## Assignment 2

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2025-10-13

```
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.5'
## (as 'lib' is unspecified)
```

### **Dataset**

The BreastCancer dataset contains cytological measurements of breast tissue samples, used to classify tumors as benign or malignant. It includes numerical features describing cell characteristics derived from microscopic examination.

To load the dataset to your environment, just run the code below:

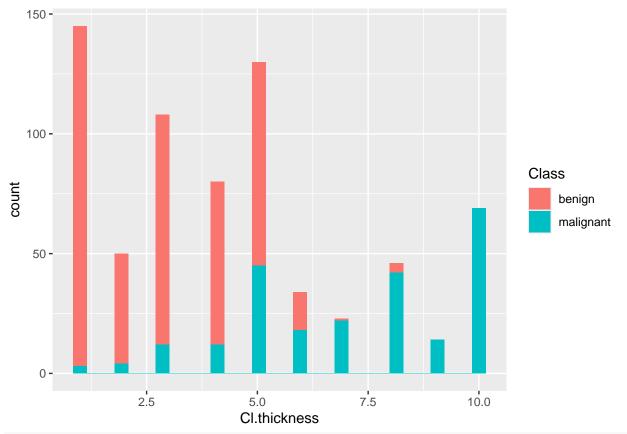
```
install.packages("mlbench")
library(mlbench)
data("BreastCancer")
BreastCancer$Cl.thickness <- as.numeric(BreastCancer$Cl.thickness)</pre>
```

### Drawing a plot for distribution

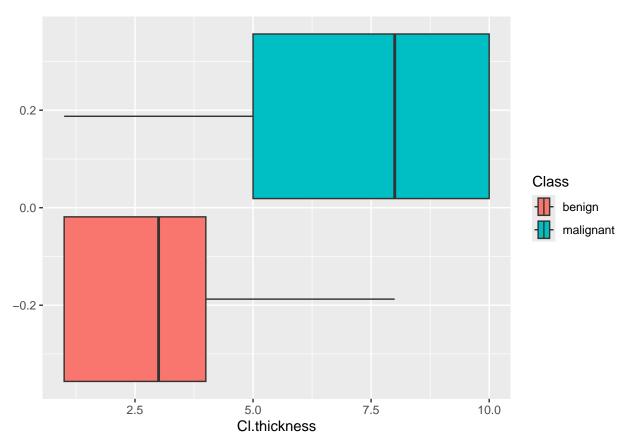
1. Please draw a plot to visualize the distribution of tissue thickness (Cl.thickness) of the patients according to Class indicating whether the tumor in the breast tissue is benign or malignant. (40 pts).

```
ggplot(BreastCancer,aes(x=Cl.thickness,fill=Class))+geom_histogram( )
```

```
## `stat_bin()` using `bins = 30`. Pick better value `binwidth`.
```



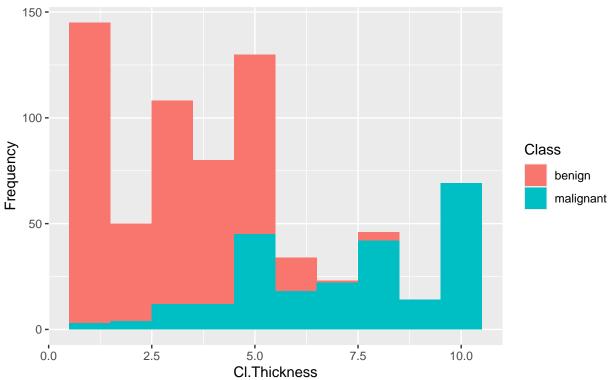
ggplot(BreastCancer,aes(x=Cl.thickness,fill=Class))+geom\_boxplot()



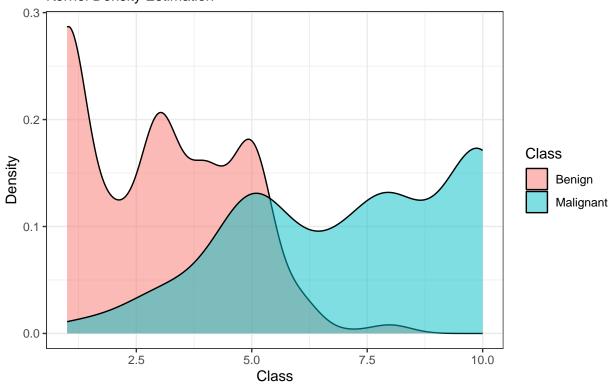
2. Please solve the problems in the plot and make it better by adding some information (30 pts). # I draw histogram. Maybe i can change binwidth = 5 but it is not good idea.

 ${\tt ggplot(BreastCancer, aes(x = Cl.thickness, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + geom\_histogram(binwidth = 1) + labs(x = "Cl.Thicknesh, fill=Class)) + labs(x = "Cl.Thicknesh$ 

# Cl.thickness Distribution of BreastCanser Histogram



## Cl.thickness Distribution of BreastCanser Kernel Density Estimation



#### 3. Interpret the plot (30 pts).

BreastCanser data has 11 variables,699 observation. One of them is Cl.thickness. We want to classify tumors as benign or malignant so you can use some graph. Maybe we can use histogram. Also we can use boxplot for distribution. You know that we find Q1,Q3 and median so we understand left skewed, right skewed. When we look at the boxplots for benign and malignant classes, the median line is on the left side of each box. This means both distributions are right-skewed. Also bening tumors between 1 and 4 while malignant tumors between 7 and 10. Benign cases mostly have Cl. thickness values between 1 and 4, while malignant cases are mostly between 5 and 10.

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