Exp8: 210701701

## Implement SVM/Decision tree classification techniques

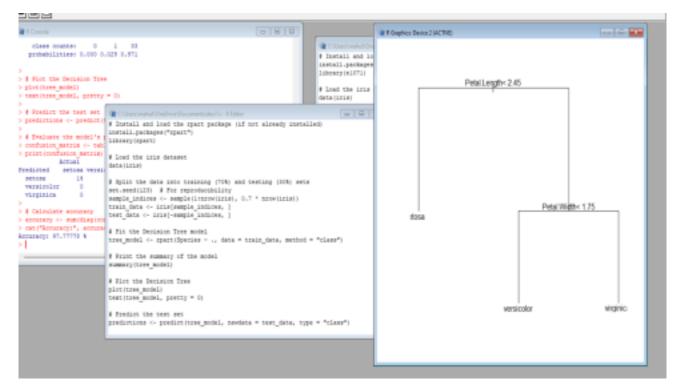
## a) SVM IN R

```
# Install and load the e1071 package (if not already installed)
install.packages("e1071")
library(e1071)
# Load the iris dataset
data(iris)
# Inspect the first few rows of the dataset
head(iris)
# Split the data into training (70%) and testing (30%) sets
set.seed(123) # For reproducibility
sample indices <- sample(1:nrow(iris), 0.7 * nrow(iris))
train data <- iris[sample indices, ]
test data <- iris[-sample indices, ]
# Fit the SVM model
svm model <- svm(Species ~ ., data = train_data, kernel = "radial")
# Print the summary of the model
summary(svm model)
# Predict the test set
predictions <- predict(svm model, newdata = test data)</pre>
# Evaluate the model's performance
confusion matrix <- table(Predicted = predictions, Actual =
test data$Species) print(confusion matrix)
# Calculate accuracy
accuracy <- sum(diag(confusion matrix)) /</pre>
sum(confusion_matrix) cat("Accuracy:", accuracy * 100, "%\n")
```

```
# Inspect the first few rows of the dataset
head(ints)
                                                                                                                                                           # Iglit the data into training (704) and testing (304) sets
set.ored(223) # For expendiculativy
sample_indices - sample(introv(1716), 0.7 * nrow(inte))
train_data <- iris[sample_indoose, ]
test_data <- iris[sample_indoose, ]
                                                                                                                                                           # Fix the SVM model
ovm_model <- svm:Species - ., data = train_data, bernel = "radial")
# Calculate accuracy accuracy = 100, "Fin") / Fun (confusion_matrix
                                                                                                                                                            # Fredict the test set
poedictions <- poedict(smm_model, newdata = test_data)
```

## b) Decision tree in R

```
# Install and load the rpart package (if not already installed)
install.packages("rpart")
library(rpart)
# Load the iris dataset
data(iris)
# Split the data into training (70%) and testing (30%) sets
set.seed(123) # For reproducibility
sample indices <- sample(1:nrow(iris), 0.7 * nrow(iris))
train data <- iris[sample indices, ]
test data <- iris[-sample indices, ]
# Fit the Decision Tree model
tree model <- rpart(Species ~ ., data = train data, method = "class")
# Print the summary of the model
summary(tree model)
# Plot the Decision Tree
plot(tree model)
text(tree model, pretty = 0)
# Predict the test set
predictions <- predict(tree model, newdata = test data, type = "class")</pre>
# Evaluate the model's performance
confusion matrix <- table(Predicted = predictions, Actual =
test data$Species) print(confusion_matrix)
# Calculate accuracy
accuracy <- sum(diag(confusion matrix)) / sum(confusion matrix)</pre>
cat("Accuracy:", accuracy * 100, "%\n")
```



## **RESULT:**

Thus the implementation of SVM/Decision tree classification techniques has been executed in the R programming language successfully.