# AI19542 - DATA SCIENCE USING R - LAB MANUAL



# DEPARTMENT OF ARTIFICIAL INTELLIGENCE AND MACHINE LEARNING

# AI19542 – DATA SCIENCE USING R LAB MANUAL

THIRD YEAR

FIFTH SEMESTER

2024 - 2025

ODD SEMESTER

Ex No:1	Basics of R – data types, vectors, factors, list and data
Date:01/08/24	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("JohnDoe","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>
gender
```

```
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:** 

AI19542 221501011

```
X File Edit Selection View Go Run Terminal Help
              PROBLEMS (73)
                                      OUTPUT DEBUG CONSOLE TERMINAL
             [1] "logical"
[1] "numeric"
[1] "integer"
[1] "character"
[1] "character"
[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"
 R
              $temperature
[1] 98.1
              $flu_status
[1] FALSE
               [1] MALE
Levels:
                           FEMALE MALE
               [1] O
Levels: A B AB O
               [1] SEVERE
Levels: MILD < MODERATE < SEVERE
               $temperature
[1] 98.1
               [1] 98.1
[1] 98.1
$temperature
[1] 98.1
              $flu_status
[1] FALSE
              101.4
                                                TRUE
```

```
File Edit Selection View Go Run Terminal Help
                                                                                                                                                  BasicsO
         PROBLEMS (T) OUTPUT DEBUG CONSOLE TERMINAL JUPYTER
         2 96.
                             FALSE
                                 TRUE
           temperature gender
        1 98.1 MALE
2 98.6 FEMALE
        [1] "John Doe"
                             "Jane Doe"
                                               "Steven Grant"
           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] [,2]
[1,] 1 3
[2,] 2 4
               [,1][,2]
         [,1] [,2]
[1,] 1 4
[2,] 2 5
[3,] 3 6
[1] 1 4
R
         [1] 1 4
[1] "apple"
[1] "cherry"
          [1] "banana"
          [1] "orange"
```

#### **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: 08/08/24	

#### 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 directoryand 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. Displaythe 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))</pre>
x_{train} \leftarrow bcd_n[1:469,] x_{test} \leftarrow 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:**

```
DEBUG CONSOLE
'data.frame':
                 569 obs. of 32 variables:
                     : int 87139402 8910251 905520 868871 9012568 906539 925291 87880 862989 89827 ...

: chr "B" "B" "B" ...

: num 12.3 10.6 11 11.3 15.2 ...
$ id
$ diagnosis
 $ radius mean
 $ texture_mean : num 12.4 18.9 16.8 13.4 13.2 ... $ perimeter mean : num 78.8 69.3 70.9 73 97.7 ...
$ points mean
                        num 0.037 0.0264 0.0248 0.048 0.0266 ...
 $ symmetry_mean
$ dimension_mean
                      $ radius_se
$ texture_se
                      $ perimeter_se
                      : num 1.67 3.43 1.34 1.85 1.34 ...
: num 17.4 27.1 13.5 26.3 17.7 ...
 $ area se
 $ smoothness se
                      $ compactness_se
                      $ 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 ...

num 13.5 11.9 12.4 11.9 16.2 ...
| Standing worst | num | 13.5 11.9 12.4 11.9 16.2 |
| Stexture_worst | num | 15.6 12.9 26.4 15.8 15.7 |
| Sperimeter_worst | num | 87 78.3 79.9 76.5 104.5 |
| Sarea_worst | num | 64.25 471 434 819 |
| Smoothness_worst | num | 0.139 0.121 0.137 0.137 0.113 | ...
$ dimension_worst : num 0.0677 0.0759 0.0788 0.0678 0.0677 ...
             Benign Malignant
  Benign
Malignant
                   4
                            35
[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: 22/08/24	

#### 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))
print(sms_corpus)
print(as.character(sms_corpus[[6]]))
sms_dtm <- DocumentTermMatrix(sms_corpus, control = list
(tolower=TRUE,removeNumbers=TRUE,stopwords=TRUE,removePunctuations=TRUE,stemm
i ng=TRUE))</pre>
```

```
x_train <- sms_dtm[1:4169, ]</pre>
x_test <- sms_dtm[4170:5572, ]</pre>
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] x_test</pre>
<- x_test[ , sms_freq_word_test]</pre>
convert_counts <- function(x) \{x <- ifelse(x > 0, "Yes",
"No")} x train <- apply(x train, MARGIN = 2,convert counts)
x test <- apply(x test, MARGIN = 2,convert counts)</pre>
library(e1071)
model <- naiveBayes(x_train, y_train,laplace=1)</pre>
                       predict(model, x test)
y_pred <-
table(y_pred, y_test) print(cm)
                            sum(diag(cm))/sum(cm)
print(paste("Accuracy:
",acc*100,"%")) OUTPUT:
PROBLEMS (73) OUTPUT DEBUG CONSOLE TERMINAL JUPYTER
                                                                                                                          R Interactive + ✓ □ 🛍 ^ X
'data.frame': 5572 obs. of 5 variables: $ v1 : chr "ham" "ham" "spam" "ham" ...
 $ v2 : chr "Go until jurong point, crazy.. Available only in bugis n great world la e buffet... Cine there got amore wat..." "Ok lar... Joking wif u oni..." "Free entry in 2 a wkly comp to win FA Cup final tkts 21st May 2005. Text FA to 87121 to receive entry question(std txt rate)T&C "U dun say so early hor... U c already then say..." ...
$ X : chr "" "" "" "" "" ""
 $ X : chr "" "" "" "" ...
$ X.1: chr "" "" "" ...
 $ X.2: chr "" "" "" ...
 <<VCorpus>>
Metadata: corpus specific: 0, document level (indexed): 0
[1] "FreeMsg Hey there darling it s been 3 week s now and no word back I d like some fun you up for it still Tb ok XXX std chgs to send 1 50 to rcv"
y_pred ham spam
  ham 1205 10
  spam 16 172
[1] "Accuracy: 98.1468282252316 %" > |
```

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

ExNo:4
Date:29/08/24

# **Risky Bank Loans using Decision Trees**

# 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. Displaythe 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 accuracyof 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)</pre>
                  train_sample
set.seed(123)
sample(1000, 800) str(train_sample)
x_train <- credit[train_sample, -17]
x_test <- credit[-train_sample, -17]
y_train <- credit[train_sample, 17]</pre>
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,"%
"))
```

# **OUTPUT:**

```
Evaluation on training data (900 cases):

Decision Tree
Size Errors

69 99(11.6%) <<

(a) (b) <-classified as

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

Attribute usage:

100.08% checking balance
54.22% credit history
48.22% months loan duration
42.22% savings balance
31.88% purpose
22.33% employment duration
9.22% years at residence
8.78% housing
8.44% jobs
6.11% other credit
```

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

$\mathbf{FvNo}$	5
L'AINU.	_

Date:05/09/24

# Medical Expense withLinear Regression.

#### 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 amore 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$region)
cor(insurance[c("age","bmi","children","expenses")])
pairs(insurance[c("age","bmi","children","expenses")])
library(psych)
pairs.panels(insurance[c("age","bmi","children","expenses")])8
ins_model <- lm(expenses ~ age + children + bmi + sex + smoker + region, data = insurance) ins_model
summary(ins_model)
insurance$age2 <- insurance$age^2</pre>
```

```
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:**

```
plot.png - Visual Studio Code
> insurance<-read.csv("E:\\Academic Docs\\Semester-5\\Data Science using R\\in$ > str(insurance)
  'data.frame': 138 obs. of 7 variables:
  's age : int 19 18 28 33 32 31 46 37 37 60 ...
  's sex : Factor w/ 2 levels "female", "male": 12 2 2 2 1 1 1 2 1 ...
  's hmi: rum 27,9 33.8 33 22,7 28,9 25,7 33.4 27,7 29.8 25.8 ...
  's children: int 0 1 3 0 0 0 1 3 2 0 ...
  's moker : Factor w/ 2 levels "no", "yes": 2 1 11 1 1 1 1 1 1 1 ...
  's region : Factor w/ 4 levels "no", "yes": 2 1 1 1 1 1 1 1 1 1 1 ...
  's region : Factor w/ 4 levels "no" theast", "northwest", ...: 4 3 3 2 2 3 3 2 1 2 ...
  's expenses: rum 1688 1726 4449 21984 3867 ...
  's summary(insuranceSexpenses)
  Hin. 1st Qu. Median Mean and Qu. Max.
  1122 4740 9382 13270 16640 63770
  >hist(insuranceSexpenses)
  *hist(insuranceSexpenses)
  *hist(insuranceSexpenses)
> table(insuranceSregion)

northeast northwest southwest

324 325 364 325

> cor(insurance[c(ap*, bmi*, children*, expenses*)])

age bmi children expenses

age bmi children expenses

bmi children expenses

bmi children (1.995410 1.00000000 0.0254070 1.01957020

children 0.0554010 1.000000000 0.01254472 1.00000000 0.01597020

children 0.055400000 0.01254472 1.00000000 0.05790200

pairs(insurance[c(ap*, bmi*, children*, expenses*)])

> library(psych)

> pairs, insurance[c(ap*, "bmi*, children*, "expenses*)])

Error: unexpected numeric constant in "pairs, panels(insurance[c("ap*, "bmi*, "children*, "expenses*)])

= ins model < lm(expenses ~ age + children + bmi + sex + smoker + region, d$

> ins_model
 call:
lm(formula = expenses ~ age + children + bmi + sex + smoker +
    region, data = insurance)
 Coefficients:
(Intercept)
-11941.6
sexmale
-131.4
regionsouthwest
-959.3
                                                                         age children bmi
256.8 475.7 339.3
smokeryes regionnorthwest
23847.5 -352.8 regionsouthes
 > summary(ins_model)
 call:
lm(formula = expenses ~ age + children + bmi + sex + smoker +
    region, data = insurance)
  Residuals:
Min 1Q Median 3Q Max
-11302.7 -2850.9 -979.6 1383.9 29981.7
Coefficients:

Estimate Sd. Error t value Pr(>|t|)

(Intercept) 11941.6 087.8 -12.089 < 2e-10 ***
age 256.8 11.9 21.586 < 2e-10 ***
bii 39.3 28.6 11.864 < 2e-10 ***
seowale -131.3 33.2, 0 -0.395 0.603275
seokeryes 238.47.5 413.1 57.723 < 4e-10 ***
regiomorthwest -352.8 476.3 -0.741 (0.458976
regiomostotheast -1815.6 476.3 -0.741 (0.458976
regiomostotheast -1815.6 476.7 -2.163 0.00863 **
regiomostotheast -1815.6 477.9 -2.163 0.00863 **
  Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
  Residual standard error: 6062 on 1329 degrees of freedom
Multiple R-squared: 0.7509, Adjusted R-squared: 0.7494
F-statistic: 500.9 on 8 and 1329 DF, p-value: < 2.2e-16
 > insuranceSage2 <- insuranceSage^2

> insuranceSami30 <- ifelse(insuranceSbmi >= 30,1,0)

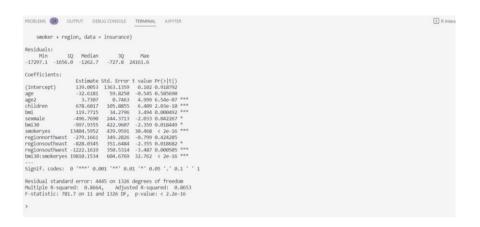
> expenses - bmi30*smoker

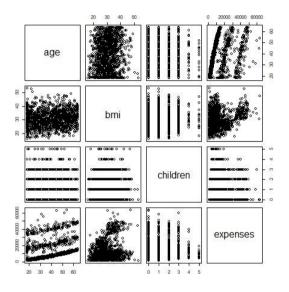
> expenses - bmi30*smokeryes+bmi30:smokeryes

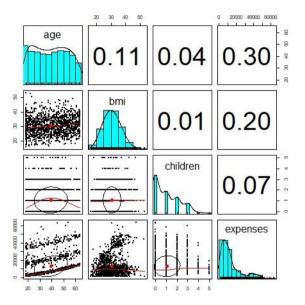
expenses - bmi30*smokeryes+bmi30:smokeryes

> ins model2 <- lm(expenses ~ age+age2+children+bmi+sex+bmi30*smoker+region, d$

> summary(ins model2)
 call:
lm(formula = expenses ~ age + age2 + children + bmi + sex + bmi30 *
smoker + region, data = insurance)
Min 1Q Median 3Q Max
-17297.1 -1656.0 -1262.7 -727.8 24161.6
```







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

ExNo: 6	
	Modeling strength of concrete.
Date:12/10/24	

# AIM:

To build a predictive model for the compressive strength of concrete based on its composition and age using linear regression in R.

#### **ALGORITHM:**

- 1. Start
- 2. Load the Insurance dataset and check its structure.
- 3. Get summary statistics and check skewness using mean, median, and histogram.
- 4. Check the distribution of "region" using a table.
- 5. Create a correlation matrix for "age," "bmi," "children," and "expenses."
- 6. Use scatterplots to examine relationships among "age," "bmi," "children," and "expenses."
- 7. Fit an initial linear model with "expenses" as the target, then improve by adding 'age2' (age squared) and 'bmi30' (1 if bmi >= 30) and re-evaluate.
- 8. Stop

# **PROGRAM:**

```
library(caret) library(ggplot2) data <- read.csv("concrete.csv")

head(data) sum(is.na(data)) set.seed(123) trainIndex <-
createDataPartition(data$CompressiveStrength, p =

0.8, list = FALSE) trainData

<- data[trainIndex,] testData

<- data[-trainIndex,] model

<- lm(CompressiveStrength

~., data = trainData)

summary(model)

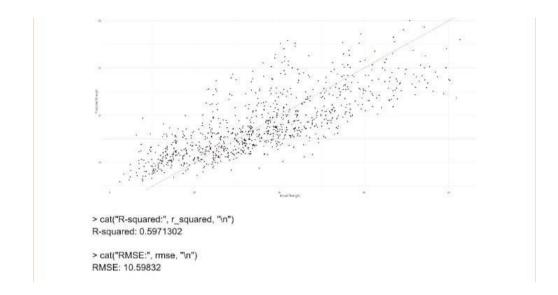
predictions <- predict(model, newdata = testData)
```

```
mae <- mean(abs(predictions - testData$CompressiveStrength))
print(paste("Mean Absolute Error:", round(mae, 2)))

ggplot() +
  geom_point(aes(x = testData$CompressiveStrength, y = predictions), color = 'blue') +
  geom_abline(slope = 1, intercept = 0, linetype = "dashed", color = "red") + labs(title
  = "Predicted vs Actual Compressive Strength", x = "Actual Strength", y =
  "Predicted Strength") + theme_minimal()</pre>
```

# **OUTPUT:**

```
> str(concrete)
 'data.frame': 1030 obs. of 10 variables:
 $ cement
               : num 540 540 332 332 199 ...
 $ slag
              : num 0 0 142 142 132 ..
          : num 0 0 142 142 132 ...
: num 0 0 0 0 0 0 0 0 0 0 ...
 $ ash
               : num 162 162 228 228 192 228 228 228 228 228 ...
 $ water
 $ superplastic : num 2.5 2.5 0 0 0 0 0 0 0 0
 : num 1040 1055 932 932 978 ...
             : int 28 28 270 365 360 90 365 28 28 28 ...
 $ age
 $ strength
                : num 80 61.9 40.3 41 44.3
 $ Predicted Strength: num 55.1 54.7 57.6 68 59.4 ...
 > summary(model)
 Call:
 Im(formula = strength ~ cement + slag + water + superplastic +
   coarseagg + fineagg + age, data = concrete)
 Residuals:
         1Q Median 3Q Max
 -30.901 -7.239 0.441 6.899 34.408
 Coefficients:
         Estimate Std. Error t value Pr(>|t|)
 (Intercept) 121.611036 17.015934 7.147 1.69e-12 ***
             0.067636 0.004135 16.357 < 2e-16 ***
 cement
          0.042550 0.005192 8.196 7.39e-16 ***
          -0.323265 0.032336 -9.997 < 2e-16 ***
 superplastic 0.371641 0.094876 3.917 9.56e-05 ***
 coarseagg -0.027502 0.006913 -3.978 7.44e-05 ***
          -0.038549 0.006777 -5.688 1.68e-08 ***
 fineagg
          0.109746 0.005514 19.903 < 2e-16 ***
age
Signif. codes: 0 **** 0.001 *** 0.01 ** 0.05 '.' 0.1 ' ' 1
Residual standard error: 10.64 on 1022 degrees of freedom
Multiple R-squared: 0.5971, Adjusted R-squared: 0.5944
F-statistic: 216.4 on 7 and 1022 DF, p-value: < 2.2e-16
> ggplot(concrete, aes(x = strength, y = Predicted_Strength)) +
+ geom_point() +
+ geom_abline(slope = 1, intercept = 0, color = "red") +
+ labs(title = "Actual vs Predicted Concrete Strength",
     x = "Actual Strength",
      y = "Predicted Strength") +
+ theme_minimal()
```



Thus the R Script program to implement Modeling strength of concrete is executed successfullyand the output is verified.

ExNo: 7

Date:19/10/24

# Identification of frequently Purchased groceries with Apriori algorithm.

# AIM:

To identify frequent itemsets of grocery items that are commonly purchased together using the Apriori algorithm. This will help in understanding customer buying patterns and optimizing store layout or inventory.

# **ALGORITHM:**

- 1. Start
- 2. Load Data: Load the transaction dataset (assume each transaction is a list of items purchased).
- 3. Data Preprocessing: Convert the data into a transactional format suitable for association rule mining.
- 4. Set Parameters: Define minimum support and confidence levels for the Apriori algorithm.
- 5. Apply Apriori Algorithm: Use the Apriori algorithm to find frequent itemsets.
- 6. Generate Association Rules: Extract association rules from the frequent itemsets based on support and confidence thresholds.
- 7. Analyze Results: Sort and filter rules to identify the most frequently purchased item combinations. 8. Stop

# **PROGRAM:**

```
if(!require(arules)) install.packages("arules", dependencies=TRUE)
library(arules) data("Groceries") summary(Groceries)
min_support <- 0.01 # Example: at least 1% of transactions
min_confidence <- 0.5 # Example: at least 50% confidence
frequent_itemsets <- apriori(Groceries, parameter = list(supp = min_support, conf =
min_confidence)) summary(frequent_itemsets) inspect(frequent_itemsets[1:10])
rules <- apriori(Groceries, parameter = list(supp = min_support, conf = min_confidence,
target = "rules")) summary(rules)
inspect(sort(rules, by = "confidence")[1:10]) # Display top 10 rules by confidence
if(!require(arulesViz)) install.packages("arulesViz", dependencies=TRUE)
library(arulesViz)
plot(rules, method = "graph", control = list(type = "items"))
OUTPUT:</pre>
```

<u>Summary of the Groceries Dataset</u> 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 rolls/buns soda yogurt (Other) 2513 1903 1809 1715 1372 34055

# **Frequent Itemsets:**

set of 50 itemsets

example of first 10 itemsets (sorted by support):

items support

[1] {whole milk} 0.25551601
[2] {other vegetables} 0.19349263
[3] {rolls/buns} 0.18393493
[4] {soda} 0.17437722
[5] {yogurt} 0.13950178
[6] {whole milk, other vegetables} 0.0751

[7] {whole milk, yogurt} 0.0561

# **Association Rules (Top 10 by Confidence)**:

set of 10 rules

example of first 10 rules (sorted by confidence):

lhs rhs support confidence lift

[1] {yogurt}  $\Rightarrow$  {whole milk} 0.0561 0.4032 1.57

[2]  $\{\text{rolls/buns}\} => \{\text{whole milk}\} \quad 0.0567 \quad 0.3084 \quad 1.21$ 

[3]  $\{soda\} => \{whole milk\} = 0.0569 = 0.3058 = 1.20$ 

[4]  $\{\text{tropical fruit}\} => \{\text{whole milk}\}\ 0.0519\ 0.2674\ 1.03$ 

[5] {other vegetables}  $\Rightarrow$  {whole milk} 0.0751 0.3926 1.53

# **RESULT:**

Thus the R program to Identification of frequently Purchased groceries with Apriori algorithm is executed successfully and the output is verified.

ExNo	): 8
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Date:26/10/24

# **Finding Teen Segments of Market.**

# AIM:

The aim of this process is to identify and segment the teen demographic in a market based on behavior, preferences, or other relevant characteristics for targeted marketing or product development.

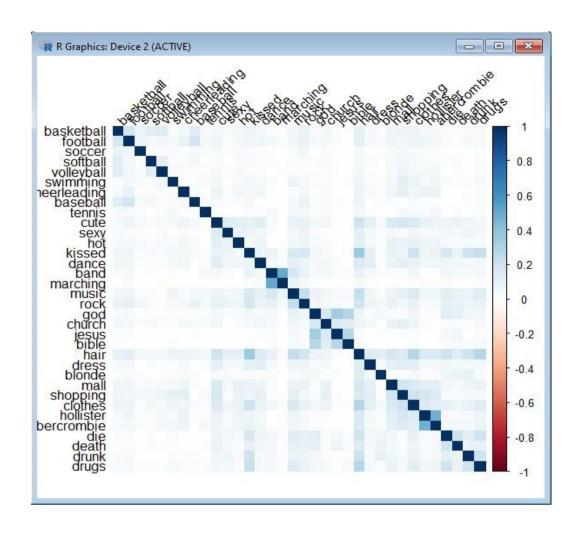
#### **ALGORITHM:**

- 1. START: Collect raw data from sources relevant to the teen market (e.g., social media data, survey responses).
- 2. PREPROCESSING: Clean the data (e.g., remove missing values, correct errors).
- 3. SELECT FEATURES: Choose features that help in segmentation (e.g., age, purchase patterns, interests).
- 4. APPLY CLUSTERING ALGORITHM: Run clustering algorithms (e.g., K-Means or DBSCAN) to create market segments.
- 5. EVALUATE MODEL: Evaluate the clustering performance using a scoring metric (e.g., silhouette score).
- 6. VISUALIZE DATA: Visualize the segmented data to understand different groups.
- 7. EXTRACT INSIGHTS: Identify unique patterns and preferences within each segment.
- 8. STOP: Develop targeted marketing strategies based on the insights from the segmentation.
- 9. This approach allows businesses to better understand the teen market and tailor their products or marketing campaigns accordingly.

# **PROGRAM:**

```
library(dplyr)
library(ggplot2)
library(corrplot)
load_data <- function(file_path) {
    df <- read.csv(file_path)
    return(df)
    }
    preprocess_data <- function(df) {
        # Check for missing values
        print(colSums(is.na(df)))
        df[is.na(df)] <- 0 # Fill missing values with 0
        return(df)
        }
    analyze_segments <- function(df) {
    # Example: Segment by gender
        Al19542
```

```
gender_counts <- table(df$gender)</pre>
print("Gender
                       Distribution:")
print(gender_counts)
 interest_features <- c('basketball', 'football', 'soccer', 'softball', 'volleyball',
                 'swimming', 'cheerleading', 'baseball', 'tennis',
                 'cute', 'sexy', 'hot', 'kissed', 'dance',
                 'band', 'marching', 'music', 'rock', 'god',
                 'church', 'jesus', 'bible', 'hair', 'dress',
                 'blonde', 'mall', 'shopping', 'clothes',
                 'hollister', 'abercrombie', 'die', 'death',
                 'drunk', 'drugs')
 corr_matrix <- cor(df[interest_features]) corrplot(corr_matrix,</pre>
method = "color", tl.col = "black", tl.srt = 45)
}
main <- function(file_path) {</pre>
df <- load_data(file_path)</pre>
df <- preprocess_data(df)</pre>
analyze_segments(df)
main('path_to_your_file.csv')
OUTPUT:
```



Thus the R program to Finding Teen Segments of Market is executed successfully and the output is verified.

ExNo: 9

Date: 26/10/24

# Tuning stock models for better performance.

# AIM:

The aim is to enhance the predictive performance of stock market models by optimizing hyperparameters, improving data features, and using techniques like crossvalidation and model selection to better forecast stock prices or trends.

# **ALGORITHM:**

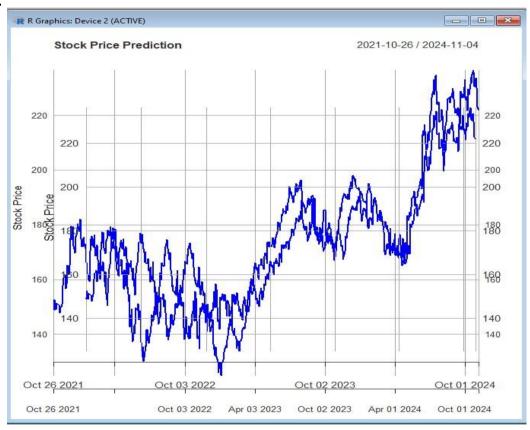
- 1. Start
- 2. Data Collection: Gather historical stock data (e.g., price, volume, market sentiment, technical indicators).
- 3. Data Preprocessing: Clean the data by handling missing values, normalizing features, and creating relevant indicators (e.g., moving averages, RSI).
- 4. Feature Engineering: Create new features based on existing data to improve model predictions (e.g., lagged values, percentage changes, or volatility).
- 5. Model Selection: Choose an appropriate model (e.g., Linear Regression, Decision Trees, Random Forest, LSTM for time series).
- 6. Hyperparameter Tuning: Tune the hyperparameters of the model using techniques like Grid Search or Random Search to optimize performance.
- 7. Cross-Validation: Implement cross-validation (e.g., k-fold) to ensure that the model generalizes well on unseen data.
- 8. Model Evaluation: Evaluate the model's performance using metrics like RMSE, MAE, or accuracy, and compare the results with different models.
- 9. Model Refinement: Refine the model by adjusting hyperparameters further, adding/removing features, or testing different algorithms to achieve better results
- 10. End.

# **PROGRAM:**

```
library(Metrics)
data <-
read.csv("C:/Users/AI_LAB/Desktop/77/stock.csv") if
(is.null(data)) { stop("Data not loaded. Please check the file
path.")
}
str(data)
data$Closing.Volume <- as.numeric(as.character(data$Closing.Volume)) # Update based on
your target variable data <- na.omit(data)
set.seed(123)
```

train\_index <- sample(1:nrow(data), 0.8 \* nrow(data)) train\_data <- data[train\_index, ] test\_data <- data[-train\_index, ] rf\_model <- randomForest(Closing.Volume ~ ., data = train\_data, ntree = 100) predictions <- predict(rf\_model, newdata = test\_data) actuals <- test\_data\$Closing.Volume mae <- mean(abs(predictions - actuals)) rmse <- sqrt(mean((predictions - actuals)^2)) cat("Mean Absolute Error:", mae, "\n") cat("Root Mean Squared Error:", rmse, "\n") plot(test\_data\$Date, actuals, type = 'l', col = 'blue', ylim = range(c(actuals, predictions)), xlab = 'Date', ylab = 'Closing Price', main = 'Actual vs Predicted Closing Prices') lines(test\_data\$Date, predictions, col = 'red') legend("topright", legend = c("Actual", "Predicted"), col = c("blue", "red"), lty = 1)

# **OUTPUT:**



# **RESULT:**

Thus the R program to Tuning stock models for better performance is executed successfully and the output is verified.