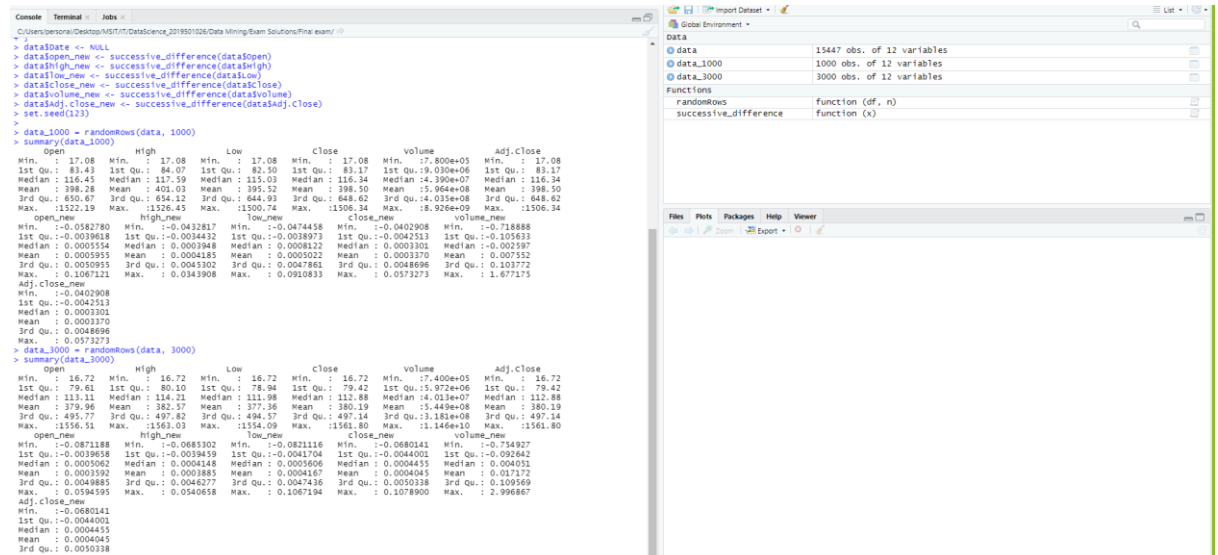


## Data Mining - Lab Exam – Remedial

1. a) For the dataset BSE\_Sensex\_Index.csv, create an extra column of successive differences for each column of numeric values in this data file. Extract two simple random samples with replacement of 1000 and 3000 observations (rows). Show your R commands for doing this. Do the same thing by using Excel. Show your Excel commands.

**Note:** Successive difference for date d1= (date d1 value-previous date of d1 value)/previous date of d1. For the last row fill up values with mean of its immediate three previous row values.



The screenshot shows the R Studio interface with the following components:

- Console:** Contains R commands and their output. The commands include:
  - `dataDate <- NULL`
  - `dataOpen_new <- successive_difference(data$open)`
  - `dataHigh_new <- successive_difference(data$high)`
  - `dataLow_new <- successive_difference(data$low)`
  - `dataClose_new <- successive_difference(data$close)`
  - `dataVolume_new <- successive_difference(data$volume)`
  - `dataAdjClose_new <- successive_difference(data$adjClose)`
  - `set.seed(123)`
  - `data_1000 <- randomRows(data, 1000)`
  - `summary(data_1000)`
  - `data_3000 <- randomRows(data, 3000)`
  - `summary(data_3000)`
- Global Environment:** Lists the datasets: `data` (15447 obs., 12 variables), `data_1000` (1000 obs., 12 variables), and `data_3000` (3000 obs., 12 variables).
- Functions:** Lists the functions: `randomRows` (Function (df, n)) and `successive_difference` (Function (x)).
- Files, Plots, Packages, Help, Viewer:** Standard R Studio tabs.

- b) For your samples, use the functions `mean()`, `max()`, `var()` and `quantile(.,25)` to compute the mean, maximum, variance and 1st quartile respectively for each column which has successive differences. Show your R code and the resulting values. Do the same thing by using Excel. Show your Excel commands.

```
> mean(data_1000$open_new)
[1] 0.0005955025
> mean(data_1000$high_new)
[1] 0.0004184797
> mean(data_1000$low_new)
[1] 0.0005022487
> mean(data_1000$close_new)
[1] 0.0003369592
> mean(data_1000$volume_new)
[1] 0.007551912
> mean(data_1000$Adj.close_new)
[1] 0.0003369592
> var(data_1000$open_new)
[1] 8.714339e-05
> var(data_1000$high_new)
[1] 6.119132e-05
> var(data_1000$low_new)
[1] 8.313995e-05
> var(data_1000$close_new)
[1] 7.637739e-05
> var(data_1000$volume_new)
[1] 0.0327711
> var(data_1000$Adj.close_new)
[1] 7.637739e-05
> max(data_1000$open_new)
[1] 0.1067121
> max(data_1000$high_new)
[1] 0.03439077
> max(data_1000$low_new)
[1] 0.09108332
> max(data_1000$close_new)
[1] 0.05732732
> max(data_1000$volume_new)
[1] 1.677175
> max(data_1000$Adj.close_new)
[1] 0.05732732
> quantile(data_1000$open_new,0.25)
      25%
-0.003961827
> quantile(data_1000$high_new,0.25)
      25%
-0.003443228
> quantile(data_1000$low_new,0.25)
      25%
-0.003897353
> quantile(data_1000$close_new,0.25)
      25%
-0.004251294
> quantile(data_1000$volume_new,0.25)
      25%
-0.1056329
> quantile(data_1000$Adj.close_new,0.25)
      25%
-0.004251294
>
```

```

>
> mean(data_3000$open_new)
[1] 0.0003591911
> mean(data_3000$high_new)
[1] 0.0003884621
> mean(data_3000$low_new)
[1] 0.0004167
> mean(data_3000$close_new)
[1] 0.0004044752
> mean(data_3000$volume_new)
[1] 0.0171718
> mean(data_3000$Adj.close_new)
[1] 0.0004044752
> var(data_3000$open_new)
[1] 8.509529e-05
> var(data_3000$high_new)
[1] 6.81047e-05
> var(data_3000$low_new)
[1] 8.768766e-05
> var(data_3000$close_new)
[1] 8.588174e-05
> var(data_3000$volume_new)
[1] 0.03939109
> var(data_3000$Adj.close_new)
[1] 8.588174e-05
> max(data_3000$open_new)
[1] 0.05945946
> max(data_3000$high_new)
[1] 0.05406578
> max(data_3000$low_new)
[1] 0.1067194
> max(data_3000$close_new)
[1] 0.10789
> max(data_3000$volume_new)
[1] 2.996867
> max(data_3000$Adj.close_new)
[1] 0.10789
> quantile(data_3000$open_new,0.25)
      25%
-0.003965834
> quantile(data_3000$high_new,0.25)
      25%
-0.003945885
> quantile(data_3000$low_new,0.25)
      25%
-0.004170403
> quantile(data_3000$close_new,0.25)
      25%
-0.00440009
> quantile(data_3000$volume_new,0.25)
      25%
-0.09264194
> quantile(data_3000$Adj.close_new,0.25)
      25%
-0.00440009
> |

```

c) Compute the same quantities in part b on the entire data set and show your answers. How much do they differ from your answers in part b? Do you find any significant difference

between two sample values like mean in comparison with entire data? If so what explanation you can give for that?

Do the same thing by using Excel. Show your Excel commands.

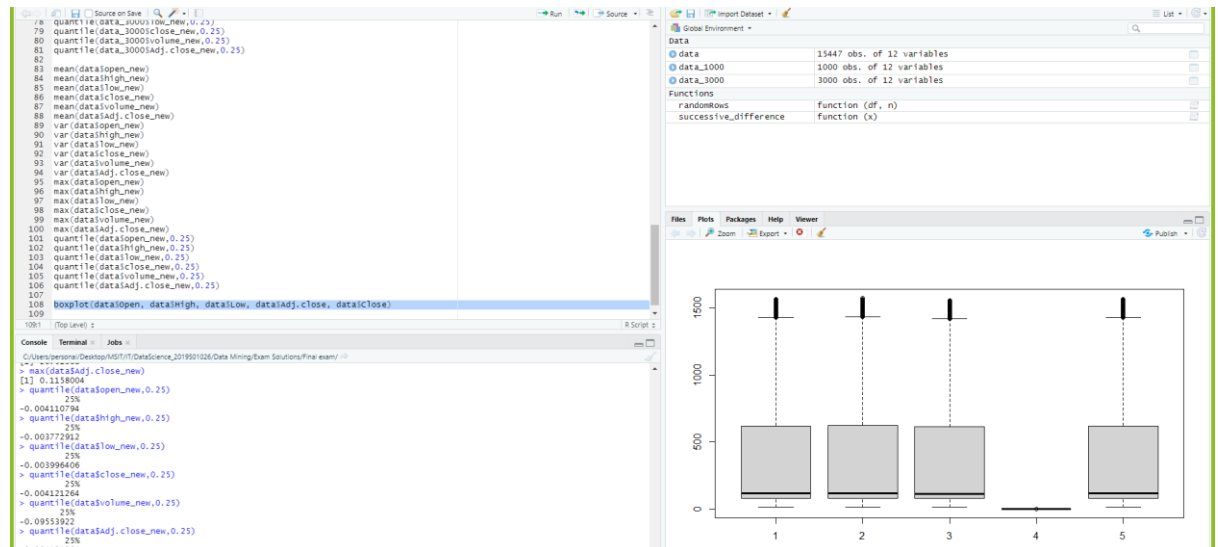
```

C:/Users/personal/Desktop/MSIT/IT/DataScience_2019501026/Data Mining/Exam Solutions/Final exam/
-0.00440009
> mean(data$open_new)
[1] 0.000329528
> mean(data$high_new)
[1] 0.0003188991
> mean(data$low_new)
[1] 0.0003266191
> mean(data$close_new)
[1] 0.0003303709
> mean(data$volume_new)
[1] 0.02062874
> mean(data$Adj.close_new)
[1] 0.0003303709
> var(data$open_new)
[1] 9.027493e-05
> var(data$high_new)
[1] 6.939914e-05
> var(data$low_new)
[1] 8.646474e-05
> var(data$close_new)
[1] 9.350347e-05
> var(data$volume_new)
[1] 0.09080738
> var(data$Adj.close_new)
[1] 9.350347e-05
> max(data$open_new)
[1] 0.1067121
> max(data$high_new)
[1] 0.08037943
> max(data$low_new)
[1] 0.1067194
> max(data$close_new)
[1] 0.1158004
> max(data$volume_new)
[1] 26.51968
> max(data$Adj.close_new)
[1] 0.1158004
> quantile(data$open_new,0.25)
25%
-0.004110794
> quantile(data$high_new,0.25)
25%
-0.003772912
> quantile(data$low_new,0.25)
25%
-0.003996406
> quantile(data$close_new,0.25)
25%
-0.004121264
> quantile(data$volume_new,0.25)
25%
-0.09553922
> quantile(data$Adj.close_new,0.25)
25%
-0.004121264
> |

```

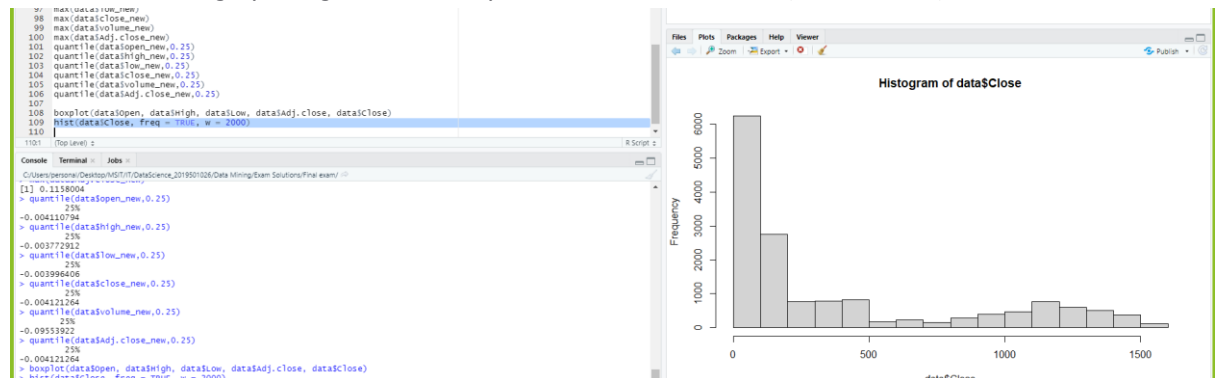
d) Use R to produce a single graph displaying a boxplot for open, close, high and low. Include the R commands and the plot.

Do the same thing by using Excel. Show your Excel commands



e) Use R to produce a frequency histogram for Close values. Use intervals of width 2000 beginning at 0. Include the R commands and the plot.

Do the same thing by using Excel. Show your Excel commands. (10+10=20M)



- Implement Apriori Algorithm or use built in packages to find out the frequent itemsets and generate rules for frequent itemsets. Trace program output for the following given dataset of transactions with a minimum support of 3 and submit. (10M)

TID, Items
101, A,B,C,D,E
102, A,C,D
103, D,E
104, B,C,E
105, A,B,D,E
106, A,B
107, B,D,E
108, A,B,D
109, A,D
110, D,E



```

> setwd("C:\\Users\\person\\Desktop\\MSIT\\IT\\DataScience_201901026\\Data Mining\\Exam Solutions\\Final exam")
> data = read.csv("C:\\Users\\person\\Desktop\\MSIT\\IT\\DataScience_201901026\\Data Mining\\Exam Solutions\\Final exam\\Liver_data.csv", header = FALSE, col.names = c("mcv", "alkphos", "sgpt", "sgot", "gammagt", "drinks", "selector"))
> data$drinks = cut(data$drinks, breaks = c(0,5,10,15,20,25), labels = c("C1", "C2", "C3", "C4", "C4"), right = FALSE)
> data = na.omit(data)
> #training and test sets
> traindata = subset(data, data$selector == 1)
> testdata = subset(data, data$selector == 2)
> x_train = subset(traindata, select = -c(selector, drinks))
> x_test = subset(testdata, select = -c(selector, drinks))
> y_train = traindata[,6, drop = TRUE]
> y_test = testdata[,6, drop = TRUE]
> #For Training Data
> library(class)
> model1 = knn(x_train, x_test, y_train, k = 1)
> 1-sum(y_train==model1)/length(y_train)
[1] 0.2827586
Warning messages:
1: In knn(x_train, x_test, y_train, model1) :
  longer object length is not a multiple of shorter object length
2: In is.na(e1) | is.na(e2) :
  longer object length is not a multiple of shorter object length
> model2 = knn(x_train, x_test, y_train, k = 2)
> 1-sum(y_train==model2)/length(y_train)
[1] 0.310945
> model3 = knn(x_train, x_test, y_train, k = 3)
> 1-sum(y_train==model3)/length(y_train)
[1] 0.2
> #For Test Data
> model4 = knn(x_train, x_test, y_test, k = 1)
> 1-sum(y_test==model4)/length(y_test)
[1] 0.445
> model5 = knn(x_train, x_test, y_test, k = 2)
> 1-sum(y_test==model5)/length(y_test)
[1] 0.43
> model6 = knn(x_train, x_test, y_test, k = 3)
> 1-sum(y_test==model6)/length(y_test)
[1] 0.42
>

```

5. Use Support Vector machine for above problem. And compare the performance of both. Explain the input parameters you provided for the classifier. (10M)

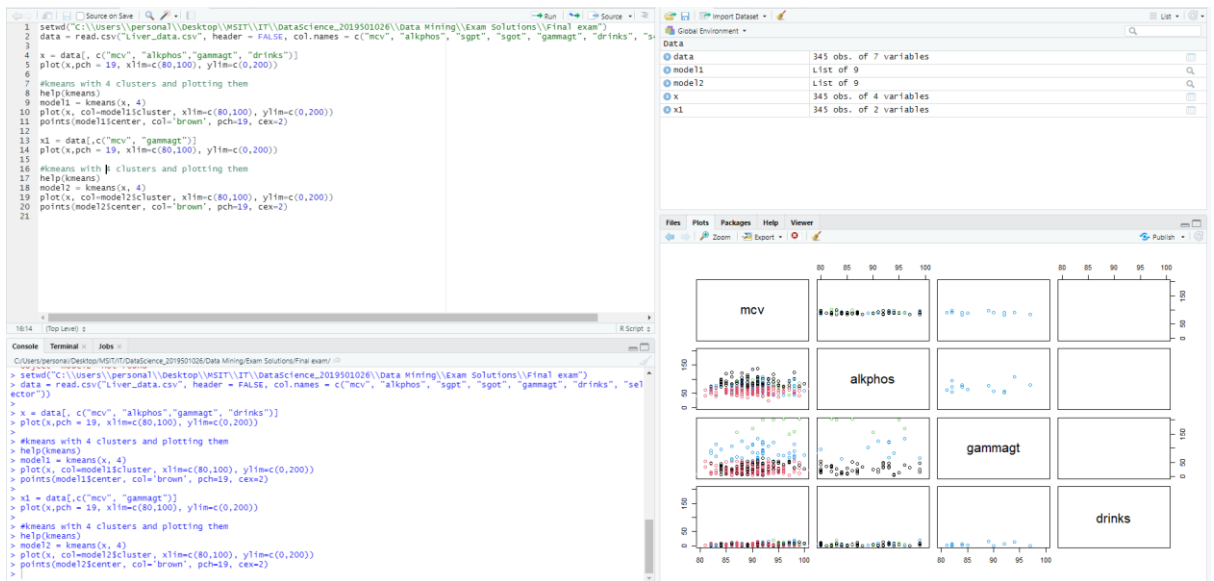
```

1 data = read.csv("Liver_data.csv", header = FALSE, col.names = c("mcv", "alkphos", "sgpt", "sgot", "gammagt", "drinks", "selector"))
2 data$drinks = cut(data$drinks, breaks = c(0,5,10,15,20,25), labels = c("C1", "C2", "C3", "C4", "C4"), right = FALSE)
3 data = na.omit(data)
4 #training and test sets
5 traindata = subset(data, data$selector == 1)
6 testdata = subset(data, data$selector == 2)
7 x_train = subset(traindata, select = -c(selector, drinks))
8 x_test = subset(testdata, select = -c(selector, drinks))
9 y_train = traindata[,6, drop = TRUE]
10 y_test = testdata[,6, drop = TRUE]
11 library(e1071)
12 #For training
13 model = svm(x_train, y_train)
14 1-sum(y_train==predict(model,x_train))/length(y_train)
15 #For test data
16 1-sum(y_test==predict(model,x_test))/length(y_test)
17

```

6. Create k-means clusters for k=4 for the Liver Disorders Data Set at <http://archive.ics.uci.edu/ml/datasets/Liver+Disorders> . Explain the input parameters you provided for the clustering algorithm. Plot the fitted cluster centers using a different color. Finally assign the cluster membership for the points to the nearest cluster center. Color the points according to their cluster membership. (10+10=20M)





7. Compute the misclassification error that would result if you used your clustering rule to classify the data by assigning the majority class of the cluster.
- (10M)
8. Consider the dataset `BSE_Sensex_Index.csv`. Create an extra column of successive growth rate for column close where the successive growth rate is defined as  
(value of day x- value of day x-1)/value of day x-1. Use a z score cut off of 3 to identify any outliers. List the respective dates from the csv file on which day these outliers fall.

(10M)

