Lab Assignment: Dimensionality Reduction Using PCA and t-SNE

Objective:

The goal of this lab is to reduce the dimensionality of a dataset using **Principal Component Analysis (PCA)** and **t-SNE**. Students will learn how dimensionality reduction helps in visualizing high-dimensional data and improving model performance.

Tasks:

1. Load the Dataset

- o Use the **Wine Dataset** from sklearn.datasets. This dataset consists of 13 features related to wine classification.
- o Display the first 5 rows and describe the dataset.
- o Split the dataset into features (X) and target labels (y).

2. Data Standardization

o Standardize the dataset using StandardScaler from sklearn.preprocessing. Standardization is important for PCA and t-SNE.

3. Principal Component Analysis (PCA)

- o Apply **PCA** to reduce the dataset to 2 components respectively.
- o For the 2-component PCA, visualize the data using a 2D scatter plot, with points colored based on their target labels.

4. t-SNE Visualization

- Use **t-SNE** to reduce the dimensionality of the dataset to 2 dimensions.
- Visualize the t-SNE transformed data using a 2D scatter plot and color the points based on their target labels.

5. K-Means Clustering on Reduced Data

- o Apply **K-Means Clustering** on the 2-component PCA reduced data.
- o Compare the clustering performance (using adjusted Rand index or silhouette score) with clustering performed on the original high-dimensional dataset.
- **Silhouette Score**: Measures how similar an object is to its own cluster (cohesion) compared to other clusters (separation). It ranges from -1 to 1, where a higher value indicates better-defined clusters.

$$s(i) = \frac{b(i) - a(i)}{\max(a(i), b(i))}$$

Where:

- o a(i)a(i)a(i) is the average intra-cluster distance (how close a point is to other points in the same cluster).
- o b(i)b(i)b(i) is the average nearest-cluster distance (how close a point is to points in the nearest neighboring cluster).

```
from sklearn.datasets import make_blobs
from sklearn.cluster import KMeans
from sklearn.metrics import silhouette_score
X, y = make_blobs(random_state=42)
kmeans = KMeans(n_clusters=2, random_state=42)
silhouette score(X, kmeans.fit predict(X))
```

```
Example t-SNE
                                      Example PCA
model = TSNE(n_components = 2,
                                      import numpy as np
random state = 0)
                                      >>> from sklearn.decomposition
# configuring the parameters
                                      import PCA
\# the number of components = 2
                                      X = np.array([[-1, -1], [-2, -1],
\# default perplexity = 30
# default learning rate = 200
                                      [-3, -2], [1, 1], [2, 1], [3, 2]])
                                      pca = PCA(n components=2)
# default Maximum number of
iterations
                                      pca.fit(X)
\# for the optimization = 1000
                                      PCA(n components=2)
tsne data =
model.fit transform(data 1000)
# creating a new data frame which
# help us in plotting the result
data
tsne data = np.vstack((tsne data.T,
labels_1000)).T
tsne_df = pd.DataFrame(data =
tsne_data,
    ______columns = ("Dim 1", "Dim 2",
"label"))
# Plotting the result of tsne
sn.scatterplot(data=tsne df,
x='Dim 1', y='Dim 2',
               hue='label',
palette="bright")
plt.show()
```