1. Implement the k-Nearest Neighbor (k-NN) algorithm to classify a given dataset using Minkowski distance for p=3. Evaluate the accuracy of the classifier. CODE: import pandas as pd from sklearn.model selection import train test split from sklearn.neighbors import KNeighborsClassifier from sklearn.metrics import accuracy score # Load dataset file path = r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3 (1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" data = pd.read csv(file path) # Prepare features and target X = data.iloc[:, :-1] # Assuming last column is the target y = data.iloc[:, -1] # Assuming last column is the target # Split dataset X train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.3, random\_state=42) # k-NN with Minkowski distance (p=3) knn = KNeighborsClassifier(n\_neighbors=3, metric='minkowski', p=3) knn.fit(X train, y train) # Predictions and accuracy

# If only some selected features have to be included in the code

y pred = knn.predict(X test)

print(f"Accuracy: {accuracy:.2f}%")

accuracy = accuracy\_score(y\_test, y\_pred) \* 100

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score

# Load dataset
file_path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e0
7d5ae3 (1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv"
data = pd.read_csv(file_path)
print(data.columns.tolist())
```

```
# Select specific features (replace with the actual column names or
indices)
selected features = ['feature 1', 'feature 2', 'feature 3'] #
Replace with your column names
X = data[selected features]
# Target column
y = data['target'] # Replace with the actual target column name
# Split dataset
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.3, random state=42)
# k-NN with Minkowski distance (p=3)
knn = KNeighborsClassifier(n neighbors=3, metric='minkowski', p=3)
knn.fit(X train, y train)
# Predictions and accuracy
y pred = knn.predict(X test)
accuracy = accuracy score(y test, y pred) * 100
print(f"Accuracy: {accuracy:.2f}%")
```

2. Implement the Iterative Dichotomiser (ID3) algorithm with entropy as the criterion to build a decision tree using a given dataset. Evaluate the classifier by computing its accuracy.

## CODE:

```
from sklearn.tree import DecisionTreeClassifier from sklearn.model_selection import train_test_split from sklearn.metrics import accuracy_score import pandas as pd
```

```
# Load dataset file_path = r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3 (1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your dataset file path data = pd.read_csv(file_path)
```

# Prepare features and target

```
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Decision Tree with entropy
dt = DecisionTreeClassifier(criterion='entropy')
dt.fit(X train, y train)
# Evaluate
y pred = dt.predict(X_test)
accuracy = accuracy score(y test, y pred)
print(f"Decision Tree (Entropy) Accuracy: {accuracy:.2f}")
3.Implement feature reduction using Principal Component Analysis by at least one dimension
for a given dataset. Evaluate the performance of Logistic regression before and after applying
PCA.
CODE:
from sklearn.decomposition import PCA
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy score
import pandas as pd
# Load dataset
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read_csv(file_path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Logistic Regression before PCA
Ir = LogisticRegression()
Ir.fit(X train, y train)
```

```
y_pred = Ir.predict(X_test)
accuracy_before = accuracy_score(y_test, y_pred)
#PCA
pca = PCA(n_components=X_train.shape[1] - 1)
X train pca = pca.fit transform(X train)
X test pca = pca.transform(X test)
# Logistic Regression after PCA
Ir pca = LogisticRegression()
Ir_pca.fit(X_train_pca, y_train)
y_pred_pca = Ir_pca.predict(X_test_pca)
accuracy_after = accuracy_score(y_test, y_pred_pca)
print(f"Logistic Regression Accuracy Before PCA: {accuracy before:.2f}")
print(f"Logistic Regression Accuracy After PCA: {accuracy_after:.2f}")
4. Implement the na ive Bayesian classifier for a given data set. Compute the accuracy of the
classifier, considering few test data. Sets.
CODE:
from sklearn.naive bayes import GaussianNB
from sklearn.metrics import accuracy_score
import pandas as pd
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read csv(file path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Naive Bayes
nb = GaussianNB()
nb.fit(X_train, y_train)
```

```
# Evaluate
y_pred = nb.predict(X_test)
accuracy = accuracy score(y test, y pred)
print(f"Naive Bayes Accuracy: {accuracy:.2f}")
```

Implement Support Vector Machine (SVM) model for a given data for Kernels :'linear rbf','sigmoid'. Plot the support vectors and with regions of classes in each case. Evaluate thei performance on the test data and suggest the best fitting kernel

```
CODE:
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
import pandas as pd
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read_csv(file_path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
kernels = ['linear', 'poly', 'rbf', 'sigmoid']
for kernel in kernels:
  svm = SVC(kernel=kernel)
  svm.fit(X_train, y_train)
  y_pred = svm.predict(X_test)
  accuracy = accuracy_score(y_test, y_pred)
  print(f"SVM Accuracy with {kernel} kernel: {accuracy:.2f}")
  # Optional: Plot decision regions if dataset is 2D
  if X_{train.shape[1]} == 2:
    plt.figure()
     plt.scatter(X_train.iloc[:, 0], X_train.iloc[:, 1], c=y_train, cmap='viridis', s=50, alpha=0.5)
```

```
plt.title(f"SVM Decision Regions ({kernel})")
plt.show()
```

6. Implement Rosenblatt's perceptron model for the Boolean expression ((p  $\land$  q)  $\lor$  r)  $\rightarrow$  (p  $\land$  r) and evaluate its accuracy.

```
CODE
import numpy as np
from sklearn.linear model import Perceptron
from sklearn.metrics import accuracy_score
# Define the truth table for the Boolean expression ((p \land q) \lor r) \rightarrow (p \land ~r)
data = []
for p in [0, 1]:
  for q in [0, 1]:
     for r in [0, 1]:
        # Calculate ((p \land q) \lor r) and (p \land ~r)
       left = (p and q) or r
        right = p and not r
        result = int(left <= right) # Implication
        data.append([p, q, r, result])
data = np.array(data)
# Input (X) and target (y)
X = data[:, :-1]
y = data[:, -1]
# Train Rosenblatt's Perceptron
perceptron = Perceptron(max iter=1000, tol=1e-3, random state=42)
perceptron.fit(X, y)
# Evaluate
y_pred = perceptron.predict(X)
accuracy = accuracy_score(y, y_pred)
print(f"Rosenblatt's Perceptron Accuracy: {accuracy:.2f}")
```

7. Implement polynomial regression using Stochastic Gradient Descent for the given dataset, plot the accuracy for different degrees and conclude the best fit.

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.preprocessing import PolynomialFeatures
from sklearn.linear model import SGDRegressor
from sklearn.metrics import mean squared error
from sklearn.model_selection import train_test_split
# Specify the file path for the Iris dataset (CSV file)
file path = r"C:\Users\tanuj\OneDrive\Desktop\iris.csv" # Replace with the actual path to your
CSV file
# Load dataset
data = pd.read csv(file path)
# Select features and target columns
# Assuming Sepal Length (1st column) as X and Petal Length (3rd column) as y
X = data.iloc[:, 0].values.reshape(-1, 1) # Feature
y = data.iloc[:, 2].values.reshape(-1, 1) # Target
# Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Evaluate polynomial regression for different degrees
degrees = range(1, 10)
train_errors = []
test errors = []
for degree in degrees:
  # Polynomial feature transformation
  poly = PolynomialFeatures(degree=degree, include bias=False)
  X train poly = poly.fit transform(X train)
  X_test_poly = poly.transform(X_test)
  # SGD Regressor
  sgd reg = SGDRegressor(max iter=1000, tol=1e-3, random state=42)
  sgd reg.fit(X train poly, y train.ravel())
  # Predictions and errors
  y train pred = sgd reg.predict(X train poly)
  y_test_pred = sgd_reg.predict(X_test_poly)
  train errors.append(mean squared error(y train, y train pred))
  test_errors.append(mean_squared_error(y_test, y_test_pred))
```

```
# Plot the errors for different polynomial degrees
plt.figure(figsize=(10, 6))
plt.plot(degrees, train_errors, label="Train Error", marker='o')
plt.plot(degrees, test_errors, label="Test Error", marker='o')
plt.xlabel("Polynomial Degree")
plt.ylabel("Mean Squared Error")
plt.title("Polynomial Degree vs. Error")
plt.legend()
plt.grid(True)
plt.show()

# Best degree conclusion
best_degree = degrees[np.argmin(test_errors)]
print(f"The best polynomial degree is {best_degree} with test error {min(test_errors):.2f}.")
```

8.Implement multiple linear regression using Stochastic Gradient Descent for the given dataset and compute the accuracy on the test data.

```
from sklearn.linear_model import SGDRegressor
from sklearn.metrics import mean squared error, r2 score
from sklearn.model selection import train test split
from sklearn.preprocessing import LabelEncoder
import pandas as pd
# Load dataset
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read csv(file path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
label_encoder = LabelEncoder()
y = label encoder.fit transform(y)
# Split dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Train Multiple Linear Regression using SGD
sgd = SGDRegressor(max iter=1000, tol=1e-3, random state=42)
sgd.fit(X train, y train)
# Predict on test data
y_pred = sgd.predict(X_test)
# Evaluate performance
mse = mean_squared_error(y_test, y_pred)
r2 = r2_score(y_test, y_pred)
# Output results
print(f"Mean Squared Error (MSE): {mse:.4f}")
print(f"R-squared (R2) Score: {r2:.4f}")
```

9. Implement the Iterative Dichotomiser (ID3) algorithm with Gini as the criterion to build a decision tree for a given dataset. Evaluate the classifier by computing its accuracy.

```
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.metrics import accuracy score
from sklearn.model_selection import train_test_split
import pandas as pd
import matplotlib.pyplot as plt
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read_csv(file_path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Train Decision Tree using Gini criterion
dt = DecisionTreeClassifier(criterion='gini', random_state=42)
dt.fit(X train, y train)
# Predict on test data
y pred = dt.predict(X test)
# Evaluate accuracy
accuracy = accuracy_score(y_test, y_pred)
print(f"Decision Tree (Gini) Accuracy: {accuracy:.4f}")
# Visualize the Decision Tree
plt.figure(figsize=(12, 8))
plot_tree(dt, feature_names=X.columns, class_names=dt.classes_.astype(str), filled=True,
rounded=True)
plt.title("Decision Tree Visualization (Gini)")
plt.show()
```

10. Implement the ADALINE model using the Delta Rule for binary classification for a given dataset. Evaluate the classifier by computing its accuracy.

```
import numpy as np
import pandas as pd # Import pandas for reading datasets
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.preprocessing import LabelEncoder
class Adaline:
  def __init__(self, lr=0.01, epochs=100):
     self.lr = Ir
     self.epochs = epochs
  def fit(self, X, y):
     self.weights = np.zeros(X.shape[1])
     self.bias = 0
    for in range(self.epochs):
       for xi, target in zip(X, y):
          y_pred = np.dot(xi, self.weights) + self.bias # Linear output
          error = target - y pred
          self.weights += self.lr * error * xi # Weight update using Delta Rule
          self.bias += self.lr * error # Bias update
  def predict(self, X):
     # Apply the sign function for binary classification
     return (np.dot(X, self.weights) + self.bias >= 0).astype(int)
# Load dataset
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read csv(file path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
label encoder = LabelEncoder()
y = label encoder.fit transform(y)
```

```
# Normalize the features
scaler = StandardScaler()
X = scaler.fit_transform(X)

# Ensure the target is binary (convert if necessary)
#y = (y == some_condition).astype(int) # Replace `some_condition` with appropriate logic if needed

# Split into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)

# Initialize and train the ADALINE model
model = Adaline(Ir=0.01, epochs=1000)
model.fit(X_train, y_train)

# Evaluate the model
accuracy = np.mean(model.predict(X_test) == y_test)
print("Accuracy:", accuracy)

11. Implement ADALINE for a regression task using sigmoid activation and delta rule, where
```

11. Implement ADALINE for a regression task using sigmoid activation and delta rule, where the model predicts continuous values for a given dataset. Compute the mean squared error on the test data.

# CODE:

## #11

from sklearn.linear\_model import SGDRegressor from sklearn.metrics import mean\_squared\_error from sklearn.model\_selection import train\_test\_split from sklearn.preprocessing import LabelEncoder import pandas as pd import numpy as np

# Sigmoid activation function
def sigmoid(x):
 return 1 / (1 + np.exp(-x))

# Load dataset

file path =

r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3

```
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read csv(file path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
# Split dataset into training and testing sets
# Split dataset into training and testing sets
label encoder = LabelEncoder()
y = label_encoder.fit_transform(y)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Train ADALINE model using SGD (Delta Rule for regression)
adaline = SGDRegressor(max_iter=1000, tol=1e-3, random_state=42)
adaline.fit(X train, y train)
# Predict using the trained model
y pred linear = adaline.predict(X test)
# Apply sigmoid activation to predictions
y_pred_sigmoid = sigmoid(y_pred_linear)
# Evaluate the model
mse = mean_squared_error(y_test, y_pred_sigmoid)
print(f"ADALINE Regression MSE (with Sigmoid Activation): {mse:.4f}")
```

12. Implement the MADALINE model for a regression task, where multiple ADALINE neurons are used in the hidden layer to predict continuous values with ReLU activations. Compute the mean squared error on the test data.

## CODE:

from sklearn.linear\_model import LinearRegression from sklearn.metrics import mean\_squared\_error from sklearn.model\_selection import train\_test\_split from sklearn.preprocessing import LabelEncoder import pandas as pd import numpy as np

# ReLU activation function

```
def relu(x):
  return np.maximum(0, x)
# Load dataset
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read csv(file path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
label encoder = LabelEncoder()
y = label_encoder.fit_transform(y)
# Split dataset
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
# First Layer: Hidden layer with ReLU
hidden layer weights = np.random.randn(X train.shape[1], 64) # Random weights
hidden_layer_bias = np.random.randn(64)
X train hidden = relu(np.dot(X train, hidden layer weights) + hidden layer bias)
X test hidden = relu(np.dot(X test, hidden layer weights) + hidden layer bias)
# Output Layer: Linear Regression
regressor = LinearRegression()
regressor.fit(X train hidden, y train)
# Predict and evaluate
y pred = regressor.predict(X test hidden)
mse = mean squared error(y test, y pred)
print(f"MADALINE Regression MSE (Alternative Approach): {mse:.4f}")
```

13. Implement the MADALINE model for a binary classification task, where multiple ADALINE neurons are used in the hidden layers with ReLU activations. Compute the accuracy on the test data.

### CODE

from sklearn.linear\_model import LogisticRegression from sklearn.metrics import accuracy\_score, classification\_report from sklearn.model\_selection import train\_test\_split from sklearn.preprocessing import StandardScaler

```
from sklearn.preprocessing import LabelEncoder
import pandas as pd
import numpy as np
# ReLU activation function
def relu(x):
  return np.maximum(0, x)
# Load dataset
file path =
r"C:\Users\jslam\Downloads\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3
(1)\8836201-6f9306ad21398ea43cba4f7d537619d0e07d5ae3\iris.csv" # Replace with your
dataset file path
data = pd.read_csv(file_path)
# Prepare features and target
X = data.iloc[:, :-1] # Assuming last column is the target
y = data.iloc[:, -1] # Assuming last column is the target
label encoder = LabelEncoder()
y = label encoder.fit transform(y)
# Split dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# First Layer: Hidden layer with ReLU activation
hidden layer weights = np.random.randn(X train.shape[1], 64) # Random weights
hidden layer bias = np.random.randn(64)
X train hidden = relu(np.dot(X train, hidden layer weights) + hidden layer bias)
X_test_hidden = relu(np.dot(X_test, hidden_layer_weights) + hidden_layer_bias)
# Scale the hidden layer output
scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_train_hidden)
X_test_scaled = scaler.transform(X_test_hidden)
# Logistic Regression: Output layer
classifier = LogisticRegression(solver='lbfgs', max_iter=1000, random_state=42) # Increased
max iter
classifier.fit(X_train_scaled, y_train)
# Predict and evaluate
y_pred = classifier.predict(X_test_scaled)
accuracy = accuracy score(y test, y pred)
print(f"\nMADALINE Classification Accuracy: {accuracy:.4f}")
```

```
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
14.Apply Linear Discriminant Analysis (LDA) for feature reduction to improve classification
performance of Logistic regression on a given dataset.
CODE:
import numpy as np
import pandas as pd
from sklearn.model selection import train test split
from sklearn.preprocessing import LabelEncoder, StandardScaler
from sklearn.discriminant analysis import LinearDiscriminantAnalysis as LDA
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, classification_report
# Load dataset from a user-defined file path
file path = r"C:\Users\tanuj\OneDrive\Desktop\iris.csv" # Replace with the actual path to your
CSV file
data = pd.read_csv(file_path)
# Extract features (X) and target (y)
X = data.iloc[:, :-1].values # Features (all columns except the last)
y = data.iloc[:, -1].values # Target (last column)
# Encode target variable if it is categorical
label encoder = LabelEncoder()
y = label_encoder.fit_transform(y)
# Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Standardize features
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X test = scaler.transform(X test)
```

# Apply Linear Discriminant Analysis (LDA)

X test Ida = Ida.transform(X test)

X\_train\_lda = Ida.fit\_transform(X\_train, y\_train)

Ida = LDA(n components=None) # Retain all components initially

```
# Check the explained variance ratio to determine retained components
explained_variance_ratio = Ida.explained_variance_ratio_
print("Explained Variance Ratio for each LDA component:", explained variance ratio)
# Train Logistic Regression on reduced features
log reg = LogisticRegression()
log reg.fit(X train Ida, y train)
# Predict and evaluate
y train pred = log reg.predict(X train lda)
y_test_pred = log_reg.predict(X_test_lda)
train accuracy = accuracy score(y train, y train pred)
test_accuracy = accuracy_score(y_test, y_test_pred)
print(f"Training Accuracy: {train_accuracy * 100:.2f}%")
print(f"Testing Accuracy: {test_accuracy * 100:.2f}%")
print("\nClassification Report (Testing):")
print(classification_report(y_test, y_test_pred))
15.15. Implement the K-means clustering algorithm for a given dataset. Plot the performance
graph Inertia vs k.
CODE:
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.datasets import make blobs
# Generate synthetic dataset
X, _ = make_blobs(n_samples=300, centers=4, cluster_std=1.0, random_state=42)
# Compute inertia for k = 1 to 10
inertias = [KMeans(n_clusters=k, random_state=42).fit(X).inertia_ for k in range(1, 11)]
# Plot Inertia vs k
plt.plot(range(1, 11), inertias, marker='o')
plt.xlabel("Number of Clusters (k)")
plt.vlabel("Inertia")
plt.title("K-means: Inertia vs k")
plt.show()
16.
import numpy as np
import pandas as pd
```

```
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import AgglomerativeClustering
from sklearn.metrics import silhouette_score
import scipy.cluster.hierarchy as sch
data = load iris()
X = data.data
y = data.target
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
clustering = AgglomerativeClustering(n_clusters=3, metric='euclidean', linkage='ward')
y pred = clustering.fit predict(X scaled)
plt.figure(figsize=(10, 7))
sch.dendrogram(sch.linkage(X_scaled, method='ward'))
plt.title('Dendrogram for Hierarchical Clustering')
plt.xlabel('Samples')
plt.ylabel('Distance')
plt.show()
sil score = silhouette score(X scaled, y pred)
print(f"Silhouette Score: {sil_score:.2f}")
```