

# **American International University of Bangladesh**

## **Mid-Term Project Report**

### **Introduction to Data Science**

**Section: C** 

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Faculty

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#### **Summary**

The Titanic dataset is a comprehensive and diverse collection of structured data that offers immense potential for research, analysis, and development within the [specific domain/subject].

Its size, coverage, and data types provide a rich resource for exploring various phenomena and building robust models.

The dataset includes various attributes for each passenger, their age, class, gender, etc. to predict if they would have survived or not. The dataset contains a total of 251 rows or instances, representing the passengers on board the Titanic. It provides valuable information for analyzing various aspects related to the survival rate of passengers, including factors such as passenger class, age, and gender.

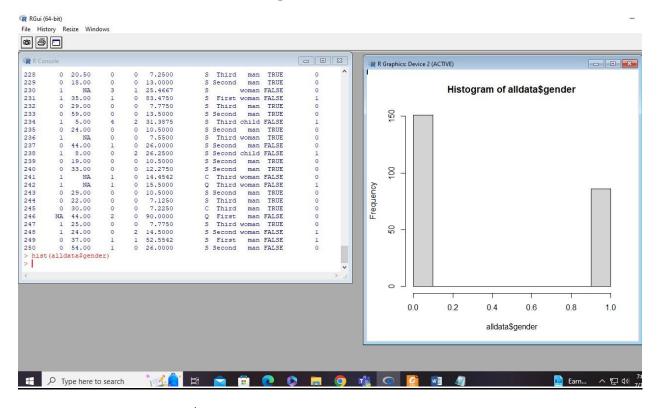
#### **Importing Titanic.csv file into R studio**



Here I have imported the dataset Titanic.csv using the "read.csv" command in the parameter I had provided the location of dataset Titanic.csv in my computer file.

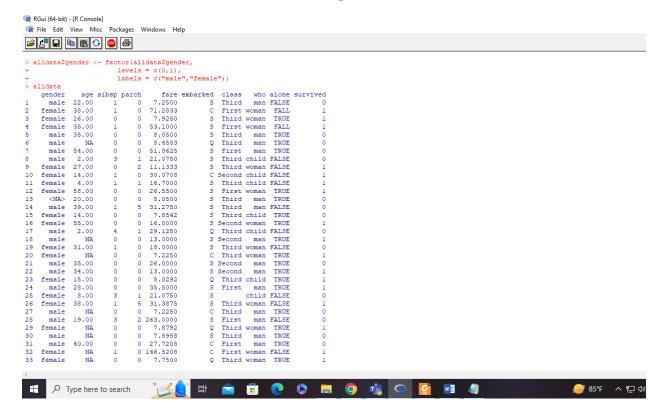
Then I ensured the parameter of a header by giving true and making it in comma Delimited Text Data Set I used sep = "," and to check all the data from the data set I wrote all data. The median is the value of the central point in the distribution.

### **Histogram of Dataset**



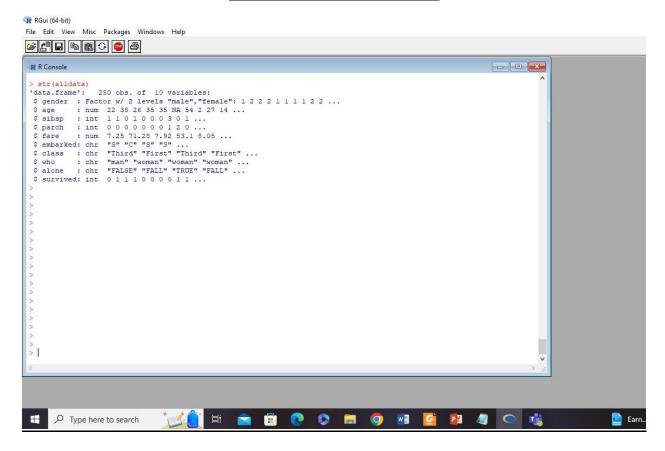
Here, using hist(alldata\$gender) command to get the histogram of values of gender attribute from Titanic.csv dataset.

#### **Annotating Dataset**



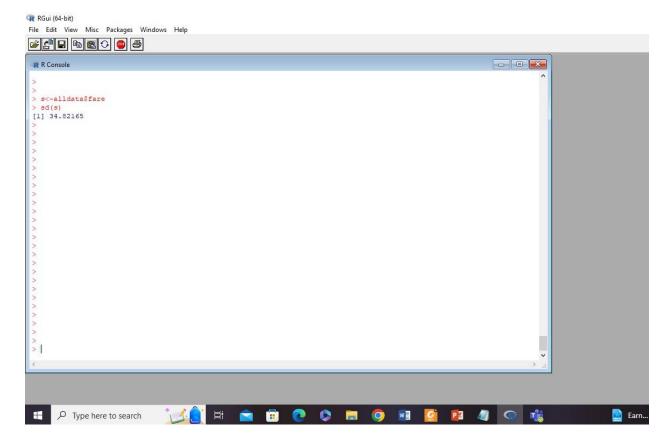
Here, annotating a dataset involves adding descriptive information, metadata, and contextual details to enhance the understanding, usability, and reliability of the data. By providing clear variable descriptions, data source information, preprocessing details, quality assessments, metadata, usage guidelines, and versioning information, dataset annotation ensures that users can effectively interpret, analyze, and make informed decisions based on the dataset. According to this code, the factor() function can be used to create value labels for categorical variables. Continuing for the above code example, say that I have a variable named gender, which is coded 0 for male and 1 for female.

#### **Structure of Dataset**



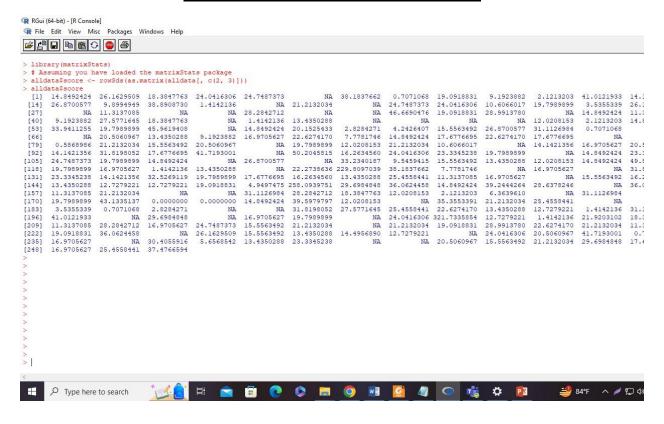
Here, I have used 'str()' command which shows the summary of the dataset and It shows from 150 observations of 10 variables.

#### **Standard Deviation**



A standard deviation (or  $\sigma$ ) is a measure of how dispersed the data is in relation to the mean. Low standard deviation means data are clustered around the mean and high standard deviation indicates data are more spread out. Here, from the fare attribute data I have calculated the deviation value is 34.82165 more spread out. So, it is a high standard deviation

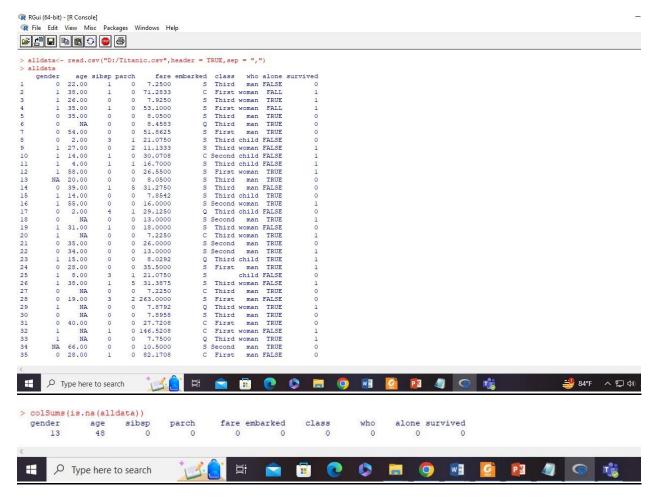
#### Raw wise standard deviation



The row-wise standard deviation in a dataset indicates the variability or spread of values within each row or observation of the dataset. It provides information about how much the individual values within a row deviate from the mean of that row.

By using alldata\$score <- rowSds(as.matrix(alldata[, c(2, 3)])) and > alldata\$score command gives the standard deviation in row wise standard deviation.

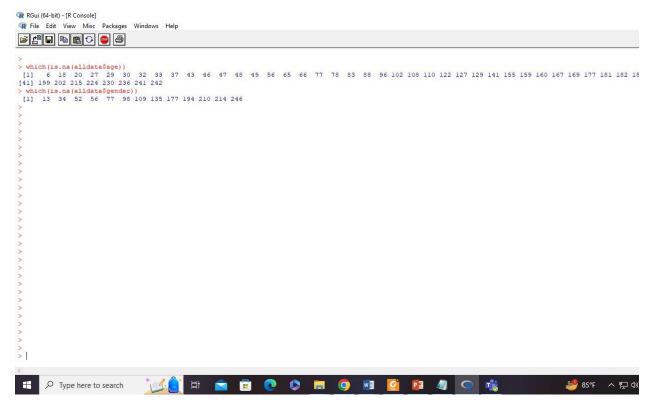
### **Counting Missing Values**



Here I have counted missing values on the dataset Titanic.csv. and found that gender has 13 and age is 48 missing values.

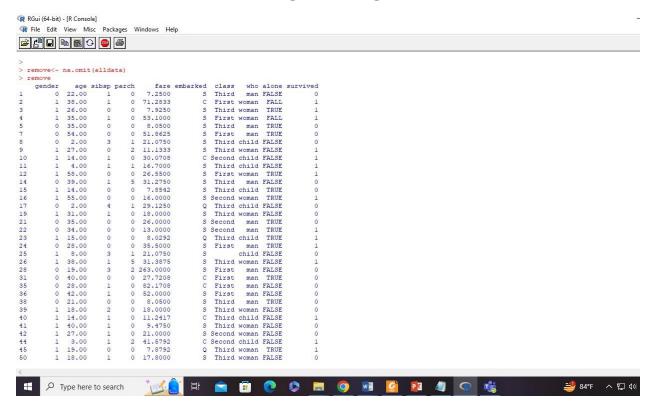
In R, the NA symbol is used to define the missing values and to represent impossible arithmetic operations (like dividing by zero) we use the NAN symbol which stands for "not a number". In simple words, we can say that both NA or NAN symbols represent missing values in R.

## **Finding Missing Values**



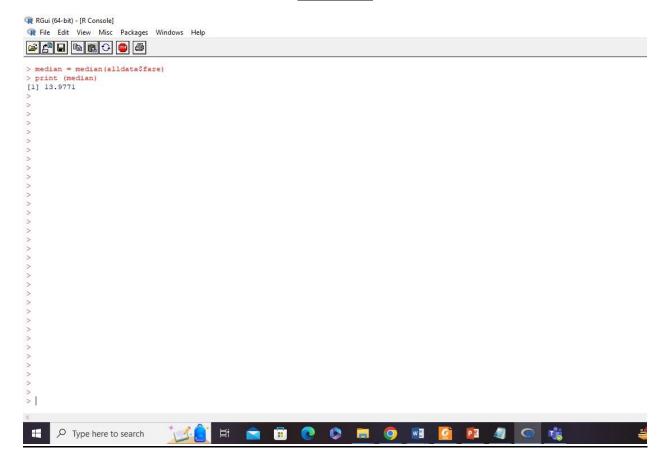
After counting missing values I got that gender has 13 and age is 48 missing values. And finding those by using >which(is.na(alldata\$age)) and using >which(is.na(alldata\$gender)) command in r studio.

#### **Removing Missing Values**



Here in R studio, I have used remove<-na. omit() command to remove all observations with missing data on ANY variable in the dataset, or use subset() to filter out cases that are missing on a subset of variables. Though there are many but I used this command to remove NA values in Titanic.csv file.

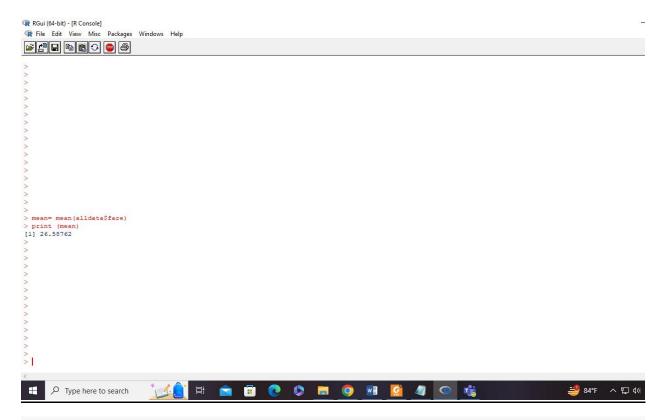
#### **Median**



Here, I have used the median = median(alldata\$fare) command to calculate the median value from the Titanic.csv file, and using print (median) I have the median value which is 13.9771 which indicates the central tendency of fare price from all the values from fare attribute

The median is useful because it is not affected by extreme values (outliers) to the same extent as the mean. It provides a robust measure of the center of the dataset, particularly in situations where the distribution is not symmetrical or when the data contains extreme values that could skew the mean.

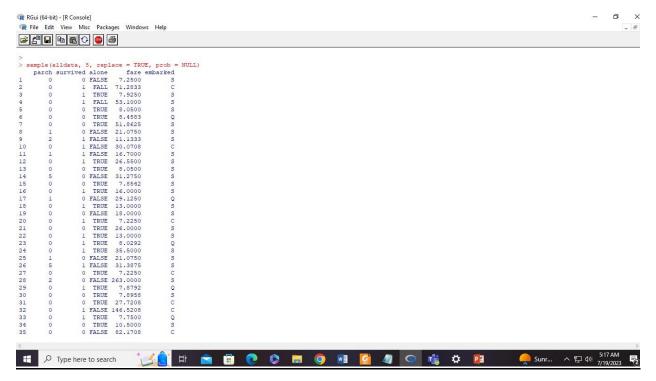
#### Mean



The mean, also known as the average, is another measure of central tendency commonly used in statistics. Unlike the median, which represents the middle value of a dataset, the mean is the sum of all the values divided by the total number of observations.

From the values of the fare attribute, we can see that it is 26.58762 which is calculated from the mean value generating command mean= mean(alldata\$fare) and printed that value with print (mean)

### **Sampling**



Sampling in a dataset refers to the process of selecting a subset of observations or data points from a larger population or dataset. The purpose of sampling is to obtain a representative sample that can provide insights or make inferences about the entire population. Thee code sample(alldata, 5, replace = TRUE, prob = NULL) randomly selects 5 observations from the titanic.csv dataset, allowing replacement (an observation can be chosen more than once), and each observation has an equal chance of being selected. The resulting output will be a sample of 5 observations from the Titanic.csv dataset.