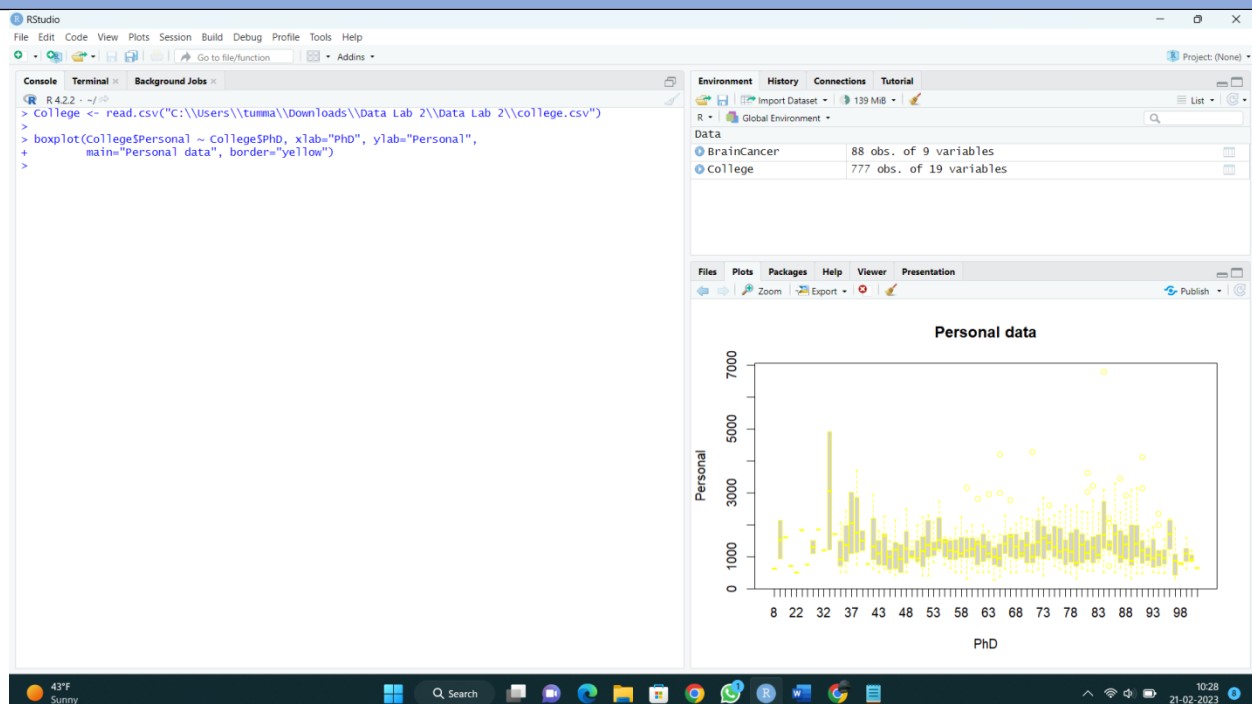
```
hist(College$Books, xlab="Books", col="green",
     main="Histogram of Books", border="black")
```



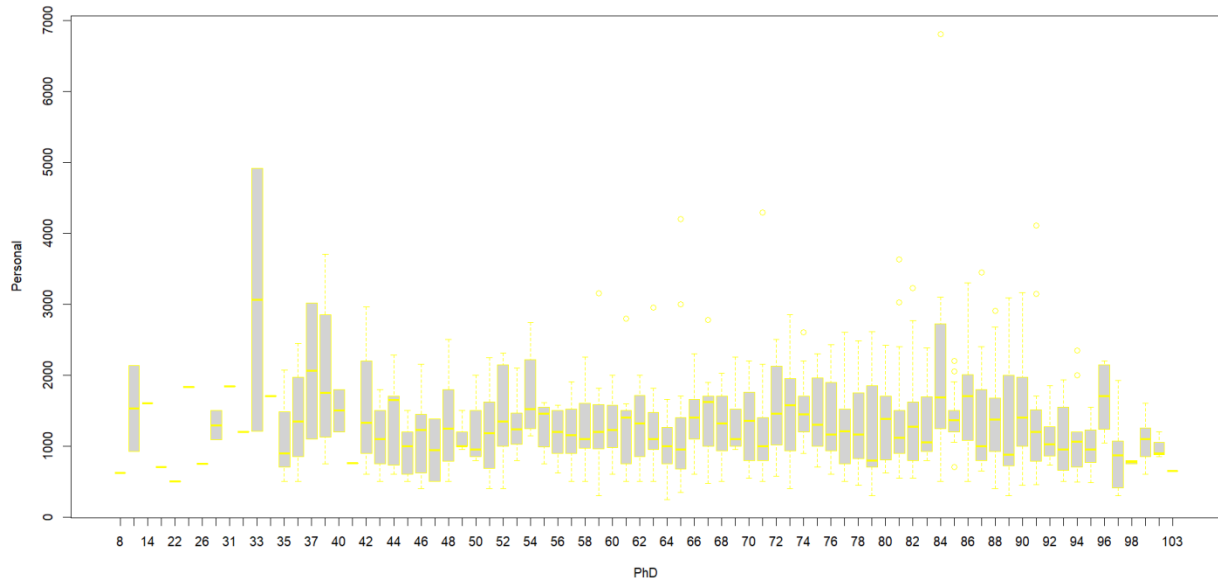


2.3 Boxplot of Personal vs PhD

```
boxplot(College$Personal ~ College$PhD, xlab="PhD", ylab="Personal",  
main="Personal data", border="yellow")
```



Personal data



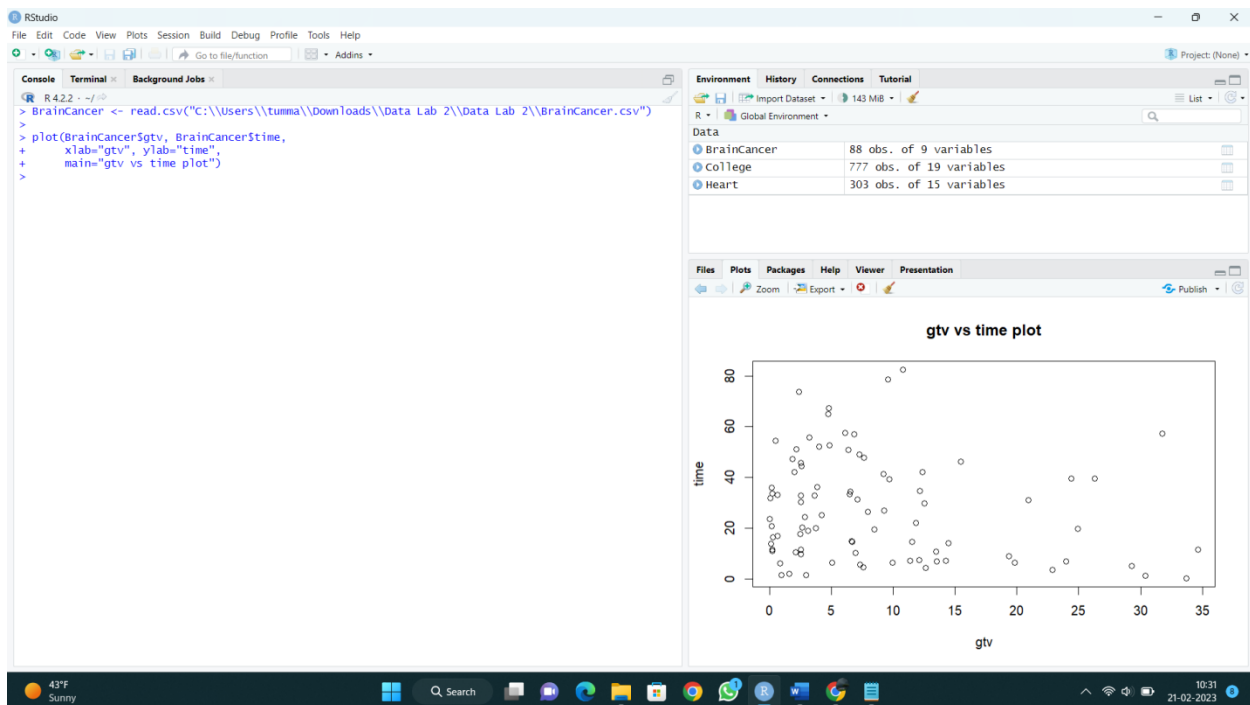
3 Visualizations from BrainCancer

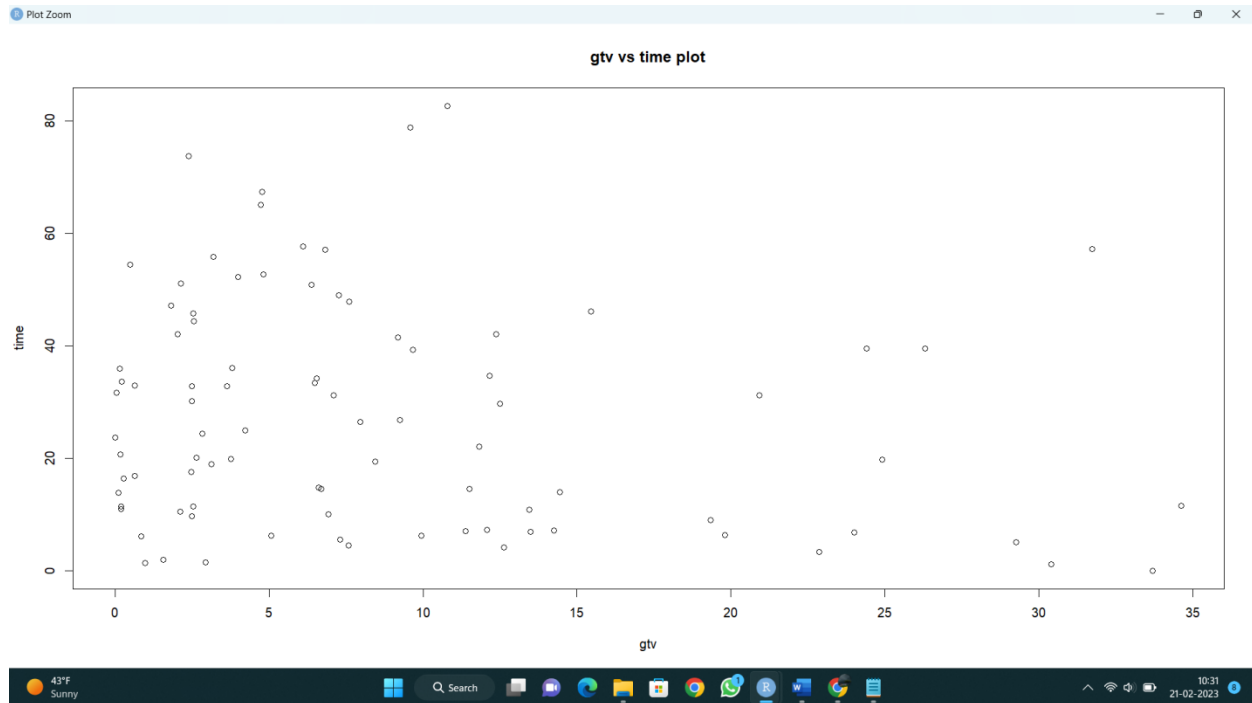
Used the read.csv function to import the BrainCancer csv file

```
BrainCancer <- read.csv("C:\\Users\\tumma\\Downloads\\Data Lab 2\\Data Lab 2\\BrainCancer.csv")
```

3.1 Scatter plot of gtv vs time

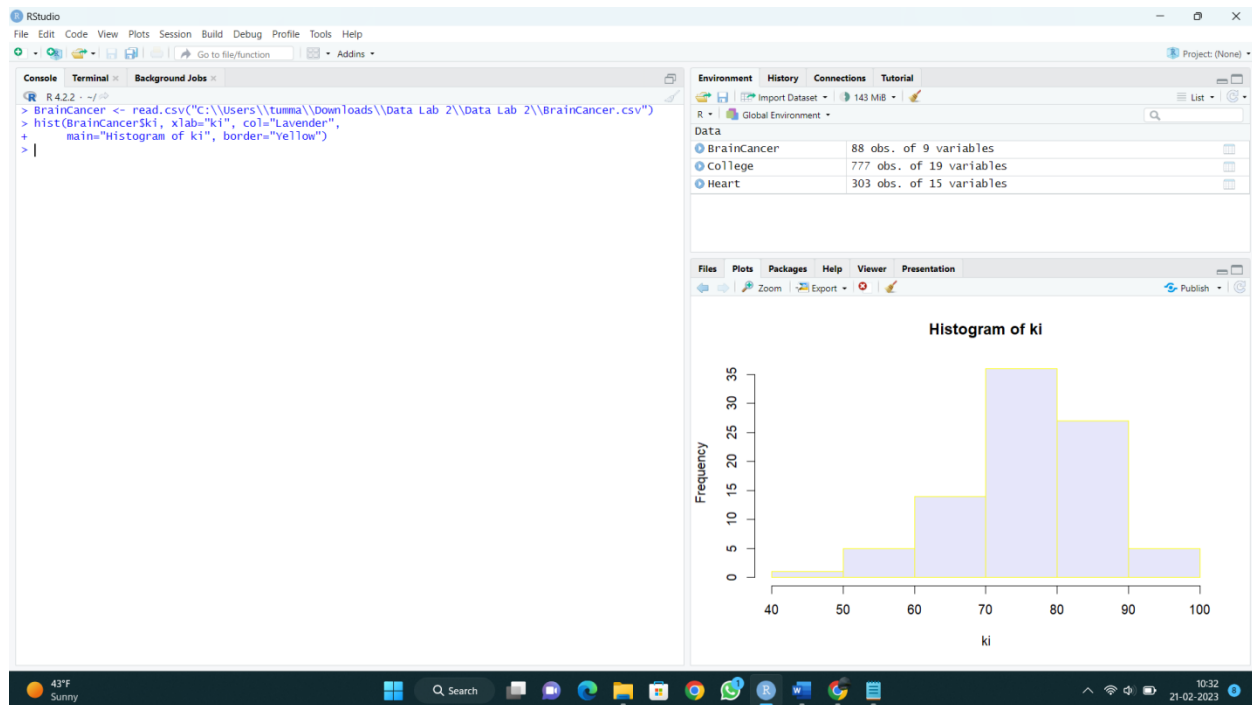
```
plot(BrainCancer$gtv, BrainCancer$time,  
     xlab="gtv", ylab="time",  
     main="gtv vs time plot")
```

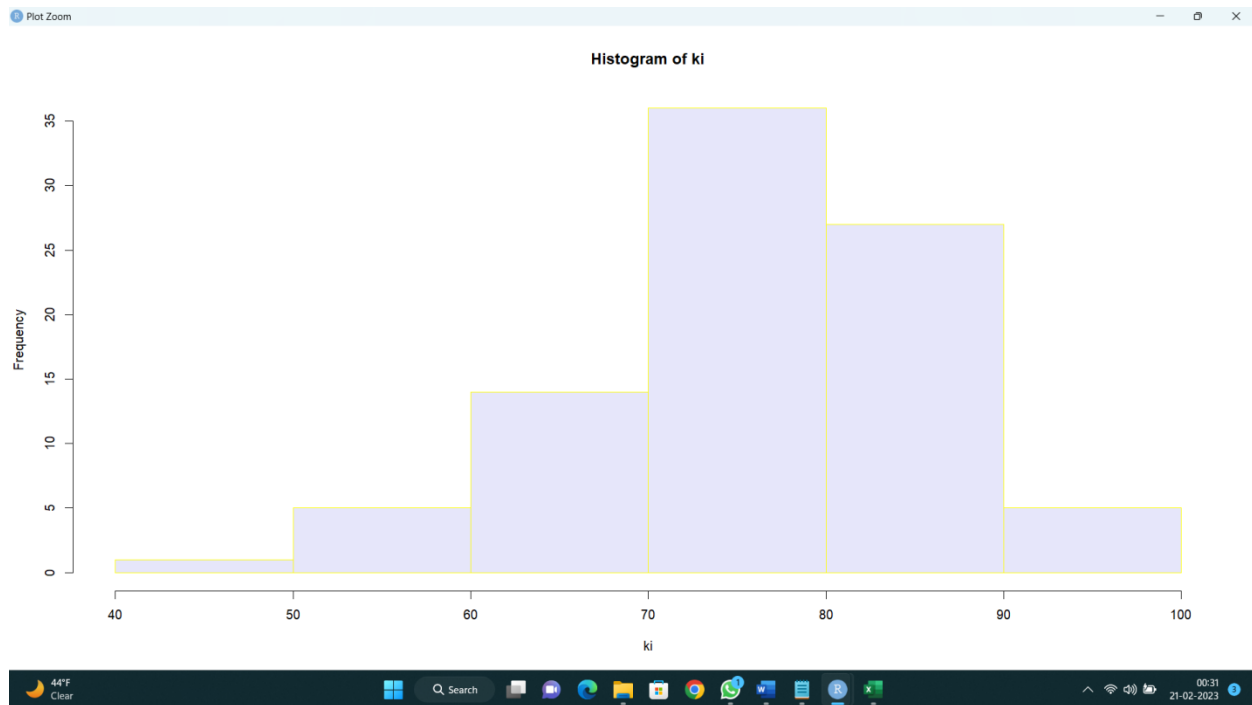




3.2 Histogram of ki

```
hist(BrainCancer$ki, xlab="ki", col="Lavender",
     main="Histogram of ki", border="Yellow")
```





3.3 Boxplot of sex vs ki

```
boxplot(BrainCancer$ki ~ BrainCancer$sex, xlab="sex", ylab="ki",  
main="ki data", border="blue")
```

