Name: Arham Sharif

Seat No.: EB21102022

Section: B

Subject: Data Warehouse & Data Mining

URL: https://www.kaggle.com/arhamsharif

DATA VISUALIZATION

EDA ON STUDENT PERFORMANCE DATASET

https://www.kaggle.com/code/arhamsharif/eda-on-student-performance-dataset

CODE

```
1. Import Libraries

import pandas as pd
import matplotlib.pyplot as plt

2. Load Dataset from Raw URL

url = "https://raw.qithubusercontent.com/selva86/datasets/master/College.csv"
df = pd.read_csv(url)
```

3. Basic Info

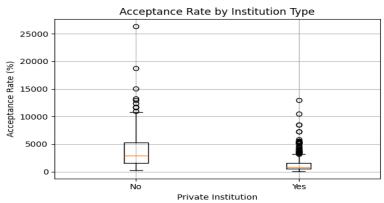
```
print("Shape:", df.shape)
print("Ghums:", df.columns:", df
```

4. Histogram: Outstate Tuition

```
| Dit.figure(figsire=(8, 4)) | 2 | pit.mix(dff'Outstate*], bins=30, colors*skyblue*, edgecolor=*black*) | 3 | pit.mix(dff'Outstate*], bins=30, colors*skyblue*, edgecolor=*black*) | 4 | pit.mix(dff'Outstate*], bins=30, colors*skyblue*, edgecolor=*black*) | 4 | pit.mix(df'Outstate*], bins=30, colors*skyblue*, edgecolor=*black*) | 5 | pit.mix(df'Outstate*), bins=30, colors*skyblue*, edgecolor=*black*) | 5 | pit.mix(df
```

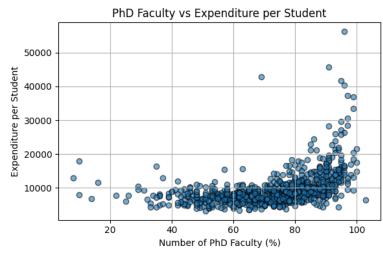
5. Boxplot: Accept Rate vs Private/Public

```
plt.figure(figsize=(6, 4))
    groups = df.groupby("Private")["Accept"]
    plt.boxplot([group for _, group in groups], tick_labels=["No", "Yes"])
    plt.title("Acceptance Rate by Institution Type")
    plt.xlabel("Private Institution")
    plt.ylabel("Acceptance Rate (%)")
    plt.grid(True)
    plt.tight_layout()
    plt.show()
```



6. Scatter Plot: Faculty vs Expenditures

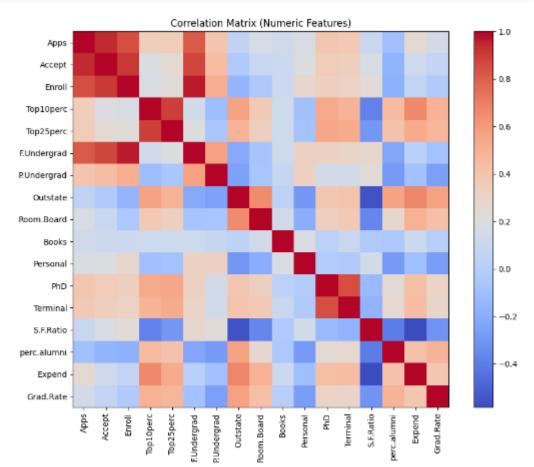




7. Correlation Heatmap with Matplotlib

```
num_df = df.select_dtypes(include="number")
corr = num_df.corr()

plt.figure(figsize=(10, 8))
plt.imshow(corr, cmap="coolwarm", interpolation="nearest")
plt.colorbar()
plt.title("Correlation Matrix (Numeric Features)")
plt.xticks(range(len(corr.columns)), corr.columns, rotation=90)
plt.yticks(range(len(corr.columns)), corr.columns)
plt.tight_layout()
plt.show()
```



REINFORCEMENT LEARNING

BASIC REINFORCEMENT LEARNING WITH CARTPOLE

https://www.kaggle.com/code/arhamsharif/basic-reinforcement-learning-with-cartpole

CODE

1. Import Libraries

```
import warnings

# Suppress the specific pkg_resources deprecation warning from pygame
warnings.filterwarnings(
    "ignore",
    message="pkg_resources is deprecated as an API*",
    category=UserWarning,
)

import numpy as np
np.bool8 = np.bool_ # Fix numpy bool8 attribute error
import gym
```

2. Create the CartPole environment with human rendering mode for visualization

```
env = gym.make("CartPole-v1", render_mode="human")

episodes = 10  # Number of episodes to run

max_steps = 280  # Max steps per episode

for episode in range(episodes):
    # Reset environment to initial state
    obs = env.reset()

# Handle new gym reset() returning (obs, info) tuple

if isinstance(obs, tuple):
    state, _ = obs

else:

state = obs
```

```
> 10
         total_reward = 0
         for step in range(max_steps):
  18
           # No need to call env.render() here, render_mode="human" handles it
  19
  28
            # Select an action randomly from the action space
            action = env.action_space.sample()
           # Take a step in the environment with the selected action
            result = env.step(action)
  25
            # Handle step() output differences between gym versions
            if len(result) == 5:
                next_state, reward, terminated, truncated, info = result
  38
                 done = terminated or truncated
            else:
  32
             next_state, reward, done, info = result
            state = next_state
            total_reward += reward
            # Break loop if episode finished
            if done:
             break
          print(f"Episode {episode + 1}: Total Reward = {total_reward}")
```

```
Episode 1: Total Reward = 42.0
Episode 2: Total Reward = 26.0
Episode 3: Total Reward = 15.0
Episode 4: Total Reward = 12.0
Episode 5: Total Reward = 39.0
Episode 6: Total Reward = 16.0
Episode 7: Total Reward = 23.0
Episode 8: Total Reward = 13.0
Episode 9: Total Reward = 12.0
Episode 10: Total Reward = 10.0
```

3. Close environment window

```
env.close()
```

CLASSIFICATION

IRIS DATASET CLASSIFICATION WITH MULTIPLE MODELS

https://www.kaggle.com/code/arhamsharif/iris-dataset-classification-with-multiple-models

CODE

~

1. Import Libraries

```
import pandas as pd
  import matplotlib.pyplot as plt
5 from matplotlib.colors import ListedColormap
5 from sklearn.preprocessing import LabelEncoder
o from sklearn.model_selection import train_test_split
7 from sklearn.pipeline import make_pipeline
8 from sklearn.preprocessing import StandardScaler
9 from sklearn.neighbors import KNeighborsClassifier
18 from sklearn.svm import SVC
11 from sklearn.gaussian_process import GaussianProcessClassifier
12 from sklearn.gaussian_process.kernels import RBF
13 from sklearn.tree import DecisionTreeClassifier
14 from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier
13 from sklearn.neural_network import MLPClassifier
10 from sklearn.naive_bayes import GaussianNB
  from sklearn.discriminant_analysis import QuadraticDiscriminantAnalysis
18 from sklearn.inspection import DecisionBoundaryDisplay
```

2. Import Data from online URL

```
url = "https://archive.ics.uci.edu/ml/machine-learning-databases/iris/iris.data"

# The iris data doesn't have headers, so add them manually

column_names = ['sepal_length', 'sepal_width', 'petal_length', 'petal_width', 'class']

df = pd.read_csv(url, header=None, names=column_names)
```

3. Preprocess / EDA / IDA

```
print(df.head())
print(df['class'].value_counts())

# Use only two features for visualization
X = df[['sepal_length', 'sepal_width']].values
y = df['class'].values

# Encode target labels
le = LabelEncoder()
y = le.fit_transform(y)
```

```
sepal_length sepal_width petal_length petal_width
  5.1 3.5 1.4 0.2 Iris-setosa
                       1.4
1.3
1.5
1.4
        4.9
                  3.0
                                        0.2 Iris-setosa
             3.0
3.2
3.1
3.6
       4.7
                                        0.2 Iris-setosa
        4.6
                                        0.2 Iris-setosa
        5.0
                                       0.2 Iris-setosa
Iris-setosa
Iris-versicolor
Iris-virginica
Name: count, dtype: int64
```

4. Train-Test Split (80% Train / 20% Test)

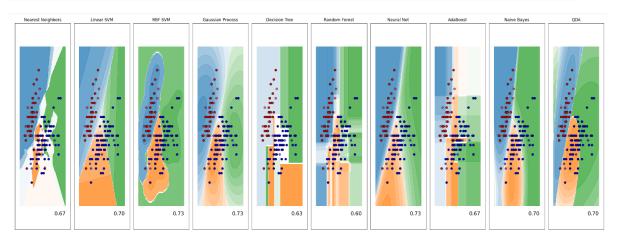
5. Execute / Feature Scaling inside pipeline

6. Define classifiers

```
1 names = [
      "Nearest Neighbors",
      "Linear SVM",
     "RBF SVM",
      "Gaussian Process",
      "Decision Tree",
      "Random Forest",
      "Neural Net",
      "AdaBoost",
      "Naive Bayes",
      "QDA",
12 ]
14 classifiers = [
     KNeighborsClassifier(3),
15
      SVC(kernel="linear", C=0.025, random_state=42),
     SVC(gamma=2, C=1, random_state=42),
      GaussianProcessClassifier(1.0 * RBF(1.0), random_state=42),
      DecisionTreeClassifier(max_depth=5, random_state=42),
      RandomForestClassifier(max_depth=5, n_estimators=10, max_features=1, random_state=42),
      MLPClassifier(alpha=1, max_iter=1000, random_state=42),
      AdaBoostClassifier(random_state=42),
      GaussianNB(),
       QuadraticDiscriminantAnalysis(),
```

7. Testing and 8. Predict + Visualize decision boundaries

```
figure = plt.figure(figsize=(27, 9))
cm_bright = ListedColormap(["#FF0000", "#0000FF"])
x_{min}, x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
y_{min}, y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
# Plot input data
for i, (name, clf) in enumerate(zip(names, classifiers), start=2):
   ax = plt.subplot(1, len(classifiers) + 1, i)
   clf_pipeline = make_pipeline(StandardScaler(), clf)
  clf_pipeline.fit(X_train, y_train)
   score = clf_pipeline.score(X_test, y_test)
   DecisionBoundaryDisplay.from_estimator(
      clf_pipeline, X, alpha=0.8, ax=ax, eps=0.5
   ax.scatter(X_train[:, 0], X_train[:, 1], c=y_train, cmap=cm_bright, edgecolors="k")
   ax.scatter(X_test[:, 0], X_test[:, 1], c=y_test, cmap=cm_bright, alpha=0.6, edgecolors="k")
   ax.set_xlim(x_min, x_max)
   ax.set_ylim(y_min, y_max)
   ax.set_xticks(())
   ax.set_yticks(())
   ax.set_title(name)
   ax.text(
      x_max - 0.5,
      y_min + 0.3
       f"{score:.2f}",
       horizontalalignment="right",
plt.tight_layout()
plt.show()
```



RECOGNIZING HANDWRITTEN DIGITS

https://www.kaggle.com/code/arhamsharif/recognizing-handwritten-digits-classification

CODE

1. Import Libraries

```
import matplotlib.pyplot as plt
import numpy as np
from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.pipeline import make_pipeline
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, accuracy_score
```

2. Load Digits Dataset

```
digits = datasets.load_digits()
X = digits.data
y = digits.target

print(f"Dataset shape: {X.shape}")
print(f"Number of classes: {len(np.unique(y))}")

Dataset shape: (1797, 64)
Number of classes: 10
```

3. Train-Test Split (80% train, 20% test)

```
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, random_state=42, stratify=y
)
```

4. Define classifier pipeline (example: KNN with scaling)

```
clf = make_pipeline(StandardScaler(), KNeighborsClassifier(n_neighbors=3))
```

5. Train classifier

6. Test classifier

?Documentation for KNeighborsClassifier

KNeighborsClassifier

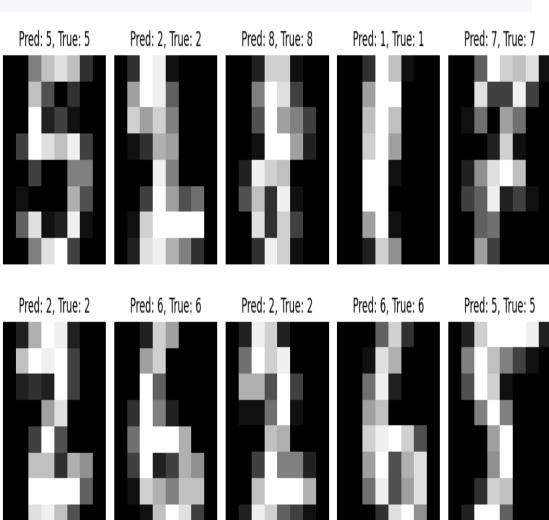
▶ Parameters

```
y_pred = clf.predict(X_test)
print(classification_report(y_test, y_pred))
print(f"Accuracy: {accuracy_score(y_test, y_pred):.4f}")
          precision recall f1-score support
                                  36
        А
            1.00 1.00 1.00
                                  36
            0.92 0.97 0.95
        1
                                  35
        2
            0.90 1.00 0.95
        3
            1.00 0.97 0.99
                                  37
        4
            0.97 0.94 0.96
                                  36
        5
            1.00 1.00 1.00
                                  37
        6
            0.97 1.00 0.99
                                  36
            0.95 0.97 0.96
        7
                                  36
        8
          0.97 0.89 0.93
                                  35
        Q
            1.00 0.92 0.96
                                  36
   accuracy
                           0.97
                                 360
  macro avg
            0.97
                 0.97 0.97
                                 360
 weighted avg
            0.97
                  0.97 0.97
                                  360
```

Accuracy: 0.9667

7. Visualize some predictions

```
fig, axes = plt.subplots(2, 5, figsize=(10, 5))
for ax, image, pred, true in zip(axes.flatten(), X_test, y_pred, y_test):
    ax.imshow(image.reshape(8, 8), cmap='gray')
    ax.set_title(f"Pred: {pred}, True: {true}")
    ax.axis('off')
plt.tight_layout()
plt.show()
```



REGRESSION

BAYESIAN REGRESSION COMPARISON (DIABETES)

https://www.kaggle.com/code/arhamsharif/bayesian-regression-comparison-diabetes

CODE

1. Import Libraries

```
import matplotlib.pyplot as plt
from sklearn.linear_model import BayesianRidge, ARDRegression, LinearRegression
from sklearn.datasets import load_diabetes
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import mean_squared_error, r2_score
```

2. Import Data (Diabetes Dataset)

```
diabetes = load_diabetes()

X = diabetes.data
y = diabetes.target

print(f"Dataset shape: {X.shape}")
print("Feature names:", diabetes.feature_names)

Dataset shape: (442, 10)
Feature names: ['age', 'sex', 'bmi', 'bp', 's1', 's2', 's3', 's4', 's5', 's6']
```

3. Preprocess / Feature Scaling (Inside Pipelines or Manual)

```
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
```

4. Train-Test Split

```
X_train, X_test, y_train, y_test = train_test_split(
    X_scaled, y, test_size=0.2, random_state=42
)
```

5. Define Regressors

```
ols = LinearRegression()
bayesian_ridge = BayesianRidge()
ard = ARDRegression()

regressors = {
    "Linear Regression (OLS)": ols,
    "Bayesian Ridge Regression": bayesian_ridge,
    "ARD Regression": ard,
}
```

6. Fit Models

```
for name, reg in regressors.items():
    reg.fit(X_train, y_train)
    print(f"{name} fitted successfully.")

Linear Regression (OLS) fitted successfully.
Bayesian Ridge Regression fitted successfully.
ARD Regression fitted successfully.
```

7. Test Models and Print Scores

```
for name, reg in regressors.items():
    y_pred = reg.predict(X_test)
    print(f"\n{name}")
    print(f"R^2 Score: {r2_score(y_test, y_pred):.4f}")
    print(f"Mean Squared Error: {mean_squared_error(y_test, y_pred):.4f}")

Linear Regression (OLS)
```

R^2 Score: 0.4526
Mean Squared Error: 2900.1936
Bayesian Ridge Regression
R^2 Score: 0.4580
Mean Squared Error: 2871.7621

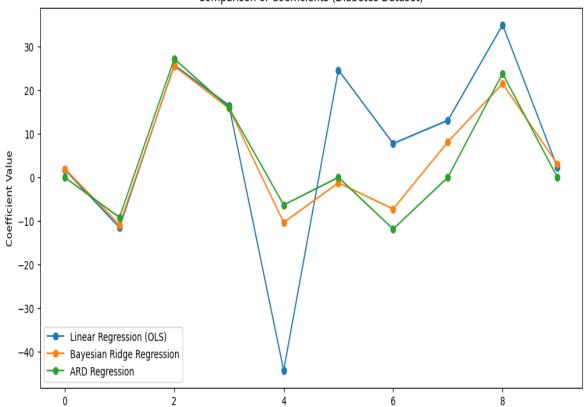
ARD Regression
R^2 Score: 0.4670
Mean Squared Error: 2823.7391

8. Visualize Coefficients

```
plt.figure(figsize=(12, 6))
for name, reg in regressors.items():
    plt.plot(reg.coef_, marker='o', label=name)

plt.title("Comparison of Coefficients (Diabetes Dataset)")
plt.xlabel("Feature Index")
plt.ylabel("Coefficient Value")
plt.legend()
plt.show()
```

Comparison of Coefficients (Diabetes Dataset)



ELASTICNET WITH AUTO PRECOMPUTED GRAM MATRIX

https://www.kaggle.com/code/arhamsharif/elasticnet-with-auto-precomputed-gram-matrix

CODE

1. Import Libraries

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import fetch_california_housing
from sklearn.linear_model import ElasticNet
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import mean_squared_error
from sklearn.model_selection import train_test_split
import time
```

2. Load Dataset

```
housing = fetch_california_housing()

X = housing.data

y = housing.target

print(f"Dataset shape: {X.shape}")

print("Feature names:", housing.feature_names)

Dataset shape: (20640, 8)
Feature names: ['MedInc', 'HouseAge', 'AveRooms', 'AveBedrms', 'Population', 'AveOccup', 'Latitude', 'Longitude']
```

3. Preprocess / Feature Scaling

```
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
```

4. Train-Test Split

5. Sample Weights (simulate weights based on distance from median target)

```
sample_weight = np.abs(y_train - np.median(y_train))
sample_weight /= sample_weight.max()
```

6. Fit ElasticNet Normally (Without Precomputed Gram)

```
start = time.time()
enet = ElastioNet(alpha=0.1, l1_ratio=0.5, fit_intercept=False, max_iter=10000)
enet.fit(X_train, y_train, sample_weight=sample_weight)
elapsed_normal = time.time() - start
print(f"ElasticNet fit without Gram took: {elapsed_normal:.4f} seconds")
```

ElasticNet fit without Gram took: 0.0535 seconds

7. Fit ElasticNet with Precomputed Gram (Automatically Handled by ElasticNet)

```
start = time.time()
enet_gram = ElasticNet(alpha=0.1, l1_ratio=0.5, fit_intercept=False, max_iter=18880, precompute=True)
enet_gram.fit(X_train, y_train, sample_weight=sample_weight)
elapsed_gram = time.time() - start
print(f"ElasticNet fit with Gram (auto-precomputed) took: {elapsed_gram:.4f} seconds")
```

ElasticNet fit with Gram (auto-precomputed) took: 0.0049 seconds

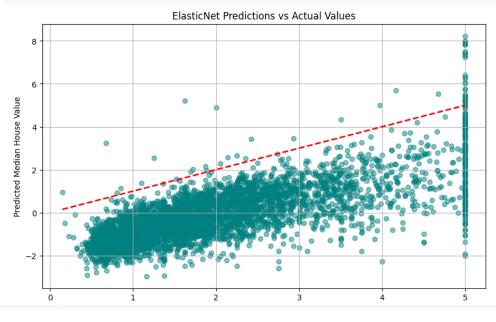
8. Predict & Evaluate

```
y_pred = enet.predict(X_test)
mse = mean_squared_error(y_test, y_pred)
print(f"Hean Squared Error (ElasticNet without Gram): {mse:.4f}")
```

Mean Squared Error (ElasticNet without Gram): 5.1821

9. Visualize Coefficients

```
> plt.figure(figsize=(18, 6))
plt.scatter(y_test, y_pred, alpha=8.5, color='teal')
plt.scatter(y_test, y_pred, alpha=8.5, color='teal')
plt.file("Last.min(), y_test.max()], [y_test.max()], 'r--', lw=2)
plt.file("ElasticMet Predictions vs Actual Values")
plt.xlabet("Predicted Median House Value")
plt.ylabet("Predicted Median House Value")
plt.spid(True)
plt.show()
```



CLUSTERING

AGGLOMERATIVE CLUSTERING ON WINE DATASET

https://www.kaggle.com/code/arhamsharif/agglomerative-clustering-on-wine-dataset

CODE

1. Import Libraries

```
i import matglotlib.pyplot as plt
from sklearn.datasets import load_mine
from sklearn.cluster import AgglomerativeClustering
from sklearn.preprocessing import StandardScaler
```

2. Load Wine Dataset

```
wine = load_wine()
X = wine.data
y = wine.data
y = wine.target
print("Dataset shape:", X.shape)
print("Flature names:", wine.feature_names)
print("Target classes:", wine.target_names)

Dataset shape: (178, 13)
Feature names: ('alcohol', 'malic_acid', 'ash', 'alcalinity_of_ash', 'magnesium', 'total_phenols', 'flavanoids', 'nonflavanoid_phenols', 'proanthocyanins', 'color_intensity', 'hue', 'od280/od315_of_diluted_wines', 'proline']
Target classes: ['class.0' 'class_1' 'class_2']
```

3. Feature Scaling (Clustering works better with scaled data)

```
scaler = StandardScaler()

X_scaled = scaler.fit_transform(X)
```

4. Apply Agglomerative Clustering with Different Linkage Methods

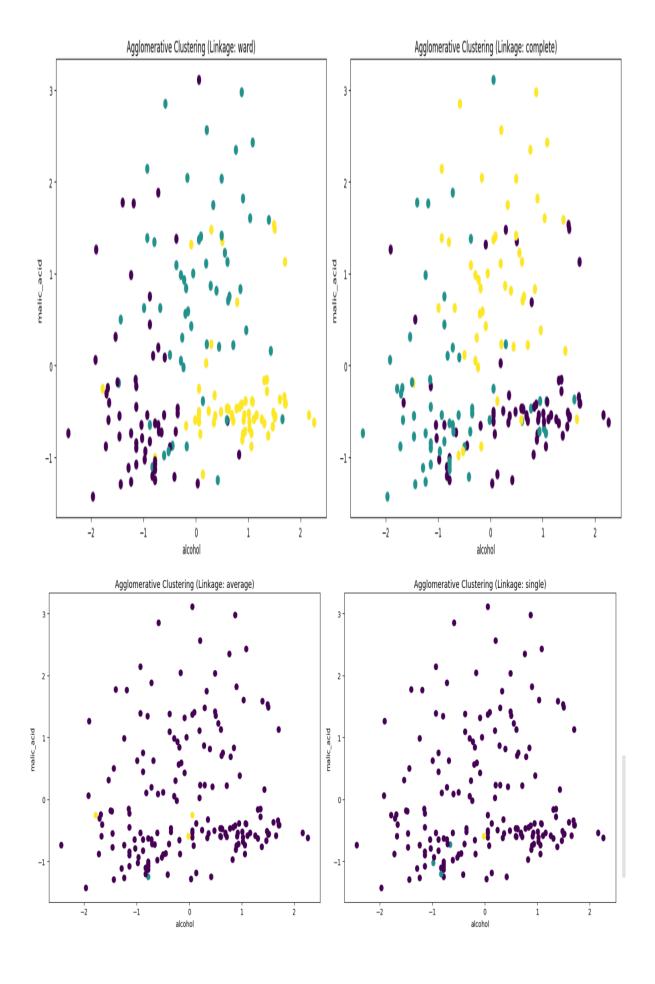
```
linkages = ['ward', 'complete', 'average', 'single']

plt.figure(figsize=(16, 12))
for i, linkage in enumerate(linkages, 1):
    # 'ward' works only with Euclidean distance
    if linkage == 'ward':
        model = AgglomerativeClustering(n_clusters=3, linkage=linkage)
    else:
        model = AgglomerativeClustering(n_clusters=3, linkage=linkage, metric='euclidean')

labels = model.fit_predict(X_scaled)

# Plot clusters using 2 selected features for visualization
    plt.subplot(2, 2, i)
    plt.scatter(X_scaled[:, 0], X_scaled[:, 1], c=labels, cmap='viridis', s=50)
    plt.title(f"Agglomerative Clustering (Linkage: {linkage})")
    plt.xlabel(wine.feature_names[0])
    plt.ylabel(wine.feature_names[1])

plt.tight_layout()
    plt.show()
```



AGGLOMERATIVE CLUSTERING WITH & WITHOUT STRUCTURE

https://www.kaggle.com/code/arhamsharif/agglomerative-clustering-with-without-structure

CODE

1. Import Libraries

```
import matplotlib.pyplot as plt
from sklearn.datasets import make_moons, make_blobs
from sklearn.cluster import AgglomerativeClustering
```

2. Generate Structured Data (Moons)

```
X_structured, _ = make_moons(n_samples=300, noise=0.05, random_state=42)
```

3. Generate Unstructured Data (Random Blobs)

```
X_unstructured, _ = make_blobs(n_samples=300, centers=3, random_state=42)
```

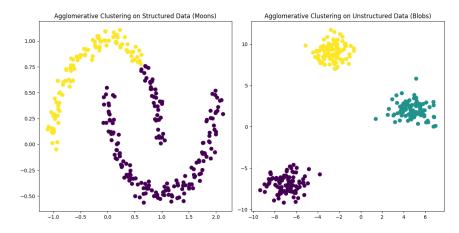
4. Apply Agglomerative Clustering

```
clustering_structured = AgglomerativeClustering(n_clusters=2)
labels_structured = clustering_structured.fit_predict(X_structured)

clustering_unstructured = AgglomerativeClustering(n_clusters=3)
labels_unstructured = clustering_unstructured.fit_predict(X_unstructured)
```

5. Visualize Results

```
plt.figure(figsize=(12, 6))
# Structured Data (Moons)
plt.subplot(1, 2, 1)
plt.scatter(X_structured[:, 0], X_structured[:, 1], c=labels_structured, cmap='viridis', s=58)
plt.title("Agglomerative Clustering on Structured Data (Moons)")
# Unstructured Data (Slobs)
plt.subplot(1, 2, 2)
plt.scatter(X_unstructured[:, 0], X_unstructured[:, 1], c=labels_unstructured, cmap='viridis', s=58)
plt.title("Agglomerative Clustering on Unstructured Data (Slobs)")
plt.tight_layout()
plt.show()
```



DIMENSIONALITY REDUCTION

KERNEL PCA WITH RBF KERNEL ON NONLINEAR CIRCLES

https://www.kaggle.com/code/arhamsharif/kernel-pca-with-rbf-kernel-on-nonlinear-circles

CODE

```
1. Import Libraries

1. Import Libraries

1. import matplotlib.pyplot as plt
2. from sklearn.detasets import make_circles
3. from sklearn.decomposition import KernelPCA
4. from sklearn.preprocessing import StandardScaler

2. Generate Nonlinear Data (Nested Circles)

1. X, y = make_circles(n_samples=600, factor=0.3, noise=0.05, random_state=62)

3. Standardize Features

1. scaler = StandardScaler()
2. X_scaled = scaler.fit_transform(X)

4. Apply Kernel PCA

1. kpca = KernelPCA(n_components=2, kernel='rbf', gamma=15)
2. X_kpca = kpca.fit_transform(X_scaled)
```

5. Plot Original Data

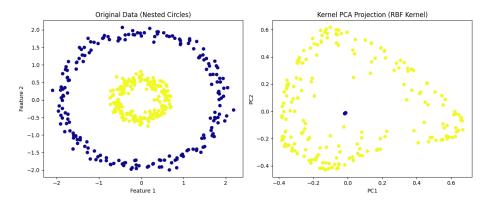
```
plt.figure(figsize=(12, 5))

plt.subplot(1, 2, 1)
plt.scatter(X_scaled[:, 0], X_scaled[:, 1], c=y, cmap='plasma', s=30)
plt.fitie("Original Data (Nested Circles)")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")

# 6. Plot Transformed Data
plt.subplot(1, 2, 2)
plt.scatter(X_kpoa[:, 0], X_kpca[:, 1], c=y, cmap='plasma', s=36)
plt.fitie("Kernel PCA Projection (RBF Kernel)")
plt.xlabel("PC1")

plt.ylabel("PC2")

plt.tight_layout()
plt.show()
```



PCA ON THE IRIS DATASET

https://www.kaggle.com/code/arhamsharif/pca-on-the-iris-dataset

CODE

1. Import Libraries import matplotlib.pyplot as plt from sklearn import datasets from sklearn.decomposition import PCA from sklearn.preprocessing import StandardScaler

2. Load Iris Dataset

```
i iris = datasets.load_iris()
2  X = iris.data
3  y = iris.target
4  target_names = iris.target_names
```

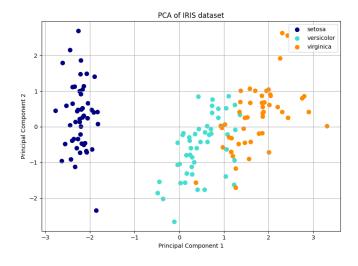
3. Standardize the Data

```
X_scaled = StandardScaler().fit_transform(X)
```

4. Apply PCA (reduce to 2 components for visualization)

```
pca = PCA(n_components=2)
z    X_pca = pca.fit_transform(X_scaled)
```

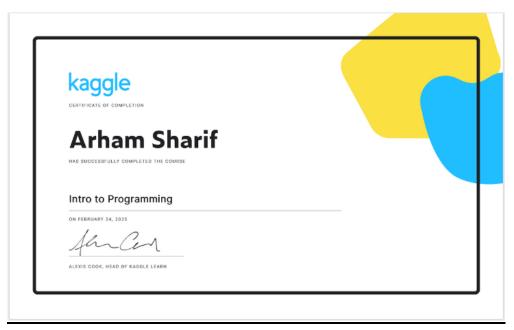
5. Plot Results



CERTIFICATES

INTRODUCTION TO PROGRAMMING

https://www.kaggle.com/learn/certification/arhamsharif/intro-to-programming



PYTHON

https://www.kaggle.com/learn/certification/arhamsharif/python



DATA VISUALIZATION

https://www.kaggle.com/learn/certification/arhamsharif/data-visualization



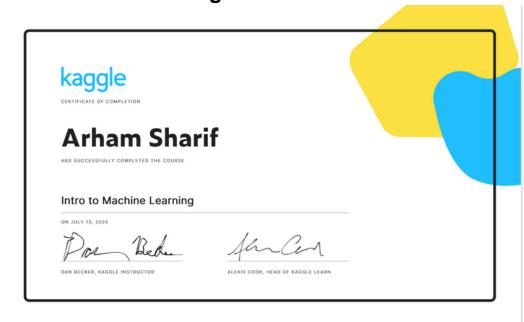
PANDAS

https://www.kaggle.com/learn/certification/arhamsharif/pandas



INTRO TO MACHINE LEARNING

https://www.kaggle.com/learn/certification/arhamsharif/intro-to-machine-learning



INTRO TO GAME AI AND REINFORCEMENT LEARNING

https://www.kaggle.com/learn/certification/arhamsharif/intro-to-game-ai-and-reinforcement-learning

