### **Practical No: 7**

#### **Feature Selection**

AIM: Data loading, feature scoring and ranking, feature selection (principal component analysis)

### **Description:**

## **Principal Component Analysis:**

Principal Component Analysis (PCA) is a handy tool in statistics for making complex data easier to understand. Before diving into it, it's crucial to standardize the data, which basically means making sure all the different measurements are on the same scale. Next, PCA calculates something called the covariance matrix, which shows how different variables in the data relate to each other. Then comes the interesting part: eigendecomposition. This involves finding special directions, called eigenvectors, and their associated importance values, called eigenvalues. The eigenvectors with the highest eigenvalues become the principal components, kind of like the main characters in our data story. These principal components capture the most important aspects of the data. By selecting a few of these components, we can simplify the data without losing much information. People often use PCA to make big datasets more manageable, create cool visualizations, and reduce noise in the data. It's like turning a complicated puzzle into a simpler picture, making it easier for us to see the patterns and trends.

# **Code with output**

import numpy as np

import pandas as pd

from sklearn.model\_selection import train\_test\_split

from sklearn.preprocessing import StandardScaler

from sklearn.decomposition import PCA

from sklearn.ensemble import RandomForestClassifier

from sklearn.metrics import confusion\_matrix, accuracy\_score

Name: Ninad Karlekar Roll no.: 22306A1012

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```
url = "https://archive.ics.uci.edu/ml/machine-learning-databases/iris/iris.data"
names = ['sepal-length', 'sepal-width', 'petal-length', 'petal-width', 'Class']
dataset = pd.read_csv(url, names=names)
dataset.head()
# Store the feature sets into X variable and the series of corresponding variables in y
x = dataset.drop('Class', axis=1)
y = dataset['Class']
x.head()
y.head()
x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.2, random_state=0)
sc = StandardScaler()
x_train1 = sc.fit_transform(x_train)
x_{test1} = sc.transform(x_{test})
y_{train} = y_{train}
y_{test1} = y_{test}
pca = PCA()
x_train1 = pca.fit_transform(x_train1)
x_{test1} = pca.transform(x_{test1})
explained_variance = pca.explained_variance_ratio_
print(explained_variance)
pca = PCA(n_components=1)
x_train1 = pca.fit_transform(x_train1)
x_{test1} = pca.transform(x_{test1})
classifier = RandomForestClassifier(max_depth=2, random_state=0)
classifier.fit(x_train1, y_train1)
```

```
y_pred = classifier.predict(x_test1)

cm = confusion_matrix(y_test, y_pred)
print(cm)
print('Accuracy:', accuracy_score(y_test, y_pred))
```

# **OUTPUT**

```
[0.72226528 0.23974795 0.03338117 0.0046056 ]
[0.72226528 0.23974795 0.03338117 0.0046056 ]
[[11 0 0]
[ 0 12 1]
[ 0 1 5]]
Accuracy 0.9333333333333333
```

Name: Ninad Karlekar Roll no.: 22306A1012

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## Learnings

The code first loads the Iris dataset, which is a collection of data about Iris flowers. The data includes four features: sepal length, sepal width, petal length, and petal width. The data also includes a label, which is the species of the Iris flower.

The code then splits the data into two sets: a training set and a test set. The training set is used to train the machine learning model, and the test set is used to evaluate the model's performance.

Before training the model, the code standardizes the features. This means that the code scales the features so that they are all on the same scale. This is important because it ensures that the model does not learn to bias towards any particular feature.

The code then uses PCA to reduce the dimensionality of the features. This means that the code combines the four features into a single feature. This can improve the model's performance because it reduces noise in the data and makes it easier for the model to learn the patterns.

The code then trains a Random Forest Classifier on the training set. A Random Forest Classifier is a type of machine learning model that is well-suited for classification tasks.

Once the model is trained, the code uses it to make predictions on the test set. The code then evaluates the model's performance using a confusion matrix and accuracy score. A confusion matrix shows how many flowers of each species were correctly and incorrectly classified by the model. An accuracy score is a measure of how often the model correctly classified a flower.

Name: Ninad Karlekar Roll no.: 22306A1012 4