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

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])
 }
```

```
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

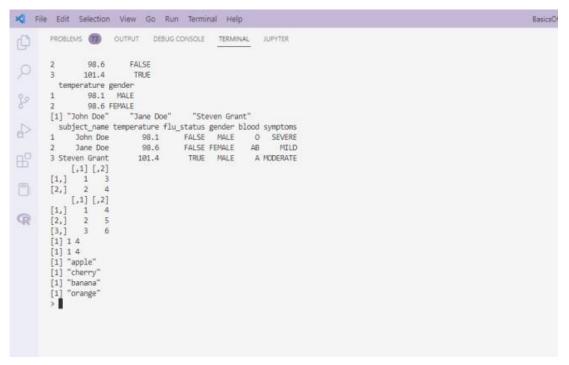
1 John Doe 98.1 FALSE MALE O SEVERE

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))</pre>
```

```
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:	

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,"%"))
```

### **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:	

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>
```

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

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
31.93% purpose
22.33% cmptoyment duration
9.22% years, at residence
8.20% housing
8.40% bits of the credit
9.13% other_credit
```

```
5.78% amount
4.98% existing_loans_count
4.287 phone
2.888 percent of income
1.888 dependents
0.78% age

Time: 0.0 secs
> y_pred <- predict(model,x_test)
> cm <- table(y_pred,y_test)
> print(cm)
> print(sm)
y_cest
y_pred no se
y_sest
y_pred no ses
y_sest
y_pred no ses
y_sest
```

## **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:	•

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)

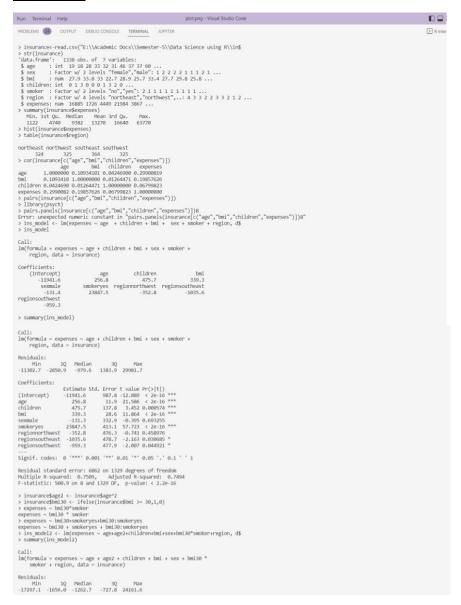
expenses ~ bmi30\*smoker

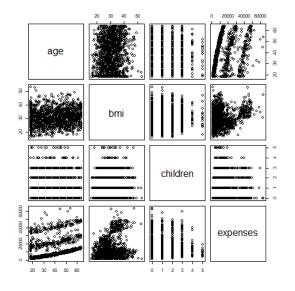
expenses ~ bmi30+smokeryes+bmi30:smokeryes

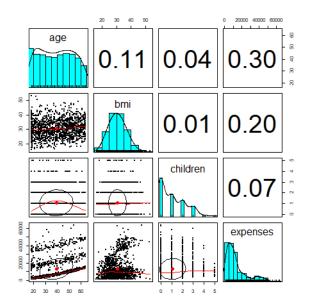
ins\_model2 <- lm(expenses ~ age+age2+children+bmi+sex+bmi30\*smoker+region, data=insurance)

summary(ins\_model2)

#### **OUTPUT:**







# **RESULT:**

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