DSBD LAB

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import pandas as pd
import matplotlib.pyplot as plt
from sklearn.decomposition import PCA
from sklearn.preprocessing import StandardScaler
# Load dataset
url = "D:\DEVENDRA\VS Code\Assignment\TY\sem
6\dsbdcode\iris.data.csv"
cols = ['sepal_length', 'sepal_width', 'petal_length',
'petal_width', 'class']
df = pd.read_csv(url, header=None, names=cols)
# Standardize features and apply PCA
x = StandardScaler().fit_transform(df[cols[:-1]])
pca = PCA(n components=2).fit transform(x)
df pca = pd.DataFrame(pca, columns=['PC1', 'PC2'])
df_pca['class'] = df['class']
# Plot
colors = {'Iris-setosa': 'r', 'Iris-versicolor': 'g', 'Iris-
virginica': 'b'}
for label, color in colors.items():
    subset = df pca[df pca['class'] == label]
    plt.scatter(subset['PC1'], subset['PC2'], c=color, label=label,
s=50)
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.title('PCA of Iris Dataset')
plt.legend()
plt.grid()
plt.show()
```

```
# Import libraries
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 squared error, r2 score
# Load dataset from local CSV
df = pd.read_csv(r"D:\DEVENDRA\VS Code\Assignment\TY\sem
6\dsbdcode\HousingData.csv")
# Drop missing values
df = df.dropna()
# Prepare X and y using DataFrame and Series
X = pd.DataFrame(df.iloc[:, :-1], columns=df.columns[:-1])
y = pd.Series(df.iloc[:, -1])
# Split data into training and testing sets (80% train, 20% test)
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.2, random state=42)
# Create and train the linear regression model
model = LinearRegression()
model.fit(X_train, y_train)
# Predict on test set
y_pred = model.predict(X test)
# Evaluate the model
mse = mean_squared_error(y_test, y_pred)
r2 = r2_score(y_test, y_pred)
print(f"Mean Squared Error: {mse}")
print(f"R2 Score: {r2}")
# Plot actual vs predicted
plt.scatter(y test, y pred)
plt.xlabel("Actual Median House Value")
plt.ylabel("Predicted Median House Value")
plt.title("Actual vs Predicted Median House Value")
plt.show()
```

```
import pandas as pd, numpy as np
from sklearn.model selection import train test split
from sklearn.preprocessing import LabelEncoder
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import confusion matrix, accuracy score,
precision_score, recall_score, f1_score
# Load & preprocess
url = "D:\DEVENDRA\VS Code\Assignment\TY\sem
6\dsbdcode\iris.data.csv"
cols = ['sepal length', 'sepal width', 'petal length',
'petal_width', 'species']
df = pd.read csv(url, header=None, names=cols)
le = LabelEncoder()
df['species'] = le.fit transform(df['species']) # setosa=0,
versicolor=1, virginica=2
# Train-test split
X_train, X_test, y_train, y_test = train_test_split(df.iloc[:, :-1],
df['species'], test_size=0.3, random_state=42)
model = KNeighborsClassifier(n neighbors=3).fit(X train, y train)
y pred = model.predict(X_test)
# Treat class 0 (setosa) as positive, others as negative
target class = 0
y_test_binary = (y_test == target_class).astype(int)
y_pred_binary = (y_pred == target_class).astype(int)
# Compute confusion matrix
cm = confusion_matrix(y_test_binary, y_pred_binary)
TP = cm[1, 1]
FP = cm[0, 1]
TN = cm[0, 0]
FN = cm[1, 0]
# Print results
print("Confusion Matrix:\n", cm)
print("\nTrue Positives (TP):", TP)
print("False Positives (FP):", FP)
print("True Negatives (TN):", TN)
print("False Negatives (FN):", FN)
acc = accuracy score(y test binary, y pred binary)
print(f"\nAccuracy: {acc:.1f}")
print(f"Error Rate: {1 - acc:.1f}")
print("Precision:", round(precision score(y test binary,
y pred binary), 1))
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print("Recall:", round(recall_score(y_test_binary, y_pred_binary),
1))
print("F1-Score:", round(f1_score(y_test_binary, y_pred_binary), 1))
```

```
import numpy as np, matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.datasets import load iris
from sklearn.preprocessing import StandardScaler
X = StandardScaler().fit transform(load iris().data)
# Elbow method
inertia = [KMeans(n_clusters=k, random state=42,
n init=10).fit(X).inertia for k in range(1, 11)]
plt.plot(range(1, 11), inertia, marker='o'), plt.xlabel('k'),
plt.ylabel('Inertia'), plt.title('Elbow Method'), plt.show()
# KMeans with k=3
labels = KMeans(n clusters=3, random state=42,
n init=10).fit predict(X)
# Cluster visualization (first 2 features)
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis',
edgecolor='k')
plt.xlabel('Feature 1'), plt.ylabel('Feature 2'), plt.title('K-Means
Clustering (k=3)'), plt.show()
```