

EXPERIMENT -2

AIM:- Create two vectors in R for Numeric Data.

PROCEDURE:-

```
#create two vectors for numeric data  x
<- c(1,2,3,4,5)

# use c() to combine elements into a numeric vector of
length 5 y <- c(6,7,8,9,10) x y class(x) class(y)

#extract the second element of x  x[2]
#update the second element of y to 100 y[2]
<-100
y
#operations between two numeric vectors
print(paste("addition", x+y)) print(paste("division",y/5))
print(paste("subtraction",y-x))

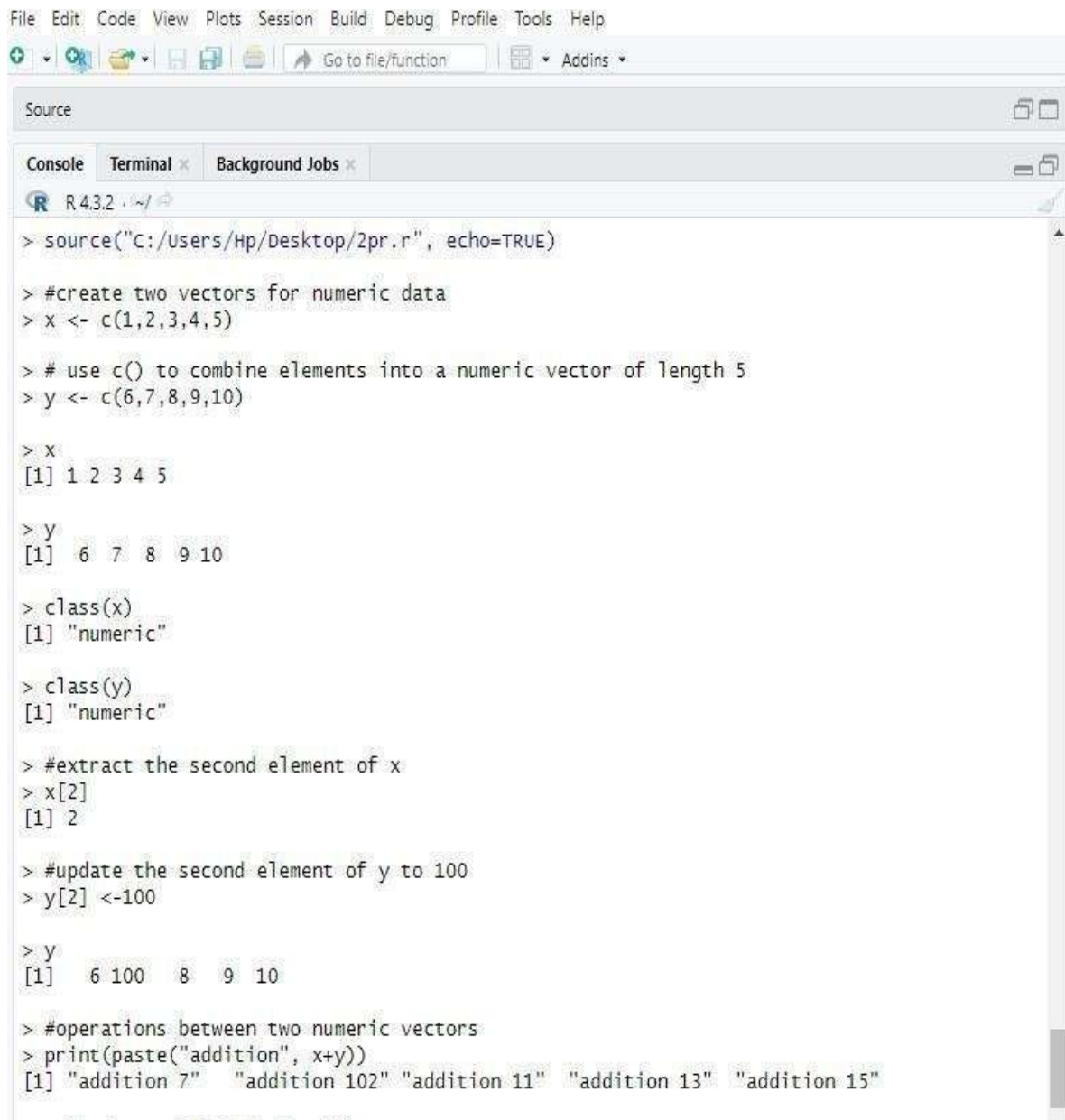
#create two vectors for numeric data  x <- c(1,2,3,4,5)

# use c() to combine elements into a numeric vector of
length 5 y <- c(6,7,8,9,10) x y class(x) class(y)

#extract the second element of x  x[2]

#update the second element of y to 100 y[2] <-
100  y

#operations between two numeric vectors
print(paste("addition", x+y))
```

INPUT/OUTPUT:-The image shows a screenshot of the R Studio interface. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. Below the menu is a toolbar with icons for file operations and a search bar labeled 'Go to file/function'. The main window is divided into three panes: 'Source' (empty), 'Console' (active), and 'Background Jobs' (empty). The Console pane shows the R prompt 'R 4.3.2 ~/' followed by a series of commands and their outputs. The commands create two numeric vectors, x and y, using the c() function. Vector x contains the values 1, 2, 3, 4, 5 and vector y contains 6, 7, 8, 9, 10. The class of both vectors is confirmed as 'numeric'. The second element of x is extracted, showing the value 2. The second element of y is updated to 100. Finally, the two vectors are added together using the + operator, and the result is printed as a character string, showing the sum of each corresponding element: 'addition 7', 'addition 102', 'addition 11', 'addition 13', and 'addition 15'.

```
> source("C:/Users/Hp/Desktop/2pr.r", echo=TRUE)

> #create two vectors for numeric data
> x <- c(1,2,3,4,5)

> # use c() to combine elements into a numeric vector of length 5
> y <- c(6,7,8,9,10)

> x
[1] 1 2 3 4 5

> y
[1] 6 7 8 9 10

> class(x)
[1] "numeric"

> class(y)
[1] "numeric"

> #extract the second element of x
> x[2]
[1] 2

> #update the second element of y to 100
> y[2] <-100

> y
[1] 6 100 8 9 10

> #operations between two numeric vectors
> print(paste("addition", x+y))
[1] "addition 7" "addition 102" "addition 11" "addition 13" "addition 15"
```

RESULT:-

The code has been executed successfully.

EXPERIMENT-3

AIM: Create a list in a data structure that has components of mixed data types.

PROCEDURE:

#a vector having elements of different type is
called list.#Use the list() function to create a list.

```
x <- list(1,"first no.",22, 2, "true")
```

```
print(x) print(paste("type of x",
```

```
class(x)))y<-list("z"=1:10) print(y)
```

```
print(paste("length of list y",
```

```
length(y))) #Multiple lists can be
```

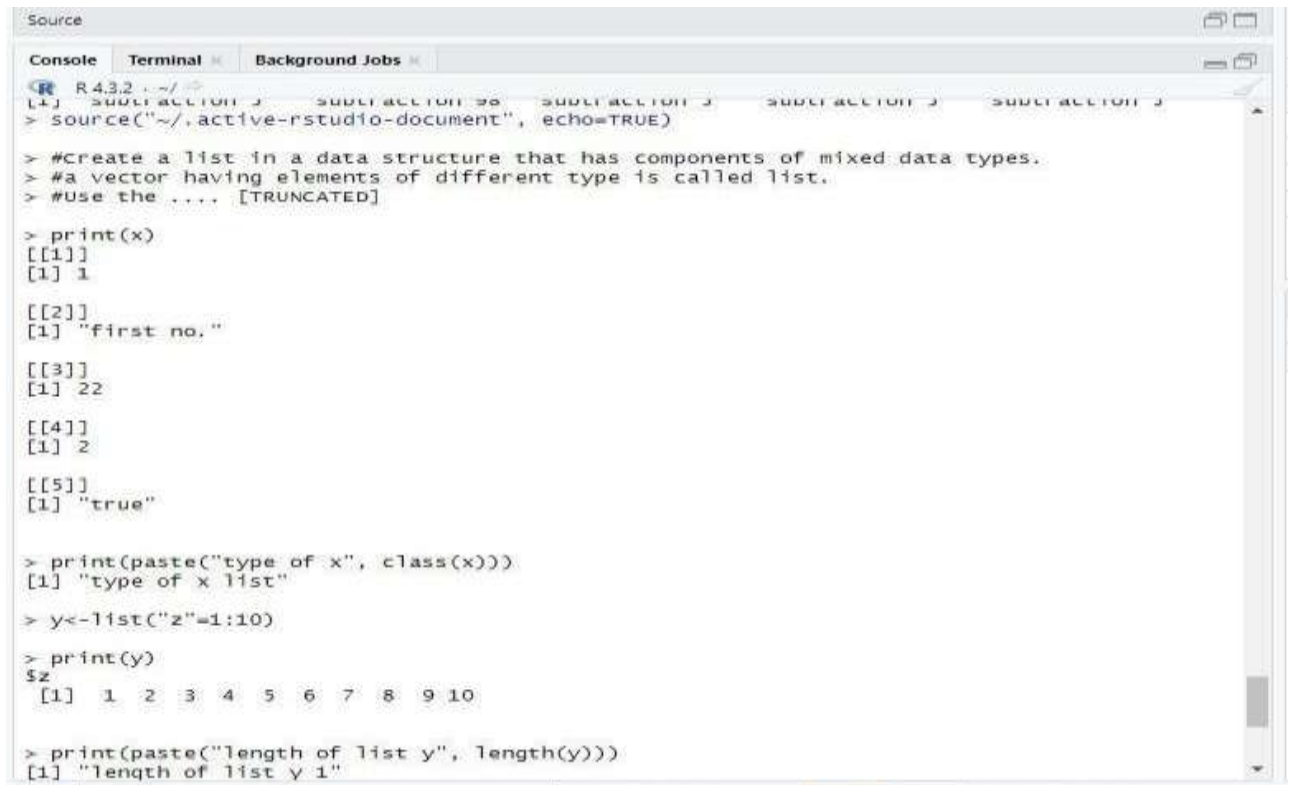
```
merged                list
```

```
combined <- c(x, y)
```

```
#create an empty list of a prespecified length with the vector()
```

```
functionz<- vector("list", length=4 )
```

```
print(z)
```

INPUT/OUTPUT:

```
Source
Console Terminal Background Jobs
R 4.3.2 ~/
[1] subtraction subtraction subtraction subtraction subtraction
> source("~/active-rstudio-document", echo=TRUE)

> #Create a list in a data structure that has components of mixed data types.
> #a vector having elements of different type is called list.
> #Use the .... [TRUNCATED]

> print(x)
[[1]]
[1] 1

[[2]]
[1] "first no."

[[3]]
[1] 22

[[4]]
[1] 2

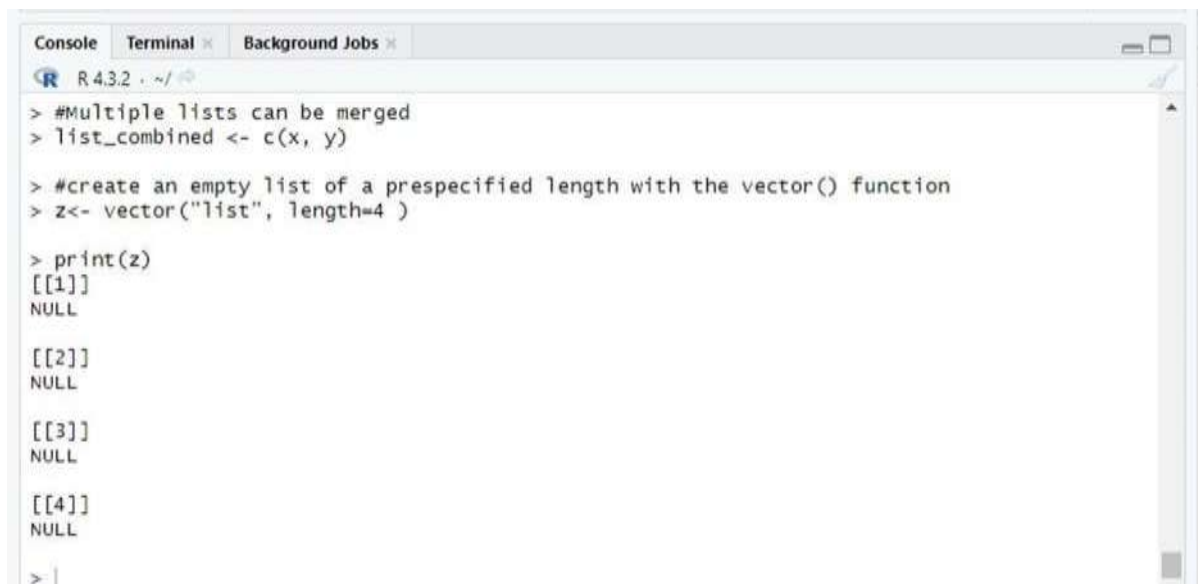
[[5]]
[1] "true"

> print(paste("type of x", class(x)))
[1] "type of x list"

> y<-list("z"=1:10)

> print(y)
$z
[1] 1 2 3 4 5 6 7 8 9 10

> print(paste("length of list y", length(y)))
[1] "length of list y 1"
```



```
R 4.3.2 . ~/
> #Multiple lists can be merged
> list_combined <- c(x, y)

> #create an empty list of a prespecified length with the vector() function
> z<- vector("list", length=4 )

> print(z)
[[1]]
NULL

[[2]]
NULL

[[3]]
NULL

[[4]]
NULL

> |
```

RESULT:

The Code Has Been Executed Successfully .

EXPERIMENT-4

AIM:- Create a code to display Fibonacci series.

PROCEDURE:-

```
print_fibonacci <-
function(n) {a <- 0
b <- 1  cat("Fibonacci Sequence:")  for
  (i in 1:n) {cat(a," ")
next_num <- a + b  a
<- b
  b <- next_num
}
}
number_of_terms<-readline("Enter the number of terms")
print_fibonacci(number_of_terms)
```

OUTPUT:-

```
R version 4.2.2 (2022-10-31 ucrt) -- "Innocent and Trusting"
Copyright (C) 2022 The R Foundation for Statistical Computing
Platform: x86_64-w64-mingw32/x64 (64-bit)

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Type 'license()' or 'licence()' for distribution details.

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Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[workspace loaded from ~/.RData]
> source("~/active-rstudio-document")
Enter the number of terms10
Fibonacci Sequence:0 1 1 2 3 5 8 13 21 34
>
```

RESULT:-

The Code has been executed successfully.

EXPERIMENT-5

Aim: Implement decision tree on credit card issue dataset (import from kaggale).

Procedure:

Implementing a decision tree on a credit card issue dataset involves several steps, including data loading, preprocessing, model training, and evaluation. Below is a simplified example using R and the rpart library for decisiontree modeling. Note that in practice, you might need to adjust the code based on the specific structure and characteristics of your dataset.

First, make sure to install and load the required libraries:
`install.packages("rpart")library(rpart)`

Now, let's assume you have a dataset named "credit_data.csv" with features and labels.

You can load the data and implement a decision tree as follows

```
# Install and load necessary libraries install.packages("rpart ")library(rpart)
```

```
# Load the dataset (replace "credit_data.csv" with your actual file path)
```

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

```
# Explore the structure of the datasetstr(credit_data)
```

```
# Split the data into training and testing sets
set.seed(123)
```

```
# Set seed for reproducibility
```

```
sample_index <- sample(1:nrow(credit_data), 0.8 * nrow(credit_data))train_data
```

```
<- credit_data[sample_index, ] test_data <- credit_data[-
sample_index, ]# Build a decision tree model using rpart
```

```
decision_tree_model <- rpart(Class ~ ., data = train_data, method = "class")
```

```
# Plot the decision tree
```

```

plot(decision_tree_model) text(decision_tree_model, cex = 0.8)

# Make predictions on the test set

predictions <- predict(decision_tree_model, test_data, type = "class")

# Evaluate the model

conf_matrix <- table(predictions, test_data$Class)print("Confusion Matrix:")
print(conf_matrix)

accuracy<- sum(diag(conf_matrix)) / sum(conf_matrix)print(paste("Accuracy:",
round(accuracy, 4)))

```

OUTPUT:-

Console

Terminal x

Background Jobs x

R 4.3.2 · ~/

> # Display the first few rows of the dataset to understand its structure

> head(data)

Time

V1

V2

V3

V4

V5

V6

1

0

-1.3598071

-0.07278117

2.5363467

1.3781552

-0.33832077

0.46238778

2

0

1.1918571

0.26615071

0.1664801

0.4481541

0.06001765

-0.08236081

3

1

-1.3583541

-1.34016307

1.7732093

0.3797796

-0.50319813

1.80049938

4

1

-0.9662717

-0.18522601

1.7929933

-0.8632913

-0.01030888

1.24720317

5

2

-1.1582331

0.87773675

1.5487178

0.4030339

-0.40719338

0.09592146

6

2

-0.4259659

0.96052304

1.1411093

-0.1682521

0.42098688

-0.02972755

V7

V8

V9

V10

V11

V12

1

0.23959855

0.09869790

0.3637870

0.09079417

-0.5515995

-0.61780086

2

-0.07880298

0.08510165

-0.2554251

-0.16697441

1.6127267

1.06523531

3

0.79146096

0.24767579

-1.5146543

0.20764287

0.6245015

0.06608369

4

0.23760894

0.37743587

-1.3870241

-0.05495192

-0.2264873

0.17822823

5

0.59294075

-0.27053268

0.8177393

0.75307443

-0.8228429

0.53819555

6

0.47620095

0.26031433

-0.5686714

-0.37140720

1.3412620

0.35989384

V13

V14

V15

V16

V17

V18

1

-0.9913898

-0.3111694

1.4681770

-0.4704005

0.20797124

0.02579058

2

0.4890950

-0.1437723

0.6355581

0.4639170

-0.11480466

-0.18336127

3

0.7172927

-0.1659459

2.3458649

-2.8900832

1.10996938

-0.12135931

4

0.5077569

-0.2879237

-0.6314181

-1.0596472

-0.68409279

1.96577500

5

1.3458516

-1.1196698

0.1751211

-0.4514492

-0.23703324

-0.03819479

6

-0.3580907

-0.1371337

0.5176168

0.4017259

-0.05813282

0.06865315

V19

V20

V21

V22

V23

V24

1

0.40399296

0.25141210

-0.018306778

0.277837576

-0.11047391

0.06692807

2

-0.14578304

-0.06908314

-0.225775248

-0.638671953

0.10128802

-0.33984648

3

-2.26185710

0.52497973

0.247998153

0.771679402

0.90941226

-0.68928096

4

-1.23262197

-0.20803778

-0.108300452

0.005273597

-0.19032052

-1.17557533

5

0.80348692

0.40854236

-0.009430697

0.798278495

-0.13745808

0.14126698

6

-0.03319379

0.08496767

-0.208253515

-0.559824796

-0.02639767

-0.37142658

V25

V26

V27

V28

Amount

Class

1

0.1285394

-0.1891148

0.133558377

-0.02105305

149.62

0

2

0.1671704

0.1258945

-0.008983099

0.01472417

2.69

0

3

-0.3276418

-0.1390966

-0.055352794

-0.05975184

378.66

0

4

-0.6433760

-0.2210280

-0.06232840

-0.06145763

133.50

0


```
1  0.1285394 -0.1891148  0.133558377 -0.02105305 149.62  0
2  0.1671704  0.1258945 -0.008983099  0.01472417   2.69  0
3 -0.3276418 -0.1390966 -0.055352794 -0.05975184 378.66  0
4  0.6473760 -0.2219288  0.062722849  0.06145763 123.50  0
5 -0.2060096  0.5022922  0.219422230  0.21515315  69.99  0
6 -0.2327938  0.1059148  0.253844225  0.08108026   3.67  0

> # Identify features and target variable
> X <- data[, -c("target_variable")] # Replace 'target_variable' with the actual target column name
```

Result:- Successfully Implemented.

EXPERIMENT-6

Aim : Implement the KNN algorithm on the Brest cancer dataset.

Procedure:

To implement the k-Nearest Neighbors (KNN) algorithm on the Breast Cancer dataset, you can use the caret and class packages in R. The Breast Cancer dataset is often available through the datasets package in R. Here's an example.

```
# Install and load necessary libraries
install.packages("caret")
install.packages("class") library(caret)
library(class)

# Load the Breast Cancer dataset
data("BreastCancer")

# Explore the structure of the dataset
str(BreastCancer)

# Split the data into training and testing sets
set.seed(123)
# Set seed for reproducibility
sample_index <- createDataPartition(BreastCancer$Class, p = 0.8, list = FALSE)
train_data <- BreastCancer[sample_index, ]
test_data <- BreastCancer[-sample_index, ]

# Preprocess the data
# In this example, we'll scale the features
preprocess_params <- preProcess(train_data[, -1], method = c("center", "scale"))

train_data_scaled <- predict(preprocess_params, train_data[, -1]) test_data_scaled <-
predict(preprocess_params, test_data[, -1])

# Train the KNN model
knn_model <- knn(train_data_scaled, test_data_scaled, train_data$Class, k = 5)

# Evaluate the model
conf_matrix <- table(knn_model, test_data$Class) print("Confusion Matrix:")
print(conf_matrix)

accuracy <- sum(diag(conf_matrix)) / sum(conf_matrix) print(paste("Accuracy:",
```

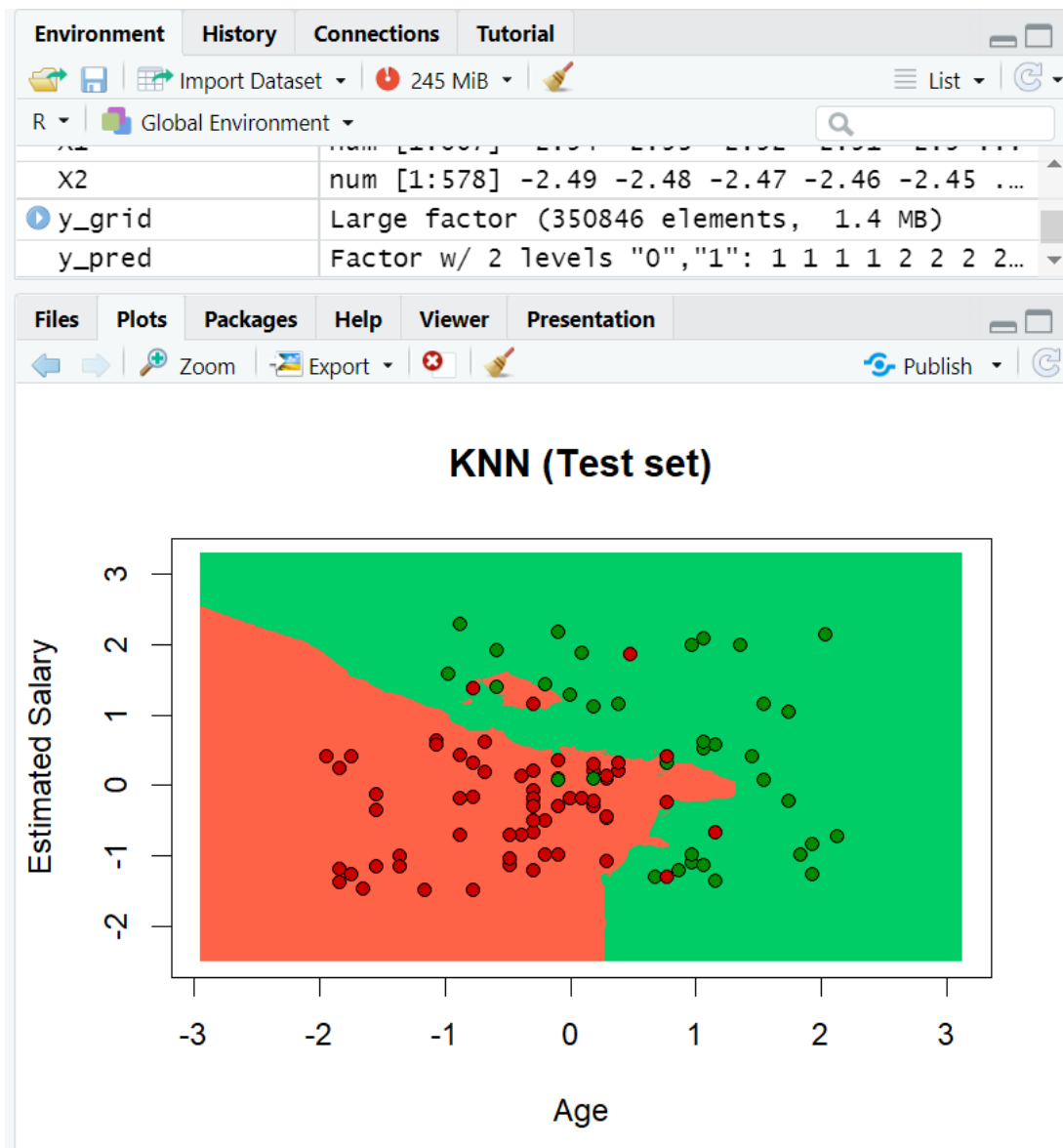
```
round(accuracy, 4)))
```

In this example, the Breast Cancer dataset is split into training and testing sets, and the features are scaled. The KNN model is then trained using the `knn` function from the `class` package, and the accuracy of the model is evaluated.

Remember to replace `BreastCancer$Class` with the actual column name representing the target variable in your dataset. Additionally, consider experimenting with different values of `k` to find the optimal number of neighbors for your specific dataset.

OUTPUT:-

```
> dataset = read.csv('social.csv')
> dataset = dataset[3:5]
>
> # Encoding the label
> dataset$Purchased = factor(dataset$Purchased, level = c(0, 1))
>
> # Splitting the dataset
> library(caTools)
> split = sample.split(dataset$Purchased, SplitRatio = 0.75)
> training_set = subset(dataset, split == TRUE)
> test_set = subset(dataset, split == FALSE)
>
> # Feature Scaling
> training_set[, 1:2] = scale(training_set[, 1:2])
> test_set[, 1:2] = scale(test_set[, 1:2])
>
> # Fitting and predicting the classifier
> library(class)
> y_pred = knn(train = training_set[, 1:2],
+             test = test_set[, 1:2],
+             cl = training_set[, 3],
+             k = 5)
>
> # Creating the confusion matrix
> cm = table(test_set[, 3], y_pred)
>
> set = training_set
> X1 = seq(min(set[, 1]) - 1, max(set[, 1]) + 1, by = 0.01)
> X2 = seq(min(set[, 2]) - 1, max(set[, 2]) + 1, by = 0.01)
> grid_set = expand.grid(X1, X2)
> colnames(grid_set) = c('Age', 'Estimated Salary')
> y_grid = knn(train = training_set[, 1:2],
+             test = grid_set,
+             cl = training_set[, 3],
+             k = 5) # - means removing the column
> plot(set[, -3],
+      main = 'KNN (Training set)',
+      xlab = 'Age', ylab = 'Estimated Salary',
+      xlim = range(X1), ylim = range(X2))
> contour(X1, X2, matrix(as.numeric(y_grid), length(X1), length(X2)), add = TRUE)
> points(grid_set, pch = '.', col = ifelse(y_grid == 1, 'springgreen3', 'tomato'))
> points(set, pch = 21, bg = ifelse(set[, 3] == 1, 'green4', 'red3'))
>
> set = test_set
> X1 = seq(min(set[, 1]) - 1, max(set[, 1]) + 1, by = 0.01)
> X2 = seq(min(set[, 2]) - 1, max(set[, 2]) + 1, by = 0.01)
> grid_set = expand.grid(X1, X2)
> colnames(grid_set) = c('Age', 'Estimated Salary')
> y_grid = knn(train = training_set[, 1:2],
+             test = grid_set,
+             cl = training_set[, 3],
+             k = 5) # - means removing the column
> plot(set[, -3],
+      main = 'KNN (Test set)',
+      xlab = 'Age', ylab = 'Estimated Salary',
+      xlim = range(X1), ylim = range(X2))
> contour(X1, X2, matrix(as.numeric(y_grid), length(X1), length(X2)), add = TRUE)
> points(grid_set, pch = '.', col = ifelse(y_grid == 1, 'springgreen3', 'tomato'))
> points(set, pch = 21, bg = ifelse(set[, 3] == 1, 'green4', 'red3'))
```



Result:- The KNN Algorithm have been implemented successfully.

EXPERIMENT-7

Aim : Implement the Naïve Bayes algorithm on the iris dataset.

Procedure:

To implement the Naive Bayes algorithm on the Iris dataset, you can use the function `naiveByes` from the `e1071` package in R

```
# Install and load necessary libraries
```

```
install.packages("e101") library(e1071)
```

```
# Load the Iris dataset data(iris)
```

```
# Explore the structure of the dataset str(iris)
```

```
# Split the data into training and testing sets
```

```
set.seed(123)
```

```
# Set seed for reproducibility
```

```
sample_index <- sample(1:nrow(iris), 0.8 * nrow(iris)) train_data <- iris[sample_index, ]
```

```
test_data <- iris[-sample_index, ]
```

```
# Train the Naive Bayes model
```

```
naive_bayes_model <- naiveBayes(Species ~ ., data = train_data)
```

```
# Make predictions on the test set
```

```
predictions <- predict(naive_bayes_model, test_data)
```

```
# Evaluate the model
```

```
conf_matrix <- table(predictions, test_data$Species)
```

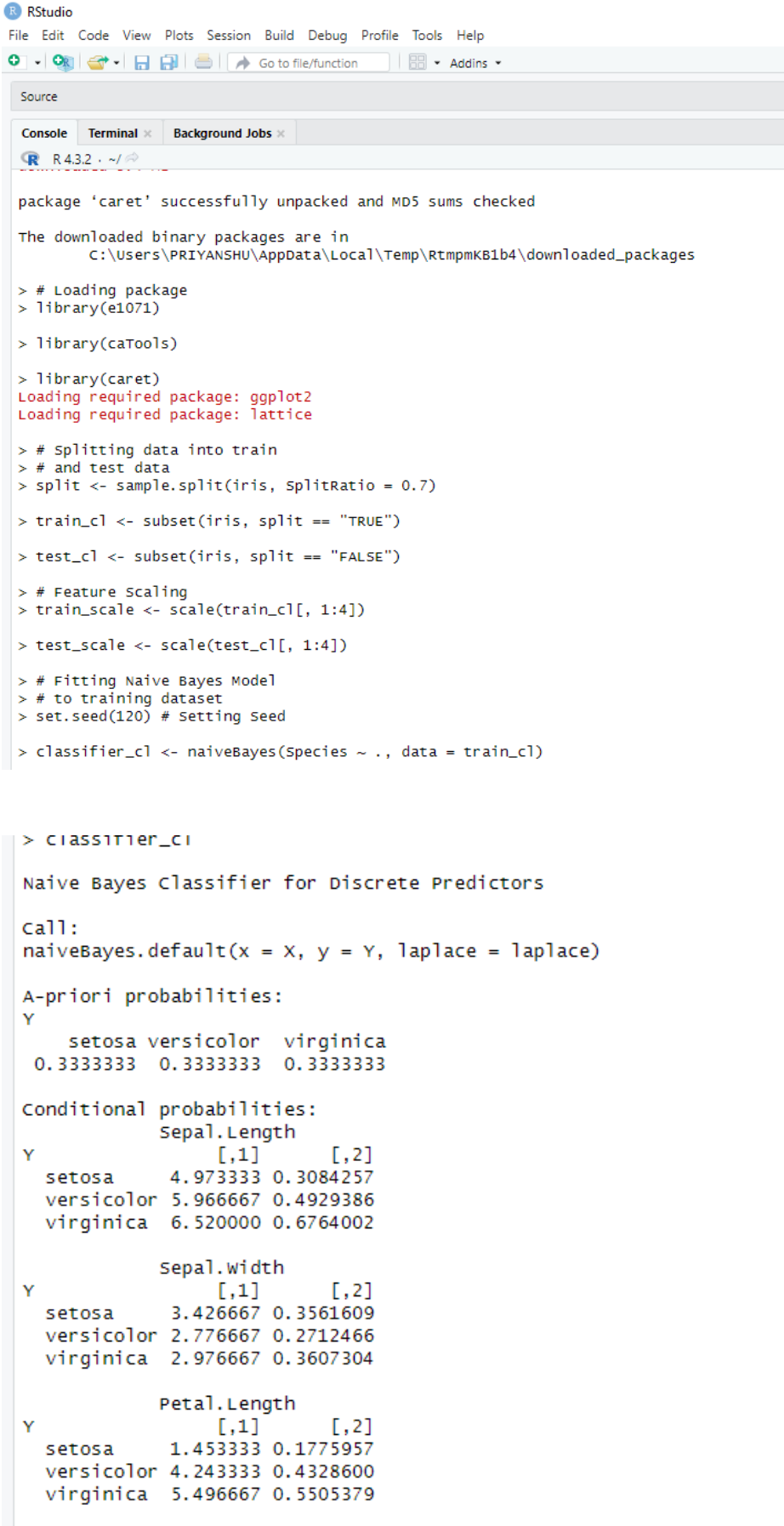
```
print("Confusion Matrix:")
```

```
print(conf_matrix)
```

```
accuracy <- sum(diag(conf_matrix)) / sum(conf_matrix)
```

```
print(paste("Accuracy:", round(accuracy, 4)))
```

OUTPUT:-



RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source

Console Terminal Background Jobs

R 4.3.2 ~ /

```

package 'caret' successfully unpacked and MD5 sums checked

The downloaded binary packages are in
  C:\Users\PRIYANSHU\AppData\Local\Temp\RtmpmKB1b4\downloaded_packages

> # Loading package
> library(e1071)

> library(caTools)

> library(caret)
Loading required package: ggplot2
Loading required package: lattice

> # Splitting data into train
> # and test data
> split <- sample.split(iris, splitRatio = 0.7)

> train_cl <- subset(iris, split == "TRUE")
> test_cl <- subset(iris, split == "FALSE")

> # Feature scaling
> train_scale <- scale(train_cl[, 1:4])
> test_scale <- scale(test_cl[, 1:4])

> # Fitting Naive Bayes Model
> # to training dataset
> set.seed(120) # Setting seed

> classifier_cl <- naiveBayes(Species ~ ., data = train_cl)

> CLASSIFIER_CL

Naive Bayes Classifier for Discrete Predictors

Call:
naiveBayes.default(x = X, y = Y, laplace = laplace)

A-priori probabilities:
Y
      setosa versicolor virginica
0.3333333  0.3333333  0.3333333

Conditional probabilities:
      Sepal.Length
Y      [,1]      [,2]
setosa  4.973333 0.3084257
versicolor 5.966667 0.4929386
virginica  6.520000 0.6764002

      Sepal.width
Y      [,1]      [,2]
setosa  3.426667 0.3561609
versicolor 2.776667 0.2712466
virginica  2.976667 0.3607304

      Petal.Length
Y      [,1]      [,2]
setosa  1.453333 0.1775957
versicolor 4.243333 0.4328600
virginica  5.496667 0.5505379

```

```

> # Model Evaluation
> confusionMatrix(cm)
Confusion Matrix and Statistics

          y_pred
          setosa versicolor virginica
setosa      20          0          0
versicolor   0         19          1
virginica    0          1         19

Overall Statistics

          Accuracy : 0.9667
          95% CI : (0.8847, 0.9959)
    No Information Rate : 0.3333
    P-Value [Acc > NIR] : < 2.2e-16

          Kappa : 0.95

    McNemar's Test P-Value : NA

Statistics by Class:

                class: setosa class: versicolor class: virginica
Sensitivity                1.0000                0.9500                0.9500
Specificity                1.0000                0.9750                0.9750
Pos Pred Value              1.0000                0.9500                0.9500
Neg Pred Value              1.0000                0.9750                0.9750
Prevalence                  0.3333                0.3333                0.3333
Detection Rate              0.3333                0.3167                0.3167
Detection Prevalence        0.3333                0.3333                0.3333
Balanced Accuracy           1.0000                0.9625                0.9625
> |

```

Result:- Successfully Implemented.