Problem 9

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

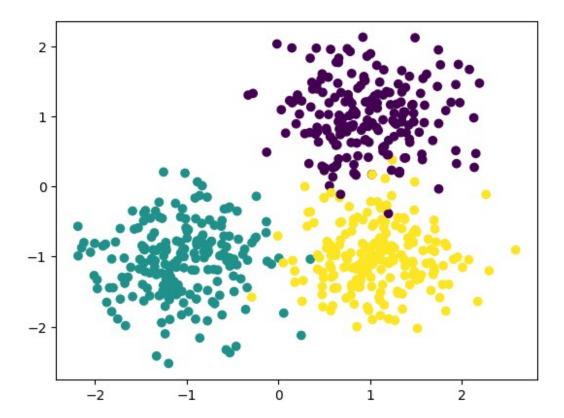
import seaborn as sns
import scipy.cluster.hierarchy as shc
from sklearn import metrics
from sklearn.datasets import make_blobs, make_circles, make_moons
from sklearn.cluster import DBSCAN
from sklearn.preprocessing import StandardScaler

seed = 0
```

Generate a dummy toy dataset with varying densities and shapes. Set the eps (Epsilon) andmin samples (MinPts) parameters, and then fit DBSCAN to the generated dataset.

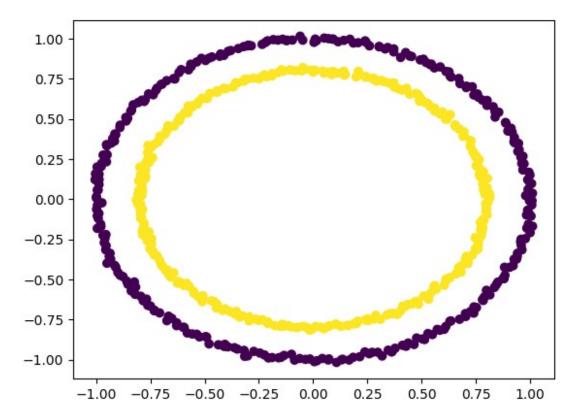
```
# blob data

centers = [[1, 1], [-1, -1], [1, -1]]
blob_X, blob_true_labels = make_blobs(n_samples = 600, centers = centers, cluster_std= 0.5, random_state = seed)
scaled_blob_X = StandardScaler().fit_transform(blob_X)
plt.scatter(blob_X[:, 0], blob_X[:, 1], c = blob_true_labels)
<matplotlib.collections.PathCollection at 0x7936dc1ae2f0>
```

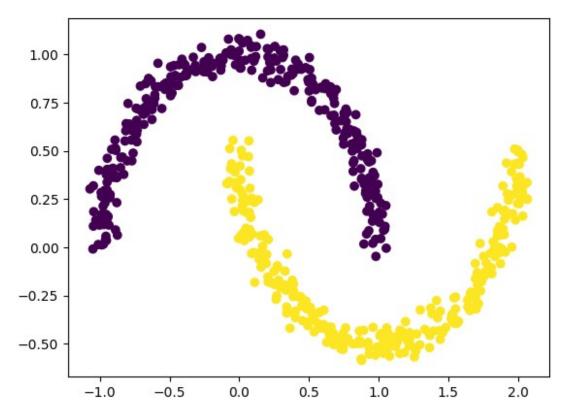


Concentric circles data

```
circle_X, circle_true_labels = make_circles(n_samples = 600, noise =
.01, random_state = seed)
scaled_circle_X = StandardScaler().fit_transform(circle_X)
plt.scatter(circle_X[:, 0], circle_X[:, 1], c = circle_true_labels)
<matplotlib.collections.PathCollection at 0x7936dbbf5600>
```



```
# Moon shaped data
moon_X, moon_true_labels = make_moons(n_samples = 600, noise = 0.05,
random_state = seed)
scaled_moon_X = StandardScaler().fit_transform(moon_X)
plt.scatter(moon_X[:, 0], moon_X[:, 1], c = moon_true_labels)
<matplotlib.collections.PathCollection at 0x7936dbc755a0>
```



```
# Helper function to plot

def plot_clusters(data, true_labels = None, cluster_labels = None,
title_true = 'True Cluster', title_cluster = 'DBSCAN clustering'):
    fig, (ax1, ax2) = plt.subplots(1, 2, figsize = (14, 6))
    ax1.scatter(data[:, 0], data[:, 1], c = true_labels)
    ax1.set_title(title_true)

if cluster_labels is not None:
    ax2.scatter(data[:, 0], data[:, 1], c = cluster_labels)
    ax2.set_title(title_cluster)

plt.show()
```

Objectives

- 1. Experiment with each combination of eps and min samples (consider at least 3 values of each) for these parameters. Report the values of the performance metrics to evaluate DBSCAN's sensitivity to parameter choices.
- 2. Visualize the clustering results using a scatter plot, where each cluster is assigned a different color. Additionally, use a different marker shape for noise points.
- 3. Calculate and report the following performance metrics: Silhouette Score, Adjusted Rand Index, Adjusted Mutual Information.

For Blob data

```
eps values blob = [0.2, 0.3, 0.5, 0.8, 1.3]
min samples values blob = [2, 3, 5, 8, 13]
for i, eps in enumerate(eps values blob):
  for j, min_samples in enumerate(min samples values blob):
   # Fit DBSCAN with current parameter combination
   dbscan blob = DBSCAN(eps=eps, min_samples=min_samples)
   dbscan blob.fit predict(scaled blob X)
   # Calculate performance metrics
   X, pred labels, true labels = scaled blob X, dbscan blob.labels ,
blob true labels
   if len(set(pred labels)) > 1:
     silhouette = metrics.silhouette score(X, pred labels)
     ari = metrics.adjusted rand score(true labels, pred labels)
     ami = metrics.adjusted mutual info score(true labels,
pred labels)
     print(f'For eps = {eps} and minimum samples = {min_samples}')
     print("Silhouette score: %0.3f " % silhouette)
     print('Adjusted Rand Index: %0.3f'% ari)
     print('Adjusted mutual information: %0.3f' % ami)
     plot clusters(scaled blob X, true labels, pred labels)
     print('\n')
Output hidden; open in https://colab.research.google.com to view.
```

Based on the scoring for different eps and min_samples we found that the eps = 0.3 and min_samples = 13 is the best combination for blob dataset.

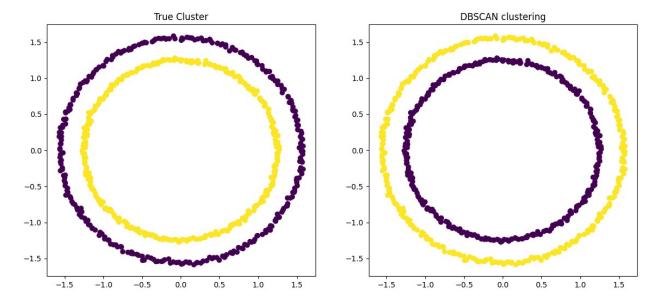
For Circle data

```
eps_values_circle = [0.2, 0.3, 0.5, 0.7]
min_samples_values_circle = [5, 10, 15, 20, 25]

for i, eps in enumerate(eps_values_circle):
    for j, min_samples in enumerate(min_samples_values_circle):
        # Fit DBSCAN with current parameter combination
        dbscan_circle = DBSCAN(eps=eps, min_samples=min_samples)
        dbscan_circle.fit_predict(scaled_circle_X)

# Calculate performance metrics
        X, pred_labels, true_labels = scaled_circle_X,

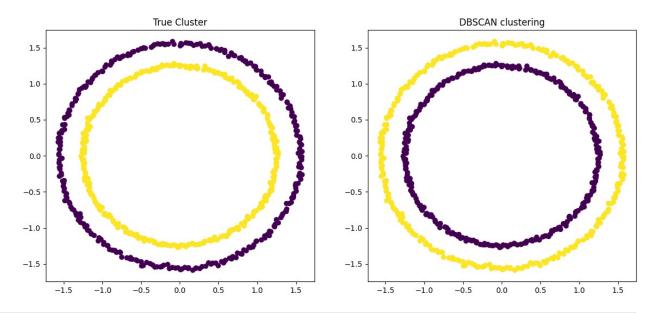
dbscan_circle.labels_, circle_true_labels
        if len(set(pred_labels)) > 1:
            silhouette = metrics.silhouette_score(X, pred_labels)
            ari = metrics.adjusted_rand_score(true_labels, pred_labels)
            ami = metrics.adjusted_mutual_info_score(true_labels,
            pred_labels)
```



For eps = 0.2 and minimum samples = 10

Silhouette score: 0.019 Adjusted Rand Index: 1.000

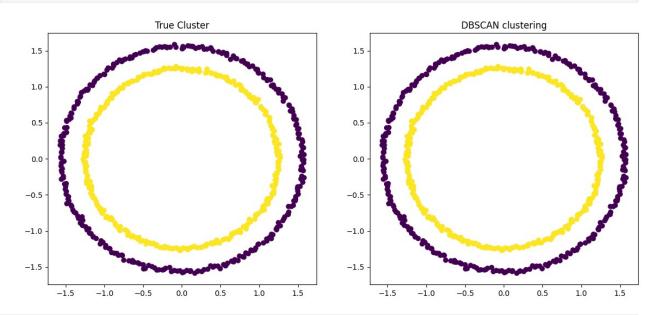
Adjusted mutual information: 1.000



For eps = 0.2 and minimum samples = 15

Silhouette score: 0.019 Adjusted Rand Index: 1.000

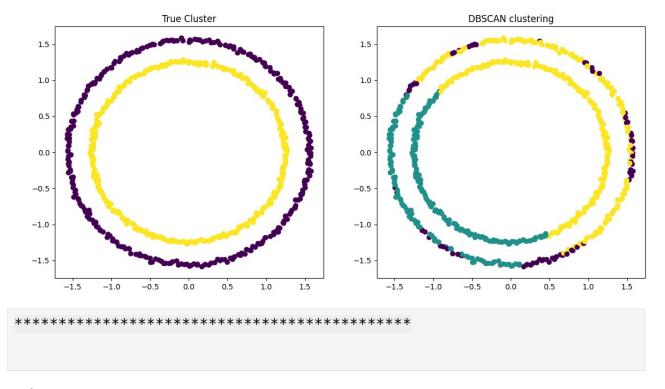
Adjusted mutual information: 1.000



For eps = 0.3 and minimum samples = 25

Silhouette score: 0.230

Adjusted Rand Index: 0.043
Adjusted mutual information: 0.110



Observations

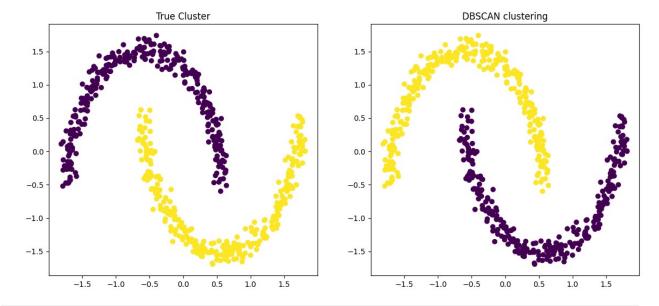
- 1. The scores are very bad for circular data.
- Intrestingly, for eps = 0.2, any number of min_samples is generating a ari and ami score =
 1, which implies that both the true cluster and predicted one agrees upon the placing of
 the points but the silhouette score is very bad.
- 3. We can conclude that DBSCAN clustering the data in proper number of clusters but it is not assigning correct datapoints to right cluster.

For moon data

```
eps_values_moon = [0.3, 0.5, 0.7]
min_samples_values_moon = [5, 10, 15, 20]

for i, eps in enumerate(eps_values_moon):
    for j, min_samples in enumerate(min_samples_values_moon):
        # Fit DBSCAN with current parameter combination
        dbscan_moon = DBSCAN(eps=eps, min_samples=min_samples)
        dbscan_moon.fit_predict(scaled_moon_X)

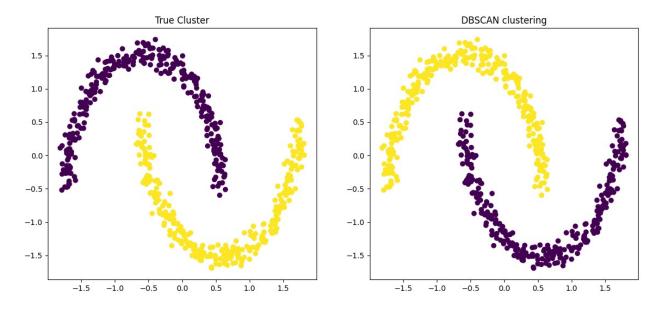
# Calculate performance metrics
        X, pred_labels, true_labels = scaled_moon_X, dbscan_moon.labels_,
moon_true_labels
    if len(set(pred_labels)) > 1:
        silhouette = metrics.silhouette_score(X, pred_labels)
```



For eps = 0.3 and minimum samples = 10

Silhouette score: 0.389 Adjusted Rand Index: 1.000

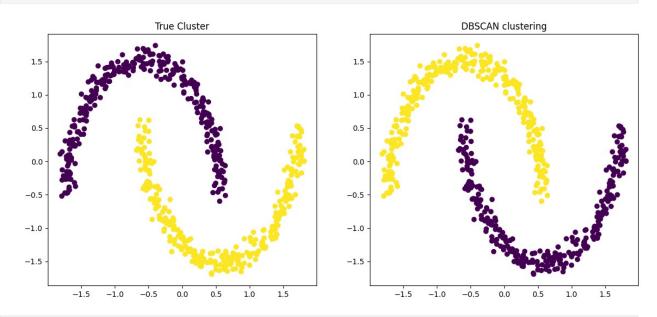
Adjusted mutual information: 1.000



For eps = 0.3 and minimum samples = 15

Silhouette score: 0.389 Adjusted Rand Index: 1.000

Adjusted mutual information: 1.000

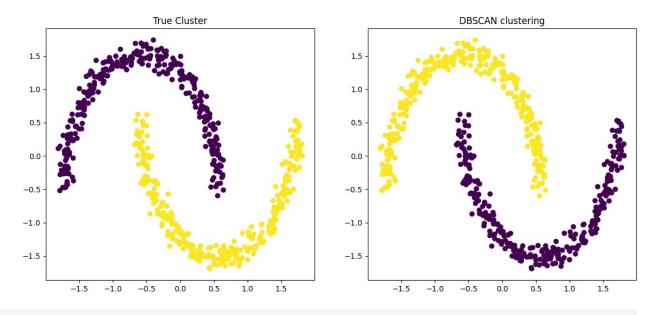


For eps = 0.3 and minimum samples = 20

Silhouette score: 0.389

Adjusted Rand Index: 1.000

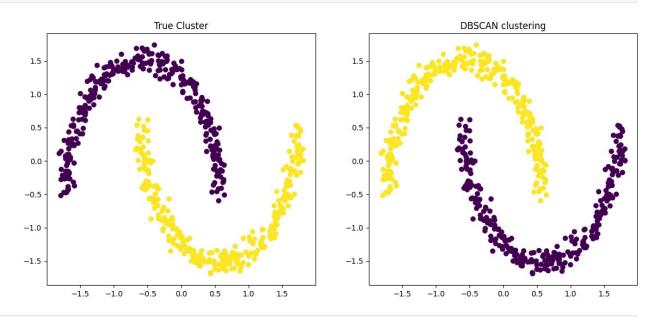
Adjusted mutual information: 1.000



For eps = 0.5 and minimum samples = 5

Silhouette score: 0.389 Adjusted Rand Index: 1.000

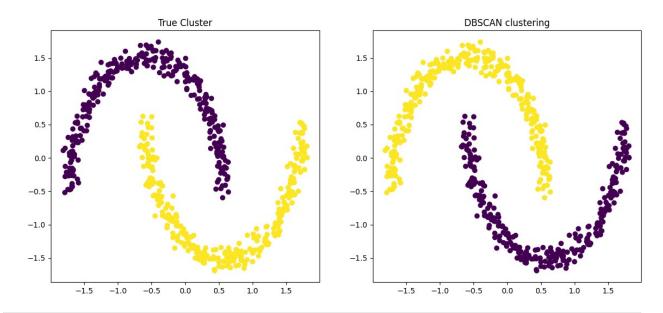
Adjusted mutual information: 1.000



For eps = 0.5 and minimum samples = 10

Silhouette score: 0.389 Adjusted Rand Index: 1.000

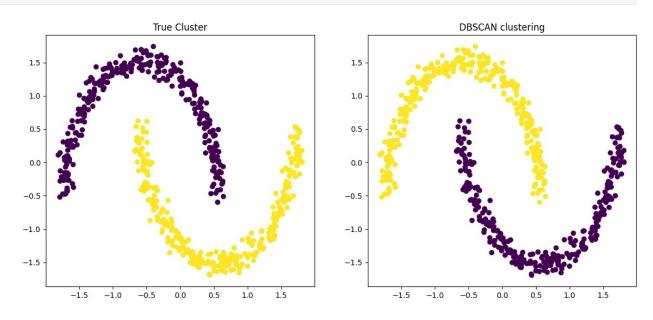
Adjusted mutual information: 1.000



For eps = 0.5 and minimum samples = 15

Silhouette score: 0.389 Adjusted Rand Index: 1.000

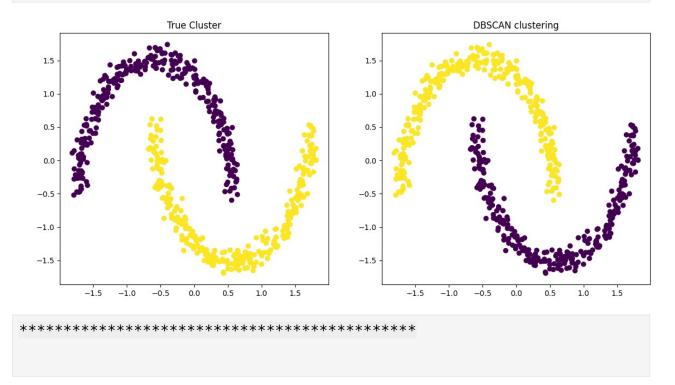
Adjusted mutual information: 1.000



For eps = 0.5 and minimum samples = 20

Silhouette score: 0.389 Adjusted Rand Index: 1.000

Adjusted mutual information: 1.000



Observations

- 1. For moon dataset, as we can see the scores are consistent irrespective of the eps and min_samples values.
- 2. Again we are getting ari and ami = 1, which implies that DBSCAN is doing the placing of the points.
- 3. So we can take any eps and min_sample value for moon dataset.