#naive bayes age, salary, purchased 1 import pandas as pd import numpy as np from sklearn.model selection import train test split from sklearn.naive bayes import GaussianNB from sklearn.metrics import accuracy_score, confusion_matrix, classification_report import matplotlib.pyplot as plt import seaborn as sns data = pd.read_csv('NaiveBayes.csv') data.head() X = data[['Age', 'Salary']] y = data['Purchased'] X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2, random_state = 42) nb_classifier = GaussianNB() y_pred = nb_classifier.predict(X_test) accuracy = accuracy_score(y_test, y_pred) print(f"Accuracy :{accuracy*100:.2f}") cm = confusion matrix(y test, y pred) print("Confusion Matrix\n", cm) cr = classification report(y test, y pred) print("Classification Report\n", cr) #naive bayes prima indians diabetes 2 import pandas as pd import numpy as np import matplotlib.pyplot as plt import seaborn as sns from sklearn.model selection import train test split from sklearn.naive_bayes import GaussianNB from sklearn.metrics import accuracy_score, confusion_matrix, classification_report from sklearn.preprocessing import StandardScaler column names = ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigree', 'Age', 'Outcome']

df = pd.read_csv('pima-indians-diabetes.csv', names = column_names , header= None, skiprows = 9) #removing the unwanted values by changing to numeric values df = df[pd.to numeric(df['Pregnancies'], errors='coerce').notnull()] df.reset index(drop=True, inplace=True) print(df.head()) # Features (X) - All columns except the last one (Outcome) X = df.iloc[:, :-1]# Target (y) - The last column (Outcome) y = df.iloc[:, -1]# Check the first few rows of features and target print(X.head()) print(y.head()) X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42) scaler = StandardScaler() X train = scaler.fit transform(X train) X test = scaler.transform(X test) model = GaussianNB() model.fit(X_train, y_train) y_pred = model.predict(X test) accuracy = accuracy_score(y_test, y_pred) print(f"Accuracy: {accuracy*100:.4f}")

#social media naive bayes 3 4

import pandas as pd import numpy as np from sklearn.model_selection import train_test_split from sklearn.preprocessing import LabelEncoder, StandardScaler from sklearn.naive_bayes import GaussianNB from sklearn.metrics import accuracy_score, confusion_matrix, classification_report import seaborn as sns

```
import matplotlib.pyplot as plt
df = pd.read_csv('data/Social_Network_Ads.csv')
print(df.head())
df = df.drop(columns=['User ID'])
df['Gender'] = LabelEncoder().fit transform(df['Gender'])
X = df[['Gender', 'Age', 'EstimatedSalary']] # Features (Gender, Age, EstimatedSalary)
y = df['Purchased'] # Target variable (Purchased)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X train = scaler.fit transform(X_train)
X_test = scaler.transform(X_test)
model = GaussianNB()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy: {accuracy * 100:.2f}%')
# Confusion Matrix
conf matrix = confusion matrix(y test, y pred)
print("Confusion Matrix:")
print(conf matrix)
# Classification Report
print("Classification Report:")
print(classification_report(y_test, y_pred))
#diabetes regression 1
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.linear model import LinearRegression
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
df = pd.read_csv('data/diabetes.csv')
df = df.dropna()
X = df[['Glucose']] # Use 'Glucose' as the predictor feature
y = df['Outcome'] # 'Outcome' is the target variable
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
model = LinearRegression()
model.fit(X_train, y_train)
y pred = model.predict(X test)
print("Coefficient (Slope):", model.coef_)
print("Intercept:", model.intercept )
rss = np.sum((y test - y pred) ** 2)
print("Residual Sum of Squares (RSS):", rss)
r2_score = model.score(X_test, y_test)
```

Plot the regression line

rmse = np.sqrt(mse)

plt.scatter(X_test, y_test, color='blue', label='Actual')

print("Root Mean Squared Error (RMSE):", rmse) accuracy = r2_score * 100 # In percentage terms print("Accuracy (R2 score in percentage):", accuracy)

print("Coefficient of Determination (R2):", r2_score) mae = mean_absolute_error(y_test, y_pred) print("Mean Absolute Error (MAE):", mae) mse = mean_squared_error(y_test, y_pred) print("Mean Squared Error (MSE):", mse)

plt.plot(X_test, y_pred, color='red', linewidth=2, label='Regression Line')

plt.title('Simple Linear Regression')

plt.xlabel('Glucose')

plt.ylabel('Outcome')

plt.legend()

plt.show()

#sat vs gpa regression 2

import pandas as pd import numpy as np

```
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
df = pd.read csv('data/1.01. Simple linear regression.csv')
X = df[['SAT']] # Independent variable (SAT score)
y = df['GPA'] # Dependent variable (GPA)
X train, X test, y train, y test = train_test_split(X, y, test_size=0.2, random_state=42)
model = LinearRegression()
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print("Coefficient (Slope):", model.coef_)
print("Intercept:", model.intercept )
# Calculate Residual Sum of Squares (RSS)
rss = np.sum((y_test - y_pred) ** 2)
print("Residual Sum of Squares (RSS):", rss)
# Coefficient of Determination (R2)
r2 = model.score(X_test, y_test)
print("Coefficient of Determination (R2):", r2)
# Mean Absolute Error (MAE)
mae = mean absolute error(y test, y pred)
print("Mean Absolute Error (MAE):", mae)
# Mean Squared Error (MSE)
mse = mean_squared_error(y_test, y_pred)
print("Mean Squared Error (MSE):", mse)
# Root Mean Squared Error (RMSE)
rmse = np.sqrt(mse)
print("Root Mean Squared Error (RMSE):", rmse)
# Plot the regression line
plt.scatter(X, y, color='blue', label='Data Points') # Scatter plot of the data
plt.plot(X, model.predict(X), color='red', label='Regression Line') # Plot the regression line
plt.title('Linear Regression: SAT vs GPA')
plt.xlabel('SAT Score')
plt.ylabel('GPA')
plt.legend()
plt.show()
#advertising regression 345
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.linear model import LinearRegression
from sklearn.model selection import train_test_split
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
df = pd.read_csv('data/advertising.csv')
y = df['Sales']
def perform_regression(feature_name):
  # Define the independent variable (X)
  X = df[[feature name]] # We are using only one feature (TV, Radio, or Newspaper)
  # Split the data into training and testing sets (80-20 split)
  X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
  # Initialize the Linear Regression model
  model = LinearRegression()
  # Train the model on the training data
  model.fit(X_train, y_train)
  # Make predictions on the test data
  y_pred = model.predict(X_test)
  # Display the coefficients
  print(f"\n{feature_name} Model Coefficients:")
  print("Coefficient (Slope):", model.coef_)
  print("Intercept:", model.intercept_)
  # Model Evaluation Metrics
  print(f"\nModel Evaluation for {feature name}:")
  # Coefficient of Determination (R2)
  r2 = r2_score(y_test, y_pred)
```

```
print("Coefficient of Determination (R2):", r2)
  # Mean Absolute Error (MAE)
  mae = mean_absolute_error(y_test, y_pred)
  print("Mean Absolute Error (MAE):", mae)
  # Mean Squared Error (MSE)
  mse = mean_squared_error(y_test, y_pred)
  print("Mean Squared Error (MSE):", mse)
  # Root Mean Squared Error (RMSE)
  rmse = np.sqrt(mse)
  print("Root Mean Squared Error (RMSE):", rmse)
  # Plot the regression line
  plt.figure(figsize=(4, 4))
  plt.scatter(X, y, color='blue', label='Data Points') # Scatter plot of the data
  plt.plot(X, model.predict(X), color='red', label=f'Regression Line ({feature name})') # Plot the regression line
  plt.title(f'Linear Regression: {feature_name} vs Sales')
  plt.xlabel(feature_name)
  plt.ylabel('Sales')
  plt.legend()
  plt.show()
perform regression('TV')
perform regression('Radio')
perform regression('Newspaper')
#cities hierarchical clustering 1
import pandas as pd
import numpy as np
from sklearn.preprocessing import StandardScaler
from scipy.cluster.hierarchy import dendrogram, linkage
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.cluster import AgglomerativeClustering
from scipy.cluster.hierarchy import fcluster
df = pd.read csv('data/cities r2.csv')
# We are interested in the 'effective literacy rate total' column for clustering
df literacy = df[['effective_literacy_rate_total']]
print(df_literacy.isnull().sum())
scaler = StandardScaler()
df literacy scaled = scaler.fit transform(df literacy)
Z = linkage(df literacy scaled, method='ward')
# Step 4: Plot the Dendrogram to decide the number of clusters
plt.figure(figsize=(5, 5))
dendrogram(Z)
plt.title('Dendrogram for Hierarchical Clustering')
plt.xlabel('Cities')
plt.ylabel('Euclidean Distance')
plt.show()
# Fit the model with 3 clusters
hc = AgglomerativeClustering(n_clusters=3, metric='euclidean', linkage='ward')
# Add the cluster labels to the original dataframe
df['Cluster'] = hc.fit predict(df literacy scaled)
# Step 6: Visualizing the clusters
# Here we plot the cities based on their effective literacy rate and color them by clusters
sns.scatterplot(x=df['effective_literacy_rate_total'],
          y=np.zeros_like(df['effective_literacy_rate_total']),
          hue=df['Cluster'],
          palette='viridis',
          s=100, alpha=0.7)
plt.title('Hierarchical Clustering - Literacy Rate')
plt.xlabel('Effective Literacy Rate Total')
plt.ylabel('Cluster')
plt.show()
print(df[['name_of_city', 'effective_literacy_rate_total', 'Cluster']])
```

```
#hitters hierarchical clustering 2
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy import dendrogram, linkage
from sklearn.preprocessing import StandardScaler
from sklearn.cluster import AgglomerativeClustering
import seaborn as sns
df = pd.read csv('data/hitters.csv')
df cruns = df[['CRuns']]
scaler = StandardScaler()
df_cruns_scaled = scaler.fit_transform(df_cruns)
Z = linkage(df_cruns_scaled, method='ward')
#dendogram
plt.figure(figsize=(4,4))
dendrogram(Z)
plt.title('Dendrogram for Hierarchical Clustering (CRuns)')
plt.xlabel('Players')
plt.ylabel('Euclidean Distance')
plt.show()
hc = AgglomerativeClustering(n_clusters=3, metric='euclidean', linkage='ward')
# Assign cluster labels to the dataframe
df['Cluster'] = hc.fit_predict(df_cruns_scaled)
# Plot the 'CRuns' values and color them based on their cluster
plt.figure(figsize=(4,4))
sns.scatterplot(x=df['CRuns'],
          y=np.zeros like(df['CRuns']),
          hue=df['Cluster'],
          palette='viridis',
          s=50, alpha=0.7)
plt.title('Hierarchical Clustering - CRuns')
plt.xlabel('CRuns (Clustered)')
plt.ylabel('Cluster')
plt.show()
#cluster count
print(df[['AtBat', 'Hits', 'HmRun', 'CRuns', 'Cluster']])
#startups hierarchical 3
import pandas as pd
from sklearn.preprocessing import LabelEncoder, StandardScaler
from scipy.cluster.hierarchy import dendrogram, linkage, fcluster
import matplotlib.pyplot as plt
import seaborn as sns
df = pd.read_csv('data/50_Startups.csv')
abel_encoder = LabelEncoder()
df['STATE'] = label encoder.fit transform(df['STATE'])
# Select the numerical columns for scaling
numerical columns = ['RND', 'ADMIN', 'MKT', 'PROFIT']
scaler = StandardScaler()
# Scale the data
df scaled = df[numerical columns]
df scaled = scaler.fit transform(df scaled)
Z = linkage(df_scaled, method='ward')
# Step 4: Plot the Dendrogram to decide the number of clusters
plt.figure(figsize=(4,4))
dendrogram(Z)
plt.title('Dendrogram for Hierarchical Clustering')
plt.xlabel('Startups')
plt.ylabel('Euclidean Distance')
plt.show()
clusters = fcluster(Z, t=3, criterion='maxclust')
df['Cluster'] = clusters
print(df[['RND', 'ADMIN', 'MKT', 'PROFIT', 'Cluster']].head())
plt.figure(figsize=(5,5))
```

sns.scatterplot(x=df['RND'], y=df['PROFIT'], hue=df['Cluster'], palette='viridis', s=50, alpha=0.7)

```
plt.title('Hierarchical Clustering - RND vs PROFIT')
plt.xlabel('RND')
plt.ylabel('PROFIT')
plt.show()
#loan eligibility decision tree 1
import pandas as pd
from sklearn.model selection import train test split
from sklearn.preprocessing import LabelEncoder, StandardScaler
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import matplotlib.pyplot as plt
from sklearn import tree
df = pd.read_csv('data/madfhantr.csv')
print(df.isnull().sum())
df['LoanAmount'].fillna(df['LoanAmount'].median(), inplace=True)
df.dropna(subset=['Loan Status'], inplace=True)
df['Dependents'] = df['Dependents'].replace('3+', 3)
# Handle missing values in 'Dependents' column
df['Dependents'] = df['Dependents'].fillna(0) # Replace NaN with 0 or another reasonable value
# Convert 'Dependents' column to integer
df['Dependents'] = df['Dependents'].astype(int)
df['Loan_Amount_Term'].fillna(df['Loan_Amount_Term'].median(), inplace=True)
df.dropna(subset=['Loan_Amount_Term'], inplace=True)
df['Credit_History'].fillna(df['Credit_History'].median(), inplace=True)
df.dropna(subset=['Credit History'], inplace=True)
for column in ['Gender', 'Married', 'Education', 'Self Employed', 'Property_Area', 'Loan_Status']:
  df[column] = LabelEncoder().fit_transform(df[column])
X = df[['Gender', 'Married', 'Dependents', 'Education', 'Self Employed', 'ApplicantIncome',
     'CoapplicantIncome', 'LoanAmount', 'Loan Amount Term', 'Credit History', 'Property Area']]
y = df['Loan Status']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
dt_classifier = DecisionTreeClassifier(random_state=42)
dt_classifier.fit(X_train_scaled, y_train)
y pred = dt classifier.predict(X test scaled)
accuracy = accuracy_score(y_test, y_pred)
print("\nAccuracy Score of the Decision Tree Classifier:", accuracy)
# Confusion Matrix
print("\nConfusion Matrix:")
print(confusion matrix(y test, y pred))
# Classification Report
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
plt.figure(figsize=(9,9))
tree.plot_tree(dt_classifier, filled=True, feature_names=X.columns, class_names=['N', 'Y'], rounded=True, proportion=False,
precision=2)
plt.title("Decision Tree - Loan Eligibility")
plt.show()
#diabetes decision tree 2
import pandas as pd
from sklearn.model selection import train test split
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import matplotlib.pyplot as plt
from sklearn import tree
column names = [
  'Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
  'BMI', 'DiabetesPedigree', 'Age', 'Outcome']
df = pd.read csv('data/pima-indians-diabetes.csv', names = column names , header= None, skiprows = 9)
```

#removing the unwanted values by changing to numeric values df = df[pd.to_numeric(df['Pregnancies'], errors='coerce').notnull()]

```
df.reset index(drop=True, inplace=True)
print(df.head())
X = df.drop('Outcome', axis=1) # Features (all columns except 'Outcome')
y = df['Outcome'] # Target variable (Outcome)
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
dt_classifier = DecisionTreeClassifier(random_state=42)
dt classifier.fit(X train, y train)
y pred = dt classifier.predict(X test)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy: {accuracy * 100:.2f}%')
# Confusion Matrix
print("\nConfusion Matrix:")
print(confusion matrix(y test, y pred))
# Classification Report
print("\nClassification Report:")
print(classification_report(y_test, y_pred))
plt.figure(figsize=(12, 8))
tree.plot_tree(dt_classifier, filled=True, feature_names=X.columns, class_names=['No Diabetes', 'Diabetes'], rounded=True)
plt.title("Decision Tree - Diabetes Prediction")
plt.show()
#random sample test
random sample = X test.sample(1) # Pick a random sample from the test set
random sample prediction = dt classifier.predict(random sample) # Predict the outcome for this sample
# Print the random sample and its prediction
print("\nRandom Sample Prediction:")
print(random sample)
print(f"Prediction (1 = Diabetes, 0 = No Diabetes): {random_sample_prediction[0]}")
#cities r2 kmeans 1
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.cluster import KMeans
from sklearn.preprocessing import LabelEncoder, StandardScaler
import matplotlib.pyplot as plt
import seaborn as sns
df = pd.read_csv('data/cities_r2.csv')
print(df.isnull().sum()) # Check for missing values
# For simplicity, let's drop rows with missing values. Alternatively, we can fill with mean or mode
df = df.dropna()
abel encoder = LabelEncoder()
# Encode categorical columns
df['state name'] = label encoder.fit transform(df['state name'])
df['location'] = label encoder.fit transform(df['location'])
X = df[['total_graduates']] # We're focusing on total_graduates for clustering
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
kmeans = KMeans(n_clusters=3, random_state=42, n_init=10)
df['Cluster'] = kmeans.fit_predict(X_scaled)
plt.figure(figsize=(5,5))
sns.scatterplot(x=df.index, y='total graduates', hue='Cluster', palette='viridis', data=df, s=100)
plt.title('K-Means Clustering based on Total Graduates')
plt.xlabel('City Index')
plt.ylabel('Total Graduates')
plt.legend(title='Cluster')
plt.show()
print(df[['name_of_city', 'total_graduates', 'Cluster']].sort_values(by='Cluster'))
#social media advertise kmeans 2
# Step 1: Import necessary libraries
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.preprocessing import LabelEncoder, StandardScaler
```

import matplotlib.pyplot as plt import seaborn as sns

df = pd.read_csv('data/Social_Network_Ads.csv')

```
label encoder = LabelEncoder()
df['Gender'] = label_encoder.fit_transform(df['Gender'])
X = df[['Age', 'EstimatedSalary']] # We can include 'Age' as well if desired
scaler = StandardScaler()
X scaled = scaler.fit transform(X)
kmeans = KMeans(n_clusters=3, random_state=42, n_init=10)
df['Cluster'] = kmeans.fit predict(X scaled)
plt.figure(figsize=(5,5))
sns.scatterplot(x='Age', y='EstimatedSalary', hue='Cluster', palette='viridis', data=df, s=100)
plt.title('K-Means Clustering based on Age and Estimated Salary')
plt.xlabel('Age')
plt.ylabel('Estimated Salary')
plt.legend(title='Cluster')
plt.show()
print(df[['User ID', 'Gender', 'Age', 'EstimatedSalary', 'Cluster']].sort_values(by='Cluster'))
```

#k-means clustering same with 'effective_literacy_rate_total' instead of total graduates 3

with 8 initial centroids kmeans 4

```
import numpy as np
import matplotlib.pyplot as plt
points = np.array([
  [0.1, 0.6], #P1
  [0.15, 0.71], #P2
  [0.08, 0.9], #P3
  [0.16, 0.85], #P4
  [0.2, 0.3], #P5
  [0.25, 0.5], #P6
  [0.24, 0.1], #P7
  [0.3, 0.2] #P8
m1 = np.array([0.1, 0.6]) # P1
m2 = np.array([0.3, 0.2]) # P8
def euclidean distance(p1, p2):
  return np.sqrt(np.sum((p1 - p2) ** 2))
def assign_clusters(points, m1, m2):
  cluster_1 = []
  cluster_2 = []
  for point in points:
     d1 = euclidean_distance(point, m1)
     d2 = euclidean_distance(point, m2)
     if d1 < d2:
       cluster_1.append(point)
     else:
       cluster_2.append(point)
  return np.array(cluster_1), np.array(cluster_2)
def calculate_new_centroids(cluster_1, cluster_2):
  new_m1 = np.mean(cluster_1, axis=0) if len(cluster_1) > 0 else m1
  new_m2 = np.mean(cluster_2, axis=0) if len(cluster_2) > 0 else m2
  return new m1, new m2
def k means clustering(points, m1, m2):
  # Step 1: Assign points to the nearest centroid
  cluster_1, cluster_2 = assign_clusters(points, m1, m2)
  # Step 2: Calculate new centroids
  new m1, new m2 = calculate new centroids(cluster 1, cluster 2)
  return cluster_1, cluster_2, new_m1, new_m2
cluster_1, cluster_2, new_m1, new_m2 = k_means_clustering(points, m1, m2)
# Question 1: Which cluster does P6 belong to?
p6 = np.array([0.25, 0.5])
d1 = euclidean distance(p6, m1)
d2 = euclidean_distance(p6, m2)
p6_cluster = 1 if d1 < d2 else 2
# Question 2: What is the population of the cluster around m2?
population_cluster_2 = len(cluster_2)
# Question 3: What are the updated values of m1 and m2?
```

```
updated m1 = new m1
updated_m2 = new_m2
# Output the results
print(f"Cluster 1 (C1):\n{cluster_1}")
print(f"Cluster 2 (C2):\n{cluster 2}")
print(f"Updated m1: {updated_m1}")
print(f"Updated m2: {updated m2}")
# Answer to the questions:
print(f"\nAnswer to the questions:")
print(f"1. P6 belongs to Cluster {p6 cluster}")
print(f"2. The population of Cluster C2 (around m2) is: {population_cluster_2}")
print(f"3. The updated centroids are m1: {updated_m1} and m2: {updated_m2}")
plt.figure(figsize=(8, 6))
# Plot Cluster 1 in red
plt.scatter(cluster_1[:, 0], cluster_1[:, 1], color='red', label='Cluster 1 (C1)', s=100)
# Plot Cluster 2 in blue
plt.scatter(cluster_2[:, 0], cluster_2[:, 1], color='blue', label='Cluster 2 (C2)', s=100)
# Plot the centroids (m1 and m2)
plt.scatter(m1[0], m1[1], color='black', marker='X', s=200, label='Centroid m1 (C1)', linewidth=3)
plt.scatter(m2[0], m2[1], color='green', marker='X', s=200, label='Centroid m2 (C2)', linewidth=3)
# Plot the updated centroids
plt.scatter(updated m1[0], updated m1[1], color='orange', marker='X', s=200, label='Updated Centroid m1', linewidth=3)
plt.scatter(updated m2[0], updated m2[1], color='purple', marker='X', s=200, label='Updated Centroid m2', linewidth=3)
# Labeling the points and adding legend
plt.title("K-Means Clustering (1 Iteration)")
plt.xlabel("X-axis")
plt.ylabel("Y-axis")
plt.legend()
plt.grid(True)
plt.show()
# mobile prices svm 1
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.svm import SVC
from sklearn.metrics import accuracy score, classification report
import matplotlib.pyplot as plt
df = pd.read_csv('data/test.csv')
# Drop the 'id' column as it's just an identifier
#df = df.drop(columns=['id'])
# Creating a target variable based on 'ram' (you can modify this to suit your needs)
def classify_price_range(row):
  if row['ram'] < 1000:
     return 0 # Low
  elif row['ram'] < 2000:
     return 1 # Medium
  else:
     return 2 # High
# Apply the classification function to create the 'price range' column
df['price_range'] = df.apply(classify_price_range, axis=1)
X = df[['ram', 'battery_power']]
y = df['price range']
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)
svm = SVC(kernel='linear') # Using linear kernel for simplicity
svm.fit(X_train_scaled, y_train)
y_pred = svm.predict(X_test_scaled)
accuracy = accuracy_score(y_test, y_pred)
print(f'Accuracy of the SVM model: {accuracy * 100:.2f}%')
```

#universal bank SVM 2

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVC
from sklearn.linear_model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy score
import matplotlib.pyplot as plt
df = pd.read csv('data/UniversalBank.csv')
X = df.drop(columns=['ID', 'Personal Loan']) # Features
y = df['Personal Loan'] # Target (Personal Loan)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Step 5: Train and test the SVM model (using linear kernel for simplicity)
svm = SVC(kernel='linear')
svm.fit(X_train_scaled, y_train)
svm_pred = svm.predict(X_test_scaled)
svm_accuracy = accuracy_score(y_test, svm_pred)
log_reg = LogisticRegression(max_iter=1000)
log_reg.fit(X_train_scaled, y_train)
log_reg_pred = log_reg.predict(X_test_scaled)
log_reg_accuracy = accuracy_score(y_test, log_reg_pred)
rf = RandomForestClassifier(n estimators=100)
rf.fit(X_train, y_train)
rf_pred = rf.predict(X_test)
rf_accuracy = accuracy_score(y_test, rf_pred)
print(f'Accuracy of SVM: {svm_accuracy * 100:.2f}%')
print(f'Accuracy of Logistic Regression: {log_reg_accuracy * 100:.2f}%')
print(f'Accuracy of Random Forest: {rf_accuracy * 100:.2f}%')
# Step 9: Plot the comparison of model accuracies
models = ['SVM', 'Logistic Regression', 'Random Forest']
accuracies = [svm_accuracy, log_reg_accuracy, rf_accuracy]
plt.bar(models, accuracies, color=['blue', 'green', 'orange'])
plt.ylabel('Accuracy')
plt.title('Model Comparison')
plt.show()
#user behaviour svm 3
import pandas as pd
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
df = pd.read csv('data/user behavior dataset.csv')
le = LabelEncoder()
df['Device Model'] = le.fit transform(df['Device Model'])
df['Operating System'] = le.fit transform(df['Operating System'])
df['Gender'] = le.fit_transform(df['Gender'])
X = df.drop(columns=['User ID', 'User Behavior Class'])
y = df['User Behavior Class']
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
svm = SVC(kernel='linear')
svm.fit(X_train_scaled, y_train)
svm_pred = svm.predict(X_test_scaled)
```

```
svm_accuracy = accuracy_score(y_test, svm_pred)
print(f'Accuracy of SVM model: {svm_accuracy * 100:.2f}%')
plt.bar(['SVM'], [svm_accuracy], color='blue')
plt.ylabel('Accuracy')
plt.title('SVM Model Accuracy')
plt.show()

#user behavior svm 4
# Step 1: Import necessary libraries
import pandas as pd
```

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder, StandardScaler
from sklearn.svm import SVC
from sklearn.metrics import accuracy score, classification report
df = pd.read csv('data/bank transactions data 2.csv')
df.dropna(inplace=True)
label encoder = LabelEncoder()
# Columns that need encoding
categorical_columns = ['TransactionType', 'Location', 'DeviceID', 'MerchantID', 'Channel', 'CustomerOccupation']
for col in categorical columns:
  df[col] = label_encoder.fit_transform(df[col])
features = ['TransactionAmount', 'CustomerAge', 'TransactionDuration', 'LoginAttempts', 'AccountBalance']
X = df[features]
# Target variable: let's predict 'TransactionType' (Debit=0, Credit=1)
v = df['TransactionType']
#Step 3: Split the data 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)
# Step 4: Standardize the features (important for SVM)
scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_train)
X test scaled = scaler.transform(X test)
svm linear = SVC(kernel='linear')
svm linear.fit(X train scaled, y train)
y_pred_linear = svm_linear.predict(X_test_scaled)
accuracy_linear = accuracy_score(y_test, y_pred_linear)
print("SVM with Linear Kernel Accuracy: ", accuracy_linear)
print("Classification Report for Linear Kernel:")
print(classification_report(y_test, y_pred_linear))
svm poly = SVC(kernel='poly', degree=3)
svm poly.fit(X train scaled, y train)
# Make predictions for Polynomial Kernel
y_pred_poly = svm_poly.predict(X_test_scaled)
accuracy_poly = accuracy_score(y_test, y_pred_poly)
print("SVM with Polynomial Kernel Accuracy: ", accuracy_poly)
print("Classification Report for Polynomial Kernel:")
print(classification_report(y_test, y_pred_poly))
if accuracy linear > accuracy poly:
  print("The Linear Kernel model performs better.")
else:
  print("The Polynomial Kernel model performs better.")
```

#apriori market basket analysis 1

```
# Import necessary libraries
import pandas as pd
from mlxtend.frequent_patterns import apriori
from mlxtend.frequent_patterns import association_rules
# Step 1: Load the dataset
data = pd.read_csv("data/Order1.csv")
# Step 2: Data preprocessing
# Renaming columns for clarity (optional)
data.rename(columns={"Member_number": "TransactionID", "itemDescription": "Item"}, inplace=True)
# Step 3: Group transactions by TransactionID
```

```
# Each transaction will be a list of items
transactions = data.groupby("TransactionID")["Item"].apply(list)
# Step 4: Create a one-hot encoded DataFrame for Apriori
# Flatten the transactions to create a binary matrix
from mlxtend.preprocessing import TransactionEncoder
te = TransactionEncoder()
te data = te.fit(transactions).transform(transactions)
df_encoded = pd.DataFrame(te_data, columns=te.columns )
# Step 5: Apply the Apriori algorithm
frequent itemsets = apriori(df encoded, min support=0.09, use colnames=True)
print("Frequent Itemsets:")
print(frequent itemsets)
# Step 6: Generate association rules
from mlxtend.frequent_patterns import association_rules
# Generate association rules from frequent itemsets
rules = association_rules(frequent_itemsets,num_itemsets=2, metric="lift", min_threshold=1.0)
# Display association rules
print("Association Rules:")
print(rules)
plt.figure(figsize=(10, 6))
sns.barplot(x='support', y='antecedents', data=rules)
plt.title('Association Rules Support')
plt.xlabel('Support')
plt.ylabel('Antecedents')
plt.show()
# Visualization 2: Plot the lift of the association rules
plt.figure(figsize=(10, 6))
sns.barplot(x='lift', y='antecedents', data=rules)
plt.title('Association Rules Lift')
plt.xlabel('Lift')
plt.ylabel('Antecedents')
plt.show()
# Visualization 3: Scatter plot of support vs. lift for the association rules
plt.figure(figsize=(10, 6))
sns.scatterplot(x='support', y='lift', data=rules)
plt.title('Support vs. Lift for Association Rules')
plt.xlabel('Support')
plt.ylabel('Lift')
plt.show()
#apriori order 2
#data shaping
transactions = []
```

transaction = [str(data.iloc[i, j]) for j in range(data.shape[1]) if pd.notnull(data.iloc[i, j])]

for i in range(data.shape[0]): # Iterate through rows

transactions.append(transaction)

```
db.createCollection("CUSTOMER")
db.CUSTOMER.insertMany([
  Cust_No: 1001,
  First_Name: "John",
  Last_Name: "Doe",
  Address: "123 Main St",
  City: "Bangalore",
  State: "KARNATAKA",
  Pincode: 560001,
  B Date: new Date("1990-01-01"),
  Status: "Married"
 // Add more documents as needed
])
db.CUSTOMER.find({ State: "KARNATAKA" })
db.CUSTOMER.deleteOne({ Pincode: 576201 })
db.CUSTOMER.updateOne(
 { Cust_No: 1003 },
 { $set: { Address: "PICT, Trimurti chowk, Dhankawadi", Pincode: 411041 } }
db.CUSTOMER.aggregate([
{ $group: { _id: "$Status", count: { $sum: 1 } } }
])
db.CUSTOMER.find().sort({ City: 1 })
db.CUSTOMER.find({
 State: { $in: ["KARNATAKA", "KERALA"] },
 Status: "Married"
db.CUSTOMER.createIndex({ City: 1 })
db.CUSTOMER.getIndexes()
db.CUSTOMER.dropIndex("City_1")
db.CUSTOMER.aggregate([
 { $group: { _id: "$City", totalCustomers: { $sum: 1 } } }
#STUDENT MANAGEMENT SYSTEM
// 1. Create collection and insert sample data
db.createCollection("STUDENTS");
db.STUDENTS.insertMany([ /* Insert documents here */ ]);
// 2. Count total number of students
db.STUDENTS.countDocuments();
// 3. Display students by CGPA (seniority)
db.STUDENTS.find().sort({ CGPA: -1 });
// 4. List students from Baroda or Ahmedabad in CSE department
db.STUDENTS.find({
 address: { $in: ["Baroda", "Ahmedabad"] },
 department: "CSE"
```

// 5. List students born on the 20th of any month

#KARNATAKA

```
db.STUDENTS.find(
 { birthdate: { $regex: "-20T" } },
 { student_id: 1, student_name: 1, department: 1, skills: 1, _id: 0 }
// 6. Calculate age of each student
db.STUDENTS.aggregate([
  $addFields: {
   age: {
     $dateDiff: { startDate: "$birthdate", endDate: new Date(), unit: "year" }
]);
// 7. List students whose name starts with 'S' or 'M', in CSE with 'Typing' skill
db.STUDENTS.find({
 student_name: { $regex: "^[SM]", $options: "i" },
 department: "CSE",
 skills: "Typing"
}, { student_name: 1, _id: 0 });
// 8. Count students in IT department from Pune
db.STUDENTS.countDocuments({ department: "IT", address: "Pune" });
// 9. List students in ETC department between ages 18 and 20, sorted alphabetically
db.STUDENTS.aggregate([
  $addFields: {
   age: {
     $dateDiff: { startDate: "$birthdate", endDate: new Date(), unit: "year" }
 { $match: { age: { $gte: 18, $lte: 20 }, department: "ETC" } },
 { $sort: { student_name: 1 } }
]);
// 10. Create index on student_name, get indexes, and drop index
db.STUDENTS.createIndex({ student_name: 1 });
db.STUDENTS.getIndexes();
db.STUDENTS.dropIndex("student_name_1");
// 11. Aggregation to count students per department
db.STUDENTS.aggregate([
 { $group: { _id: "$department", totalStudents: { $sum: 1 } } }
]);
#ORDER MANAGEMENT SHORT
// 1. Retrieve all orders from the collection
db.ORDERS.find();
// 2. List customers in ascending order of their age (assuming customer info in another collection)
db.CUSTOMER.find().sort({ age: 1 });
// 3. Display total number of orders
db.ORDERS.countDocuments();
// 4. Display the mobile number of customers who have purchased product "Shoes" (if available in customer collection)
db.CUSTOMER.find({
 "orders.products.product_name": "Shoes"
}, { mobile_number: 1, _id: 0 });
// 5. Display total number of customers
db.CUSTOMER.countDocuments();
```

```
// 6. Display total products purchased in `order_id: 2`
db.ORDERS.aggregate([
 { $match: { order_id: 2 } },
 { $unwind: "$products" },
 { $group: { _id: "$order_id", totalProducts: { $sum: "$products.quantity" } } }
1);
// 7. Add another product with quantity 2 in order id: 3 for customer "ABC"
db.ORDERS.updateOne(
 { order_id: 3, customer_name: "ABC" },
 { $push: { products: { product_name: "New Product", quantity: 2, price: 1000 } } }
// 8. Create index on customer_name
db.ORDERS.createIndex({ customer_name: 1 });
// 9. Get the list of indexes on the collection
db.ORDERS.getIndexes();
// 10. Drop the index on customer name
db.ORDERS.dropIndex("customer name 1");
// 11. Aggregate function to calculate total order per customer
db.ORDERS.aggregate([
  $group: {
   _id: "$customer_name",
   totalOrders: { $sum: "$total amount" }
// OR: MapReduce function to calculate total order per customer
db.ORDERS.mapReduce(
 function() {
  emit(this.customer_name, this.total_amount);
 function(key, values) {
  return Array.sum(values);
 { out: "totalOrdersPerCustomer" }
#ORDER MANAGEMENT BIG
// 1. Display all orders
db.ORDERS.find();
// 2. List customers in ascending order of their names
db.ORDERS.find({}, { customer_name: 1, _id: 0 }).sort({ customer_name: 1 });
// 3. Display orders placed before April 2022
db.ORDERS.find({ order date: { $lt: new Date("2022-04-01") } });
// 4. Display customer names who spent more than 25000
db.ORDERS.find({ total_amount: { $gt: 25000 } }, { customer_name: 1, _id: 0 });
// 5. Display all orders that contain "PenDrive"
db.ORDERS.find({ "products.product_name": "PenDrive" });
// 6. Update order_date of an order placed by "ABC"
db.ORDERS.updateOne(
 { customer_name: "ABC" },
 { $set: { order_date: new Date("2023-01-01") } }
```

```
// 7. List orders with products having quantity < 10
db.ORDERS.find({ "products.quantity": { $lt: 10 } });
// 8. Display mobile number of customers with the highest buying total
db.ORDERS.aggregate([
 { $group: { _id: "$mob_no", totalSpent: { $sum: "$total_amount" } } },
 { $sort: { totalSpent: -1 } },
 { $limit: 1 }
]);
// 9. Create index on customer_name
db.ORDERS.createIndex({ customer_name: 1 });
// 10. Get indexes on the collection
db.ORDERS.getIndexes();
// 11. Drop index on customer_name
db.ORDERS.dropIndex("customer_name_1");
// 12. Aggregation to display total order per customer
db.ORDERS.aggregate([
  $group: {
    _id: "$customer_name",
   totalOrders: { $sum: "$total_amount" }
// 13. MapReduce to calculate total order per customer
db.ORDERS.mapReduce(
 function() {
  emit(this.customer_name, this.total_amount);
 function(key, values) {
  return Array.sum(values);
 { out: "totalOrdersPerCustomer" }
#EMPLOYEE MANAGEMENT SMALL
// 1. List employees who earn between 30,000 and 45,000
db.EMPLOYEE.find({ salary: { $gte: 30000, $lte: 45000 } });
// 2. List departments with at least four employees
db.EMPLOYEE.aggregate([
 { $group: { _id: "$department", employeeCount: { $sum: 1 } } },
 { $match: { employeeCount: { $gte: 4 } } },
 { $project: { department: "$_id", _id: 0, employeeCount: 1 } }
// 3. Count the number of employees in the "IT" department
db.EMPLOYEE.countDocuments({ department: "IT" });
// 4. Display name of employee who gets the maximum salary
db.EMPLOYEE.find().sort({ salary: -1 }).limit(1).project({ ename: 1, _id: 0 });
// 5. Display name of department with the maximum number of employees
db.EMPLOYEE.aggregate([
 { $group: { _id: "$department", employeeCount: { $sum: 1 } } },
 { $sort: { employeeCount: -1 } },
 { $project: { department: "$_id", _id: 0, employeeCount: 1 } }
]);
// 6. Update department name from 'IT' to "Information Technology"
```

```
db.EMPLOYEE.updateMany(
 { department: "IT" },
 { $set: { department: "Information Technology" } }
// 7. Create index on the 'department' field
db.EMPLOYEE.createIndex({ department: 1 });
// 8. Get indexes for the collection
db.EMPLOYEE.getIndexes();
// 9. Drop index on the 'department' field
db.EMPLOYEE.dropIndex("department 1");
// 10. Aggregate total number of employees per department
db.EMPLOYEE.aggregate([
 { $group: { _id: "$department", totalEmployees: { $sum: 1 } } }
// 11. MapReduce to calculate total number of employees per department
db.EMPLOYEE.mapReduce(
 function() {
  emit(this.department, 1); // Emit the department as the key and count 1 for each employee
 function(key, values) {
  return Array.sum(values); // Sum all the values to get the total number of employees per department
 { out: "employeeCountByDepartment" } // Output the results to a new collection
#EMPLOYEE MANAGEMENT BIG
// 1. List all employees
db.EMPLOYEE.find();
// 2. List employees from Baroda or Ahmedabad in CSE department
db.EMPLOYEE.find({
 city: { $in: ["Baroda", "Ahmedabad"] },
 department: "CSE"
// 3. List empid, ename, department and skills of employees whose join date is 20th of any month
db.EMPLOYEE.find(
 { $expr: { $eq: [{ $dayOfMonth: "$join_date" }, 20] } },
 { empid: 1, ename: 1, department: 1, skills: 1, _id: 0 }
// 4. Calculate total experience of employees considering today's date
db.EMPLOYEE.aggregate([
  $project: {
   ename: 1,
   totalExperience: {
     $cond: {
      if: { $eq: ["$leaving date", null] },
      then: { $dateDiff: { startDate: "$join_date", endDate: new Date(), unit: "year" } },
      else: { $dateDiff: { startDate: "$join_date", endDate: "$leaving_date", unit: "year" } }
]);
// 5. List employees whose name starts with 'S' or 'M', in FE department, with "Programming" skill
db.EMPLOYEE.find({
 ename: { $regex: "^[sSmM]" },
```

```
department: "FE",
 skills: "Programming"
// 6. Count the number of employees in ETC department in Pune
db.EMPLOYEE.countDocuments({ department: "ETC", city: "Pune" });
// 7. Calculate department-wise total salary and show the department with the highest total salary
db.EMPLOYEE.aggregate([
 { $group: { _id: "$department", totalSalary: { $sum: "$salary" } } },
 { $sort: { totalSalary: -1 } },
 { $limit: 1 }
]);
// 8. Create index, get index, and drop index on the department field
db.EMPLOYEE.createIndex({ department: 1 });
db.EMPLOYEE.getIndexes();
db.EMPLOYEE.dropIndex("department_1");
// 9. Using MapReduce to display total number of employees per department
db.EMPLOYEE.mapReduce(
 function() {
  emit(this.department, 1);
 function(key, values) {
  return Array.sum(values);
 { out: "employeeCountByDepartment" }
// 10. Using aggregation to display total number of employees per department
db.EMPLOYEE.aggregate([
 { $group: { _id: "$department", totalEmployees: { $sum: 1 } } }
]);
```