Exp No: 8

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)) / sum(confusion matrix) cat("Accuracy:",
accuracy * 100, "%\n")
```

```
Console Terminal × Background Jobs ×
 R 4.4.1 · ~/ ≈
 setosa versicolor virginica
> # 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)</pre>
> print(confusion_matrix)
             Actual
Predicted
               setosa versicolor virginica
  setosa
                   14
                                0
                                            0
                    0
                                17
  versicolor
  virginica
                                 1
                                           13
> # Calculate accuracy
> accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
> cat("Accuracy:", accuracy * 100, "%\n")
Accuracy: 97.77778 %
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) cat("Accuracy:",
accuracy * 100, "%\n")
```

