## 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, ]</pre>
test_data <- iris[-sample_indices, ]</pre>
# Fit the SVM model
svm_model <- svm(Species ~ ., data = train_data, kernel = "radial")</pre>
# 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)</pre>
print(confusion_matrix)
# Calculate accuracy
accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
cat("Accuracy:", accuracy * 100, "%\n")
```

```
> # 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))</pre>
> train_data <- iris[sample_indices, ]</pre>
> test_data <- iris[-sample_indices, ]</pre>
> # Fit the SVM model
> svm_model <- svm(Species ~ ., data = train_data, kernel = "radial")</pre>
> # Print the summary of the model
> summary(svm_model)
svm(formula = Species ~ ., data = train_data, kernel = "radial")
Parameters:
  SVM-Type: C-classification
 SVM-Kernel: radial
       cost: 1
Number of Support Vectors: 45
 (7 18 20)
Number of Classes: 3
Levels:
 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
              14 0
                                   0
  setosa
               0
  versicolor
                            17
                                        0
  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, ]</pre>
test_data <- iris[-sample_indices, ]</pre>
# 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)</pre>
print(confusion_matrix)
# Calculate accuracy
accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
cat("Accuracy:", accuracy * 100, "%\n")
```

#### 

