

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)

# 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")
```

```

> # 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)

Call:
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)
> # Evaluate the model's performance
> confusion_matrix <- table(Predicted = predictions, Actual = test_data$Species)
> print(confusion_matrix)
      Actual
Predicted setosa versicolor virginica
setosa      14           0           0
versicolor  0          17           0
virginica   0           1          13
> # Calculate accuracy
> accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)
> 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")
# 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")

```

