Ex No:1	Basics of R – data types, vectors, factors, list and data
Date:	frames

AIM:

To implement and understand the basics of R programming with its data types, vectors, factors, list and data frames.

ALGORITHM:

- 1. Start
- **2.** Assign values in logical, numerical, character, complex and character in raw form to a variable v.
- 3. Print the class of v.
- **4.** Assign a vector for subject Names, temperature and flu_status for three patients using c() function and access the elements.
- **5.** Create a factor using factor() with duplicate values and assign level with distinct values.
- **6.** Display the specific element and check for certain values in factor.
- 7. Create a list using list() from the patient details and access the multiple elements.
- **8.** Create a data frame using data.frame() with multiple vectors as features. Access the elements.
- **9.** Create a matrix using matrix() with different allocations and access the elements.
- **10.** Stop.

PROGRAM:

```
#Data Types
v<-TRUE
print(class(v))
v < -23.5
print(class(v))
v < -2L
print(class(v))
v < -2 + 5i
print(class(v))
v<-"TRUE"
print(class(v))
v<-charToRaw("Hello")
print(class(v))
#Vectors
subject_name<-c("John Doe","Jane Doe","Steven Grant")</pre>
temperature <- c(98.1,98.6,101.4)
flu_status<-c(FALSE,FALSE,TRUE)
temperature[2]
temperature[2:3]
temperature[-2]
#Factors
gender<-factor(c("MALE","FEMALE","MALE"))</pre>
blood<-factor(c("O","AB","A"),levels=c("A","B","AB","O"))
```

```
blood[1:2]
symptoms<-factor(c("SEVERE","MILD","MODERATE"),
         levels=c("MILD","MODERATE","SEVERE"),
         ordered=TRUE)
symptoms>"MODERATE"
#Lists
subject1<-list(fullname=subject_name[1],</pre>
        temperature=temperature[1],
        flu_status=flu_status[1],
        gender=gender[1],
        blood=blood[1],
        symptoms=symptoms[1])
subject1
subject1[2]
subject1[[2]]
subject1$temperature
subject1[c("temperature","flu_status")]
#Data Frames
pt_data<-data.frame(subject_name, temperature, flu_status,
           gender, blood, symptoms)
pt_data
pt_data$subject_name
pt_data[c("temperature","flu_status")]
pt_data[c(1,2),c(2,4)]
pt_data[,1]
pt_data[,]
#Matrices
m < -matrix(c(1,2,3,4),ncol=2)
print(m)
m < -matrix(c(1,2,3,4,5,6),nrow=3)
print(m)
print(m[1,])
print(m[1,])
thismatrix <- matrix(c("apple", "banana", "cherry", "orange"), nrow = 2, ncol = 2)
for (rows in 1:nrow(thismatrix)) {
 for (columns in 1:ncol(thismatrix)) {
  print(thismatrix[rows, columns])
 }
```

OUTPUT:

```
File Edit Selection View Go Run Terminal Help
                   PROBLEMS 73
 d)
                  [1] "logical"
[1] "numeric"
[1] "integer"
[1] "complex"
[1] "character"
[1] "raw"
[1] 98.6
[1] 98.6 101.4
[1] 98.1 101.4
[1] MALE FEMALE MALE
Levels: FEMALE MALE
[1] O AB
Levels: A B AB O
[1] TRUE FALSE FALSE
$fullname
[1] "John Doe"
 ရှ
                                                FEMALE MALE
 R
                    $temperature
[1] 98.1
                    $flu_status
[1] FALSE
                    [1] MALE
Levels:
                                       FEMALE MALE
                     [1] O
Levels: A B AB O
                    $symptoms
[1] SEVERE
Levels: MILD < MODERATE < SEVERE
                     $temperature
[1] 98.1
                    [1] 98.1
[1] 98.1
$temperature
[1] 98.1
                    $flu_status
[1] FALSE
                    subject_name temperature flu_status gender blood symptoms

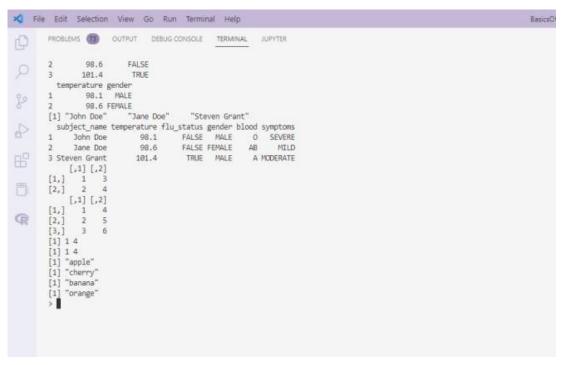
1 John Doe 98.1 FALSE MALE O SEVERE

2 Jane Doe 98.6 FALSE FEMALE AB MILD

3 Steven Grant 101.4 TRUE MALE A MODERATE

[1] "John Doe" "Jane Doe" "Steven Grant"

temperature flu_status
                  [1]
                                 98.1 FALSE
98.6 FALSE
101.4 TRUE
```



Result:

Thus the R Script program to implement various data types, vectors, factors, lists and data frames is executed successfully and the output is verified.

Ex no: 2	Diagnosis of Breast Cancer using KNN.
Date:	

Aim:

To implement a R program to predict and diagnose Breast Cancer using KNN algorithm.

Algorithm:

- 1. Start
- 2. Read the csv file from the directory and store it in bcd variable.
- 3. Drop the first column id.
- 4. Change the diagnosis feature with categorical values B and M in a factor
- 5. Normalize the dataset.
- 6. Split the dataset for training and testing, with diagnosis as the response variable and the rest as the predictor variables.
- 7. Import the library "class" for knn classification.
- 8. Predict the knn model using knn() with 5 clusters with the corresponding training and testing data.
- 9. Display the confusion matrix and accuracy of the knn model.
- 10. Stop

PROGRAM:

```
bcd<-read.csv("../input/breast-cancer-dataset/Breast_Cancer.csv", stringsAsFactors=FALSE)
bcd<-bcd[-1]
bcd$diagnosis<-factor(bcd$diagnosis, levels=c("B","M"), labels=c("Benign","Malignant"))
normalize<-function(x){
    return (x-min(x)) / (max(x)-min(x))
}
bcd_n <- as.data.frame(lapply(bcd[2:31], normalize))
x_train <- bcd_n[1:469,]
x_test <- bcd_n[470:569,]
y_train <- bcd[1:469,1]
y_test <- bcd[470:569,1]
library(class)
y_pred<-knn(train=x_train,test=x_test,cl=y_train,k=5)
tbl=table(x=y_test,y=y_pred)
tbl
accuracy = sum(diag(tbl))
```

OUTPUT:

```
PROBLEMS 37 OUTPUT DEBUG CONSOLE
                                                                            TERMINAL

        $ perimeter mean
        : num
        78.8 69.3 76.9 73 97.7 ...

        $ area mean
        : num
        464 346 373 385 712 ...

        $ smoothness mean
        : num
        0.1028 0.0969 0.1077 0.1164 0.0796 ...

        $ compactness mean
        : num
        0.0698 0.1147 0.078 0.1136 0.0693 ...

        $ concavity mean
        : num
        0.0399 0.0639 0.0305 0.0464 0.0339 ...

  $ points_mean
$ symmetry_mean
                                          num 0.037 0.0264 0.0248 0.048 0.0266 ...
num 0.196 0.192 0.171 0.177 0.172 ...
                                      : num 0.8595 0.0649 0.0634 0.0667 0.0554 ...
: num 0.236 0.451 0.197 0.338 0.178 ...
: num 0.666 1.197 1.387 1.343 0.412 ...
: num 1.67 3.43 1.34 1.85 1.34 ...
  $ dimension_mean
  $ radius se
  $ texture se
  $ perimeter_se
                                       : num 17.4 27.1 13.5 26.3 17.7 ...

: num 0.00805 0.00747 0.00516 0.01127 0.00501 ...

: num 0.0118 0.03581 0.00936 0.03498 0.01485 ...
  $ area se
  $ compactness se
                                       num 0.0168 0.0335 0.0106 0.0219 0.0155 ...
: num 0.01241 0.01365 0.00748 0.01965 0.00915 ...
  $ concavity_se
  $ points se
  $ symmetry_se
$ dimension_se
                                      : num 0.0192 0.035 0.0172 0.0158 0.0165 ...
: num 0.00225 0.00332 0.0022 0.00344 0.00177 ...
  $ radius_worst
$ texture_worst
                                       : num 13.5 11.9 12.4 11.9 16.2 ...
: num 15.6 22.9 26.4 15.8 15.7 ...
 $ perimeter_worst : num 87 78.3 79.9 76.5 104.5 ... $ area_worst : num 549 425 471 434 819 ... $ smoothness_worst : num 0.139 0.121 0.137 0.137 0.113 ... $ compactness_worst : num 0.127 0.252 0.148 0.182 0.174 ...
 Benign Malignant
Malignant 4
[1] "Accuracy 96"
```

Result:

Thus the R Script program to implement diagnosis of Breast Cancer using K-Nearest Neighbour algorithm is executed successfully and the output is verified.

Ex No: 3	Filtering Mobile phone spam using Naïve Bayes
Date:	

AIM:

To implement a R program to Filter Mobile phone spam using Naïve Bayes.

ALGORITHM:

- 1. Start
- **2.** Import the csv file and store the dataframe in "Sms". Have a glimpse at the structure of the data frame.
- 3. Remove the unneccesary columns which is from column 3 to 5.
- **4.** Convert the labels as factors.
- **5.** Remove special characters from the dataset and retain only alpha numeric characters using alnum in str_replace_all() from "stringr" package.
- **6.** Create a volatile corpus VCorpus() for text mining from the source object of "v2" which is extracted using VectorSource().
- **7.** Create a DocumentTermMatrix() to split the SMS message into individual Components.
- **8.** Create training and testing dataset with the split ratio 0.75.
- **9.** Find the frequent terms which appear for atleast 5 times in DocumentTermMatrix in training and testing dataset respectively.
- 10. Train the model using naiveBayes() from e1071 library.
- 11. Evaluate the model Performance.
- **12.** Print the confusion matrix and Accuracy of the model.
- **13.** Stop.

PROGRAM:

```
sms <- read.csv("../input/spam-ham-dataset/spam.csv", stringsAsFactors=FALSE)
str(sms)
sms <-sms[-3:-5]
sms$v1 <- factor(sms$v1)
library(stringr)
sms$v2 = str_replace_all(sms$v2, "[^[:alnum:]]", " ") %>% str_replace_all(.,"[]+", " ")
library(tm)
sms_corpus <- VCorpus(VectorSource(sms$v2))</pre>
```

```
print(sms_corpus)
print(as.character(sms_corpus[[6]]))
sms_dtm <- DocumentTermMatrix(sms_corpus, control = list</pre>
(tolower=TRUE, removeNumbers=TRUE, stopwords=TRUE, removePunctuations=TRUE, stemmi
ng=TRUE))
x_train <- sms_dtm[1:4169, ]</pre>
x_test <- sms_dtm[4170:5572, ]
y_train <- sms[1:4169, ]$v1</pre>
y_test <- sms[4170:5572, ]$v1</pre>
sms_freq_word_train <- findFreqTerms(x_train, 5)</pre>
sms freq word test <- findFreqTerms(x test, 5)</pre>
x_train<- x_train[ , sms_freq_word_train]</pre>
x_test <- x_test[ , sms_freq_word_test]</pre>
convert_counts <- function(x) \{x \leftarrow ifelse(x > 0, "Yes", "No")\}
x_train <- apply(x_train, MARGIN = 2,convert_counts)</pre>
x_test <- apply(x_test, MARGIN = 2,convert_counts)</pre>
library(e1071)
model <- naiveBayes(x_train, y_train,laplace=1)</pre>
y_pred <- predict(model, x_test)</pre>
cm = table(y_pred, y_test)
print(cm)
acc = sum(diag(cm))/sum(cm)
print(paste("Accuracy: ",acc*100,"%"))
```

OUTPUT:

RESULT:

Thus the R program to implement filtering of Mobile phone spam using Naïve Bayes is executed successfully and the output is verified.

Ex No:4	Risky Bank Loans using Decision Trees
Date:	

AIM:

To implement a R program to find Risky Bank loans using Decision Tree.

ALGORITHM:

- 1. Start
- 2. Import the dataset credit.csv and display the structure of the dataset.
- 3. Display the table to find the range of values and find the missing values.
- **4.** Factorise the default column and set seed of 123.
- **5.** Split the dataset for training and testing in the ratio of 0.8, with "default" as the response variable, and the rest as predictor variables.
- **6.** Import the library C5.0 for implementing decision tree.
- 7. Train the decision tree model using C5.0 function for the training dataset.
- **8.** Test the model to predict using predict(). Print the confusion matrix.
- **9.** Print the accuracy of the decision tree model.
- **10.** Stop

PROGRAM:

```
credit <- read.csv("credit.csv")

str(credit)

table(credit$savings_balance)

summary(credit$amount)

credit$default <- factor(credit$default)

set.seed(123)

train_sample <- sample(1000, 800)

str(train_sample)

x_train <- credit[train_sample, -17]

x_test <- credit[-train_sample, -17]

y_train <- credit[train_sample, 17]

y_test <- credit[-train_sample, 17]

library(C50)

model <- C5.0(x_train,y_train)
```

```
summary(model)

y_pred <- predict(model,x_test)

cm = table(y_pred,y_test)

print(cm)

acc=sum(diag(cm))/sum(cm)

print(paste("Accuaracy: ",acc*100,"%"))</pre>
```

OUTPUT:

```
: : ...checking_balance = < 0 DM: yes (A)
: : crbcking_balance = < 0 DM: yes (A)
: : crbcking_balance = - 200 DM: no (3/1)
: purpose = furniture/appliances:
: ...sardings_balance in (so = 100 DM: yes (6)
: sardings_balance = < 100 DM: yes (6)
: ...months_loan_duration < 22: yes (12/1)
: months_loan_duration < 22: yes (12/1)
: months_loan_duration < 22: yes (12/1)
: amount < 2325: yes (3)
: amount < 2325: yes (3)
: amount < 2325: yes (6)
: sardings_balance in (los = 500 DM; no (6)
: sardings_balance in (los = 500 DM; no (8)
: sardings_balance = (100 DM: 900 DM; no (8)
: ...job in (management_unskilled,
: : ...dependents > 1: no (3/1)
: dependents > 1: no (3/1)
: dependents < -1:
: ....months_loan_duration < 22: yes (8)
: months_loan_duration > 22: yes (8)
: months_loan_duration > 22: yes (8)
: months_loan_duration > 7 years:
: ...order_duration > 7 years:
: ...order_curation > 7 years:
:
```

```
Decision Tree

Size Errors

69 99(11.0%) <<

(a) (b) <-classified as

625 10 (a): class no
89 176 (b): class yes

Attribute usage:

100.00% checking balance
54.22% credit, history
48.22% months loan duration
42.22% savings, balance
31.00% purpose
21.33% employment duration
9.22% years, at residence
8.70% housing
8.40% jbs
6.11% other_credit
```

RESULT:

Thus the R program to find Risky Bank loans using Decision Tree is executed successfully and the output is verified.

Ex No: 5	
	Medical Expense with Linear Regression.
Date:	•

AIM:

To implement a R program to predict Medical Expense using Linear Regression

ALGORITHM:

- 1. Start
- **2.** Load the Insurance dataset and analyse the structure of the dataset.
- **3.** Get the summary statistics. Check whether the distribution is right-skewed or left skewed by comapring the mean and median. Verify the same using histogram.
- **4.** Check the distribution of "region" using table.
- **5.** Create a correlation matrix of "age", "bmi", "children", "expenses".
- **6.** To determine the pattern of the dataset, use scatterplot using pairs() for "age", "bmi", "children", "expenses".
- 7. To display a more informative scatterplot use pairs.panel() from "psych" library.
- **8.** Fit the linear regression model using lm() with expenses as the dependent variable.
- **9.** Evaluate the model performance using summary().
- **10.** To improve the model performance, square the age variable as age2 and bmi30 is 1 if bmi>=30 else 0.
- **11.** Train the model with age + age2+bmi30 as also as the independent variables.
- **12.** Evaluate the model performance for model2 using summary().
- **13.** Stop.

PROGRAM:

```
insurance<-read.csv("insurance.csv",stringsAsFactors = TRUE)
str(insurance)
summary(insurance$expenses)
hist(insurance$expenses)
table(insurance$region)
cor(insurance[c("age","bmi","children","expenses")])
pairs(insurance[c("age","bmi","children","expenses")])
library(psych)
pairs.panels(insurance[c("age","bmi","children","expenses")])8
ins_model <- lm(expenses ~ age + children + bmi + sex + smoker + region, data = insurance)
ins_model</pre>
```

summary(ins_model)

insurance\$age2 <- insurance\$age^2

insurance\$bmi30 <- ifelse(insurance\$bmi >= 30,1,0)

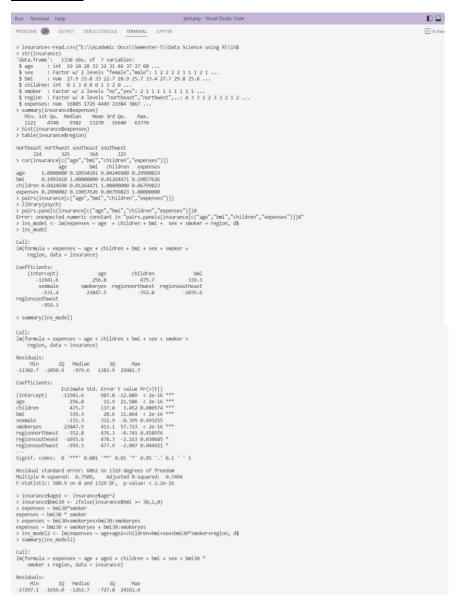
expenses ~ bmi30*smoker

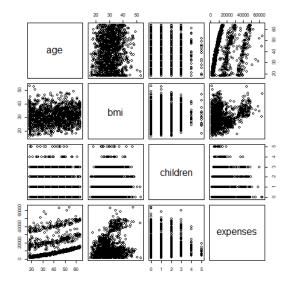
expenses ~ bmi30+smokeryes+bmi30:smokeryes

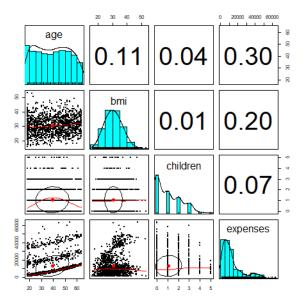
ins_model2 <- lm(expenses ~ age+age2+children+bmi+sex+bmi30*smoker+region, data=insurance)

summary(ins_model2)

OUTPUT:







RESULT:

Thus the R program to predict medical expenses using linear regression is executed successfully and the output is verified.

Ex No:6 Date:	Modeling the strength of concrete with ANNs

AIM:

To create a R program to implement Modeling steength of concrete with ANNs.

ALGORITHM:

- 1. Start
- 2. Read the dataset "Concrete" using read.csv() function and store it in concrete variable.
- **3.** To perform calculations with ease, normalise the data using an user-defined function 'normalise' and apply it to the dataset using lapply() and store it to concrete norm.
- **4.** Compare the strength values of concrete and concrete_norm to check if the data is normalised.
- 5. Split the dataset for training and testing in the ratio of 75:25.
- **6.** To use ANN, import the library 'neuralnet'.
- 7. Train the model using the neuralnet() with strength as the response variable and the rest as predictor variables from the training dataset, with only one hidden node.
- **8.** Visualise the neural network topology by plotting the model.
- **9.** To generate predictions on the test dataset, use the compute() which returns \$neurons and \$net.result which stores the predicted values.
- 10. Store the predicted values from model result\$net.result into predicted strength.
- 11. To analyse the relationship between predicted strength and the true value, display the correlation matrix using cor().
- **12.** To improve the model performance, develop a new neuralnet() model with 5 hidden nodes.
- 13. Plot the network again to see a drastic increase in no of connections.
- **14.** Evaluate the second model and display the correlation matrix.
- **15.** Stop.

CODE:

```
concrete<-read.csv(".\Concrete_Data.csv") View(concrete)
str(concrete)
normalize<-function(x){
    return ((x-min(x))/(max(x)-min(x)))
concrete_norm<-as.data.frame(lapply(concrete,normalize))
summary((concrete_norm$strength))
summary(concrete$strength) concrete_train<-
concrete_norm[1:773,] concrete_test<-concrete_norm[774:1030,]
library(neuralnet)</pre>
```

```
concrete_model<-neuralnet(strength~cement+slag+ash+water+superplasticizer

+coarseagg+fineagg+age, data=concrete_train)

plot(concrete_model)

model_results<-compute(concrete_model,concrete_test[1:8]) predicted_strength<-
model_results$net.result cor(predicted_strength,concrete_test$strength)

concrete_model2<-neuralnet(strength~cement+slag+ash+water+superplasticizer

+coarseagg+fineagg+age, data=concrete_train,hidden=5)

plot(concrete_model2)

model_results2 <- compute(concrete_model2, concrete_test[1:8])

predicted_strength2 <- model_results2$net.result

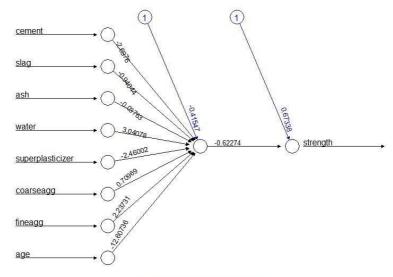
cor(predicted_strength2, concrete_test$strength) OUTPUT:
```

```
File Edit Selection View Go Run Terminal Help
                                                                                                                                             • concrete.r - Data Science using R - Visual Studio Code
            PROBLEMS 49 OUTPUT DEBUG CONSOLE TERMINAL
             > concrete<-read.csv("E:\\Academic Docs\\Semester-5\\Data Science using R\\Dat$
                str(concrete)
                               e': 1030 obs. of 9 variables:

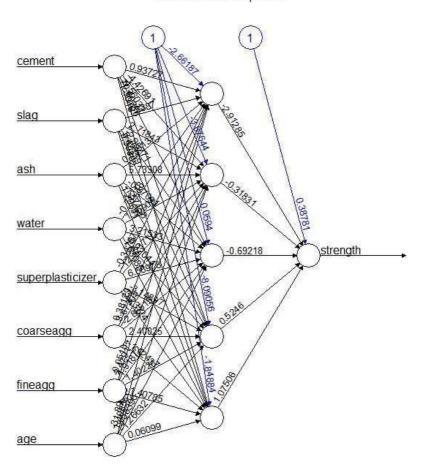
: num 540 540 332 332 199

: num 0 0 142 142 132 ...

: num 0 0 0 0 0 0 0 0 0
              'data.frame':
               $ cement
$ slag
               $ ash
              $ water : num 162 162 228 228 192 228 228 228 228 228 ...
$ superplasticizer: num 2.5 2.5 0 0 0 0 0 0 0 0 ...
$ coarseagg : num 1040 1055 932 932 978 ...
$ fineagg : num 676 676 594 594 826 ...
$ age : num 28 28 270 365 360 90 365 28 28 28 ...
$ strength : num 80 61.9 40.3 41 44.3 ...
             > normalize<-function(x){
             + return ((x-min(x))/(max(x)-min(x)))
+ }
              > concrete_norm<-as.data.frame(lapply(concrete,normalize))
            > library(neuralnet)
> concrete_model<-neuralnet(strength-cement+slag+ash+water+superplasticizer
                                                          +coarseagg+fineagg+age,
data=concrete_train)
             plot(concrete_model)
> model_results<-compute(concrete_model,concrete_test[1:8])
> predicted_strength
> cor(predicted_strength,concrete_test$strength)
               oncrete_model2<-neuralnet(strength~cement+slag+ash+water+superplasticizer
+coarseagg+fineagg+age,
data=concrete_train,hidden=5)
           > plot(concrete_model2)
> model_results2 <- compute(concrete_model2, concrete_test[1:8])
> predicted_strengtp1 <- model_results2$net.result
> cor(predicted_strengtp2, concrete_test$strength)
```



Error: 5.680718 Steps: 999



Error: 1.616487 Steps: 18311

RESULT:

Thus the R Program to implement modeling the strength of concrete using ANNs is executed successfully and the output is verified.

Ex	No:7
Da	te:

Identification of frequently Purchased groceries with Apriori

AIM:

To implement a R program to identify frequently purchased groceries with Apriori algorithm.

ALGORITHM:

- 1. Start
- 2. Import the 'arules' library to use Apriori algorithm and the Groceries dataset.
- **3.** Import the dataset 'Groceries' from arules library using data(). The preloaded dataset is a sparse matrix
- **4.** To do EDA, use the summary() function for the dataset.
- **5.** To see the contents of the sparse matrix, use inspect().
- **6.** To view the proportion of transactions of the items, use itemFrequency().
- 7. To view the proportion of transactions visually, we use itemFrequencyPlot() with 10 percent support.
- **8.** To limit the plot a specific number of items, use the topN parameter.
- **9.** To visualize the entire sparse matrix we use image() function for random 100 transactions.
- **10.** Train the apriori model using apriori(), with confidence threshold of 25%, set support level of 0.006 which is 60 transactions(2 per day) out of 9835, and to eliminate rules fewer than 2.
- 11. Summarise the groceryrules from the model.
- 12. Inspect the first 3 groceryrules and deduce the meaning of it.
- 13. To inspect which customer is most likely to buy a item relative to average customer we sort the top 5 groceryrules in the order of lift.
- **14.** To find the rules of only berries use the subset() in the transactions, items or rules and save it as a dataframe.
- **15.** Stop

CODE:

```
library(arules) data(Groceries)
summary(Groceries)
inspect(Groceries[1:5])
itemFrequency(Groceries[,1:3])
itemFrequencyPlot(Groceries,
support=0.1)
itemFrequencyPlot(Groceries, topN = 20)
image(Groceries[1:5])
image(sample(Groceries,100))
apriori(Groceries)
groceryrules <- apriori(Groceries, parameter=list(support=0.006, confidence = 0.25, minlen=2)) groceryrules
summary(groceryrules) inspect(groceryrules[1:3])
AIRSPEC(sort(groceryrules, by = "lift")[1:5]) berryrules <-
```

subset(groceryrules, items %in% "berries")
inspect(berryrules) write(groceryrules, file =
"groceryrules.csv") groceryrules_df <- as(groceryrules,
"data.frame") str(groceryrules_df)</pre>

OUTPUT:

```
PROBLEMS OUTPUT DEBUG CONSOLE TERMINAL JUPYTER
> library(arules)
> data(Groceries)
> summary(Groceries)
transactions as itemMatrix in sparse format with
9835 rows (elements/itemsets/transactions) and
169 columns (items) and a density of 0.02609146
most frequent items:
          whole milk other vegetables
                   2513
                                                 1903
                                                                              1809
                                  (Other)
                 yogurt
element (itemset/transaction) length distribution:
Sizes

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16

2159 1643 1299 1005 855 645 545 438 350 246 182 117 78 77 55 46

17 18 19 20 21 22 23 24 26 27 28 29 32

29 14 14 9 11 4 6 1 1 1 1 3 1
  Min. 1st Qu. Median Mean 3rd Qu. Max.
1.000 2.000 3.000 4.409 6.000 32.000
includes extended item information - examples:
            labels level2
Iabels level2 level1

frankfurter sausage meat and sausage
sausage sausage meat and sausage
liver loaf sausage meat and sausage
inspect(Groceries[1:5])
[1] {citrus fruit, semi-finished bread,
         margarine,
ready soups}
[2] {tropical fruit,
         coffee}
[3] {whole milk}
[4] {pip fruit,
yogurt,
cream cheese,
meat spreads}
[5] {other vegetables,
         whole milk,
        long life bakery product}
> itemFrequency(Groceries[,1:3])
frankfurter sausage liver loaf
0.058973055 0.093950178 0.005083884
> itemFrequencyPlot(Groceries, support=0.1)
> itemFrequencyPlot(Groceries, topN = 20)
> image(Groceries[1:5])
> image(sample(Groceries, 100))
> apriori(Groceries)
Apriori
Parameter specification:
ranameter specification;

confidence minval smax arem aval originalSupport maxtime support minlen
0.8 0.1 1 none FALSE TRUE 5 0.1 1

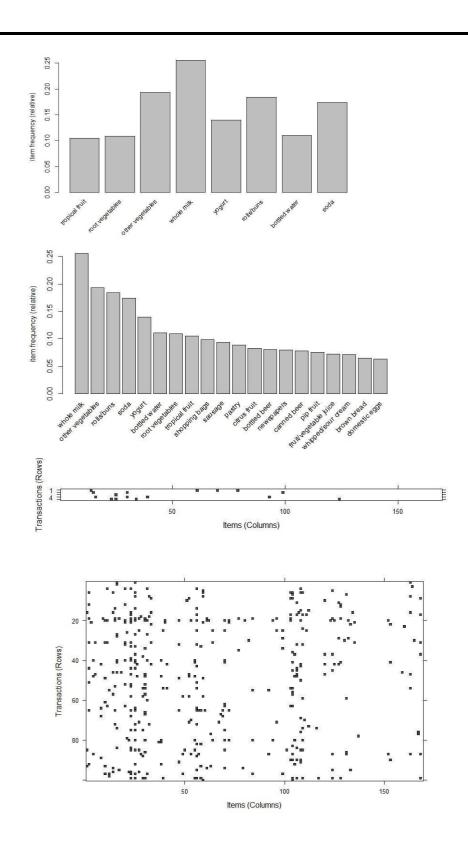
maxlen target ext
10 rules TRUE
Algorithmic control:
 filter tree heap memopt load sort verbose
0.1 TRUE TRUE FALSE TRUE 2 TRUE
Absolute minimum support count: 983
set item appearances ...[0 item(s)] done [0.00s].
set transactions ...[169 item(s), 9835 transaction(s)] done [0.01s].
sorting and recoding items ... [8 item(s)] done [0.00s].
creating transaction tree ... done [0.00s].
checking subsets of size 1 2 done [0.00s].
writing ... [0 rule(s)] done [0.00s].
```

```
creating transaction tree ... done 10.00sl.
 checking subsets of size 1 2 done [0.00s].
 writing ... [0 rule(s)] done [0.00s]. creating S4 object ... done [0.00s].
  set of 0 rules
 > groceryrules <- apriori(Groceries, parameter=list(support=0.006, confidence $
 Apriori
 Parameter specification:
  confidence minval smax arem aval originalSupport maxtime support minlen
   0.25 0.1
maxlen target ext
                                1 none FALSE
                                                                       TRUE
        10 rules TRUE
 Algorithmic control:
  filter tree heap memopt load sort verbose
0.1 TRUE TRUE FALSE TRUE 2 TRUE
 Absolute minimum support count: 59
set item appearances ...[0 item(s)] done [0.00s].
set transactions ...[169 item(s), 9835 transaction(s)] done [0.01s].
sorting and recoding items ... [109 item(s)] done [0.00s].
creating transaction tree ... done [0.00s].
checking subsets of size 1 2 3 4 done [0.01s].
writing ... [463 rule(s)] done [0.00s].
creating S4 object ... done [0.00s].
> groceryrules
 set of 463 rules
> summary(groceryrules)
set of 463 rules
rule length distribution (lhs + rhs):sizes
150 297 16
   Min. 1st Qu. Median Mean 3rd Qu. Max. 2.000 2.000 3.000 2.711 3.000 4.000
 summary of quality measures:
                            confidence coverage
Min. :0.2500 Min. :0.009964
1st Qu.:0.2971 1st Qu.:0.018709
     support
n. :0.006101
                                                                                       lift
 Min. :0.006101
1st Ou.:0.007117
                                                                                 Min. :0.9932
                                                                                 1st Ou.:1.6229
                            Median :0.3554
Mean :0.3786
3rd Qu.:0.4495
  Median :0.008744
                                                     Median :0.024809
                                                                                 Median :1.9332
           :0.011539
                                                    Mean :0.032608
3rd Qu.:0.035892
                                                                                 Mean :2.0351
  3rd Qu.:0.012303
                                                                                 3rd Qu.:2.3565
 Max. .. count
           :0.074835 Max.
                                     :0.6600 Max. :0.255516
 Min. : 60.0
1st Qu.: 70.0
  Median: 86.0
 Mean
           :113.5
  3rd Qu.:121.0
 Max.
           :736.0
mining info:
        data ntransactions support confidence
                           9835 0.006
  apriori(data = Groceries, parameter = list(support = 0.006, confidence = 0.25, minlen = 2))
> inspect(groceryrules[1:3])
                                                       support
                                                                       confidence coverage
      0.006914082 0.4000000 0.01728521
0.006100661 0.4054054 0.01504830
 [3] {herbs}
      lift count
1.565460 68
      1.586614 60
3.956477 69
 inspect(sort(groceryrules, by = "lift")[1:5])

lhs rhs
                                  rhs support confidence coverage lift

=> {root vegetables} 0.007015760 0.4312500 0.01626843 3.956477

=> {whipped/sour cream} 0.009049314 0.2721713 0.03324860 3.796886
      {herbs}
 [2] {berries}
[3] {tropical fruit,
        other vegetables,
whole milk} => {root vegetables} 0.007015760 0.4107143 0.01708185 3.768074
 [4] {beef, other vegetables} => {root vegetables} 0.007930859 0.4020619 0.01972547 3.688692
 [5] {tropical fruit,
other vegetables} => {pip fruit} 0.009456024 0.2634561 0.03589222 3.482649
> berryrules <- subset(groceryrules, items %in% "berries")
 > inspect(berryrules)
    lhs    rhs
 count
 [1] 89
[2] 104
[3] 101
[4] 116
        89
 [4] 116
> write(groceryrules, file = "groceryrules.csv")
> groceryrules_df <- as(groceryrules, "data.frame")
> str(groceryrules_df)
'data.frame': 463 obs. of 6 variables:
$ rules : chr "{pot plants} => {whole milk}" "{herbs} => {root vegetables}" "{herbs} => {other vegetables}" ...
$ support : num    0.00691    0.0061    0.00702    0.00773    0.00773    ...
$ confidence: num     0.4     0.405     0.431    0.475    0.475    ...
   $ confidence: num 0.4 0.405 0.431 0.475 0.475 ...
$ coverage : num 0.0173 0.015 0.0163 0.0163 0.0163 ...
                : num 1.57 1.59 3.96 2.45 1.86 ...
: int 68 60 69 76 76 69 70 67 63 88 ...
   $ lift
      count
```



RESULT:

Thus the R program to identify frequently purchased groceries using Apriori algorithm is executed successfully and the output is verified.

Ex. No: 8	
	Identification of frequently Purchased groceries with Apriori algorithm
Date:	ruentification of frequentry furchased groceries with Apriori algorithm

Aim:

To identify frequently purchased groceries using apriori.

Algorithm:

Step 1: Import the 'arules' package.

Step 2: Create a list of transaction datasets. Each transaction is represented as a list of items.

Step 3: Convert the transaction data into a binary format suitable for association rule mining.

Step 4: Use the Apriori algorithm to mine association rules with specified support and confidence thresholds.

Step 5: Display the frequent itemsets and association rules.

Coding:

```
library(arules)

transactions <- list(
    c("bread", "milk", "cereal"),
    c("bread", "milk"),
    c("bread", "cereal"),
    c("bread", "milk", "cereal"),
    c("eggs", "milk", "cereal")
)

transactions <- as(transactions, "transactions")

rules <- apriori(transactions, parameter = list(support = 0.3, confidence = 0.7))

inspect(rules)
```

Output:

```
Apriori
Parameter specification:
confidence minval smax arem aval originalSupport maxtime support minlen m
       0.7 0.1 1 none FALSE TRUE 5 0.3
Algorithmic control:
 filter tree heap memopt load sort verbose
Absolute minimum support count: 1
set item appearances ...[0 item(s)] done [0s].
set transactions ...[5 item(s), 5 transaction(s)] done [0s].
sorting and recoding items ... [4 item(s)] done [0s].
creating transaction tree \dots done [0s].
checking subsets of size 1 2 done [0s].
writing ... [3 rule(s)] done [0s].
creating S4 object ... done [0s].
               rhs
                        support confidence lift count
[1] {bread} => {cereal} 0.4 1 2.5 2
[2] {cereal} => {bread} 0.4 1 2.5 2
[3] {milk} => {bread} 0.6 1 2.5 3
```

Result:

Thus, the identification of frequently purchased groceries is executed successfully.

Ex. No: 9	
Date:	Finding Teen Segments of Market

Aim:

To implement Teen segments of market in R.

Algorithm:

Step 1: Load the 'ggplot2' library to enable data visualization.

Step 2: Set a random seed for reproducibility.

Step 3: Generate sample data for demonstration purposes. In this case, create a data frame called 'data' with 100 random values for "Age" and "Spending."

Step 4: Specify the number of clusters (k) for K-Means clustering. In this example, k is set to 2.

Step 5: Perform K-Means clustering on the 'data' using the specified number of clusters (k).

Step 6: Add a new column, "Cluster," to the data frame, representing the cluster assignments for each data point.

Step 7: Display the data in a tabular format. Print the column headers for "Age," "Spending," and "Cluster."

Step 8: Iterate through each row of the data frame and print the "Age," "Spending," and "Cluster" values in a tabular format.

Coding:

```
\label{eq:set_sed} \begin{subarray}{l} library(ggplot2) \\[-2pt] set.seed(123) \\[-2pt] data <- data.frame( \\[-2pt] Age = runif(100, min = 13, max = 19), \\[-2pt] Spending = rnorm(100, mean = 50, sd = 10) \\[-2pt] ) \\[-2pt] k <- 2 \\[-2pt] kmeans\_result <- kmeans(data, centers = k) \\[-2pt] data Cluster <- as.factor(kmeans\_result Cluster) \\[-2pt] \end{subarray}
```

Output:

Result:

Thus, the R program for teen segment of market is executed successfully.

Ex. No: 10	
Date :	TUNING STOCK MODELS

Aim:

To implement tuning stock models for better performance.

Algorithm:

- Step 1: Load the 'quantmod' and 'caret' libraries, which are necessary for stock data retrieval and model tuning.
- Step 2: Download historical stock data using the `getSymbols` function, specifying the desired stock symbol (e.g., "AAPL").
- Step 3: Create a data frame named 'stock_data' to store the historical stock data, including dates and adjusted closing prices.
- Step 4: Define the training and test data sets by selecting a specific range of rows from the 'stock data' data frame.
- Step 5: Create time series objects ('train_ts' and 'test_ts') using the 'xts' function to work with the adjusted closing prices of the training and test data.
- Step 6: Define a control object ('ctrl') for model tuning, specifying parameters like the resampling method (cross-validation with 5 folds).
- Step 7: Tune a stock prediction model (in this example, a simple linear regression) using the 'train' function. The model is trained on the 'train_ts' data, with the dependent variable as 'train_ts' and the independent variable as the index of 'train_ts.' The model tuning is performed under the specified control settings.
- Step 8: Print the results of the tuned model, including information about the model's performance, such as root mean squared error (RMSE) and R-squared values, as well as any tuned parameters.

Coding:

```
library(quantmod)
library(caret)
getSymbols("AAPL")
stock_data <- data.frame(Date = index(AAPL), Adjusted = Ad(AAPL))</pre>
```

```
train_data <- stock_data[1:200, ]

test_data <- stock_data[201:nrow(stock_data), ]

train_ts <- xts(train_data$Adjusted, order.by = train_data$Date)

test_ts <- xts(test_data$Adjusted, order.by = test_data$Date)

ctrl <- trainControl(method = "cv", number = 5)

tune_results <- train(train_ts ~ index(train_ts), data = data.frame(train_ts), method = "lm", trControl = ctrl)

print(tune_results)</pre>
```

Output:

```
Linear Regression

200 samples
    1 predictor

No pre-processing
Resampling: Cross-Validated (5 fold)
Summary of sample sizes: 160, 160, 160, 160
Resampling results:

RMSE Rsquared
    10.98207 0.8478949

Tuning parameter 'intercept' was held constant at a value of TRUE
```

Result:

Thus, the implementation of tuning the stocks for better performance is executed successfully.