# Clustering

## 1. DBSCAN

Using DBSCAN iterate (for-loop) through different values of min\_samples (1 to 10) and epsilon (.05 to .5, in steps of .01) to find clusters in the road-data used in the Lesson and calculate the Silohouette Coeff for min\_samples and epsilon . Plot **one** line plot with the multiple lines generated from the min\_samples and epsilon values. Use a 2D array to store the SilCoeff values, one dimension represents min\_samples , the other represents epsilon.

Expecting a plot of epsilon vs sil\_score.

```
import pandas as pd
import numpy as np
%matplotlib notebook
import matplotlib.pyplot as plt
import seaborn
from mpl_toolkits.mplot3d import Axes3D
plt.rcParams['font.size'] = 14
# plt.rcParams['figure.figsize'] = (20.0, 10.0)
from sklearn.cluster import DBSCAN
from sklearn.metrics import silhouette_score
```

```
In [2]: # Reading in road data
X = pd.read_csv('3D_spatial_network.txt.gz', header=None, names=['osm'
X = X.drop(['osm'], axis=1).sample(1000)
X.head()
```

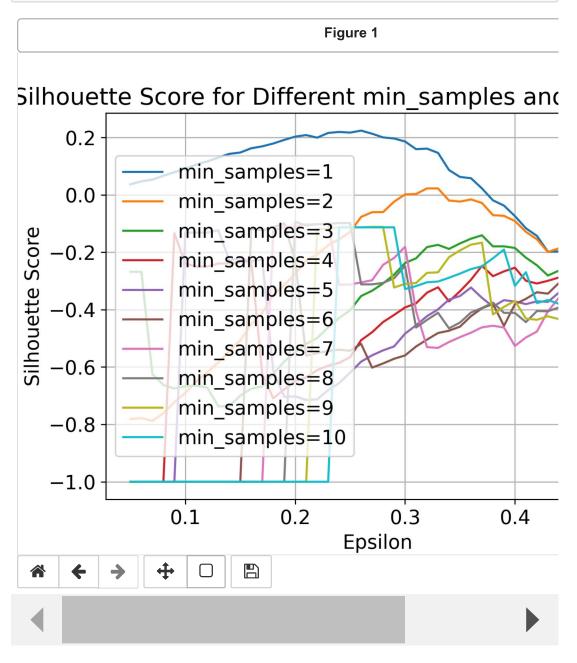
### Out[2]:

	lat	lon	alt
234401	8.449426	56.855887	13.494604
286099	9.976817	57.323641	36.328023
48980	8.789672	56.890996	46.130555
145711	9.593939	57.251441	10.187147
39950	8.150701	56.601929	2.199211

```
In [3]: ▶ # Setting up ranges for min_samples and epsilon
            min_samples = np.arange(1,11,1)
            epsilons = np.arange(0.05, 0.51, 0.01)
            # Checking work
            print(min_samples, epsilons)
            [ 1 2 3 4 5 6 7 8 9 10] [0.05 0.06 0.07 0.08 0.09 0.1 0.11
            0.12 0.13 0.14 0.15 0.16 0.17 0.18
             0.19 0.2 0.21 0.22 0.23 0.24 0.25 0.26 0.27 0.28 0.29 0.3 0.31 0.3
            2
             0.33 0.34 0.35 0.36 0.37 0.38 0.39 0.4 0.41 0.42 0.43 0.44 0.45 0.4
             0.47 0.48 0.49 0.5 ]
In [6]: ▶ # Initializing list to store silhouette scores
            all_scores = []
            for min_sample in min_samples:
                scores = []
                for epsilon in epsilons:
                    # Applying DBSCAN
                    dbscan = DBSCAN(eps=epsilon, min samples=min sample)
                    labels = dbscan.fit predict(X)
                    # Calculate silhouette score
                   try:
                        score = silhouette_score(X, labels)
                    except:
                        score = -1 # Set to -1 if silhouette score calculation fa
                    scores.append(score)
                all_scores.append(scores)
            # Converting the list to a numpy array for easier manipulation
            silhouette_scores = np.array(all_scores)
```

```
In [12]:  # Plotting the results
    plt.figure(figsize=(7,5))
    for i, min_sample in enumerate(min_samples):
        plt.plot(epsilons, silhouette_scores[i, :], label=f'min_samples={m

    plt.xlabel('Epsilon')
    plt.ylabel('Silhouette Score')
    plt.title('Silhouette Score for Different min_samples and Epsilon Valu
    plt.legend()
    plt.grid(True)
    plt.show()
```



# 2. Clustering your own data

Using your own data, find relevant clusters/groups within your data (repeat the above). If your data is labeled with a class that you are attempting to predict, be sure to not use it in training and clustering.

You may use the labels to compare with predictions to show how well the clustering performed using one of the clustering metrics (http://scikitlearn.org/stable/modules/clustering.html#clustering-performance-evaluation (http://scikit-

<u>learn.org/stable/modules/clustering.html#clustering-performance-evaluation)</u>).

If you don't have labels, use the silhouette coefficient to show performance. Find the optimal fit for your data but you don't need to be as exhaustive as above.

Additionally, show the clusters in 2D or 3D plots.

As a bonus, try using PCA first to condense your data from N columns to less than N.

Two items are expected:

- Metric Evaluation Plot (like in 1.)
- · Plots of the clustered data

```
In [18]:
         penguin = pd.read_csv('penguins.csv')
            # Dropping rows with missing data
            penguin.dropna(inplace=True)
            # Converting 'sex' to binary variable
            penguin['sex'] = penguin['sex'].apply(lambda x: 1 if x == 'FEMALE' els
            # Previewing data
            penguin.head()
```

#### Out[18]:

	culmen_length_mm	culmen_depth_mm	flipper_length_mm	body_mass_g	sex
0	39.1	18.7	181.0	3750.0	0
1	39.5	17.4	186.0	3800.0	1
2	40.3	18.0	195.0	3250.0	1
4	36.7	19.3	193.0	3450.0	1
5	39.3	20.6	190.0	3650.0	0

## In [28]: import os import pandas as pd import numpy as np import matplotlib.pyplot as plt from sklearn.cluster import KMeans from sklearn.metrics import silhouette\_score from sklearn.decomposition import PCA from sklearn.preprocessing import StandardScaler # Set OMP NUM THREADS environment variable to avoid memory leak warnin os.environ['OMP\_NUM\_THREADS'] = '4'

```
In [30]:  

# Standardizing the features
             scaler = StandardScaler()
             X_scaled = scaler.fit_transform(X)
             # Applying PCA to reduce dimensionality for visualization
             pca = PCA(n_components=3)
             X_pca = pca.fit_transform(X_scaled)
             # Finding optimal number of clusters using silhouette score
             silhouette scores = []
             for n clusters in range(2, 8): # trying clusters from 2 to 7
                 kmeans = KMeans(n clusters=n clusters, n init=10, random state=42)
                 cluster labels = kmeans.fit predict(X scaled)
                 silhouette avg = silhouette score(X scaled, cluster labels)
                 silhouette scores.append(silhouette avg)
             # Plotting silhouette scores
             plt.figure(figsize=(10, 6))
             plt.plot(range(2, 8), silhouette scores, marker='o', linestyle='--')
             plt.title('Silhouette Scores for Different Numbers of Clusters')
             plt.xlabel('Number of Clusters')
             plt.ylabel('Silhouette Score')
             plt.xticks(range(2, 8))
             plt.grid(True)
             plt.show()
             # Choosing optimal number of clusters based on silhouette score
             optimal_n_clusters = silhouette_scores.index(max(silhouette_scores)) +
             print(f"Optimal number of clusters based on silhouette score: {optimal
             # Performing K-means clustering with optimal number of clusters
             kmeans = KMeans(n_clusters=optimal_n_clusters, n_init=10, random_state
             cluster_labels = kmeans.fit_predict(X_scaled)
             # Visualizing clusters in 2D using PCA components
             plt.figure(figsize=(10, 6))
             plt.scatter(X_pca[:, 0], X_pca[:, 1], c=cluster_labels, cmap='viridis'
             plt.title(f'Clusters of Penguins (PCA-reduced Data, {optimal_n_cluster
             plt.xlabel('Principal Component 1')
             plt.ylabel('Principal Component 2')
             plt.colorbar(label='Cluster')
             plt.grid(True)
             plt.show()
```

C:\Users\kaoui\anaconda3\lib\site-packages\sklearn\cluster\ kmeans.p y:1382: UserWarning: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks than available threads. You can avoid it by setting the environment variable OMP\_NUM\_THREADS=4. warnings.warn(

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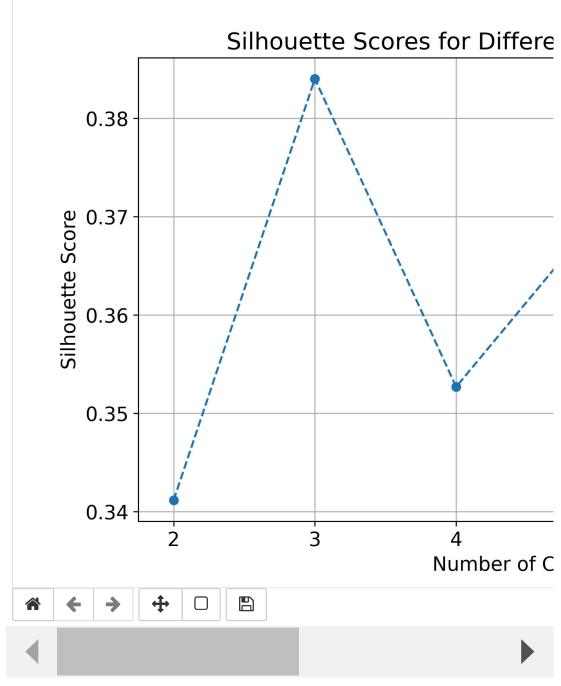
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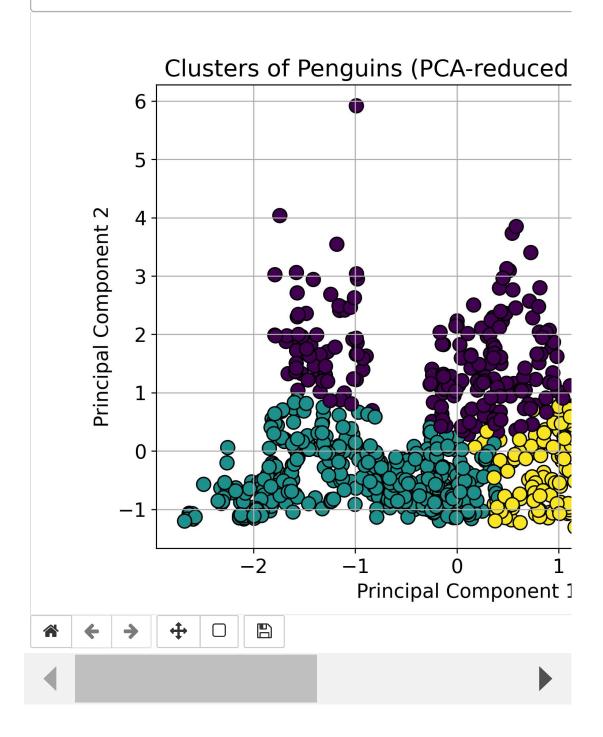




Optimal number of clusters based on silhouette score: 3

C:\Users\kaoui\anaconda3\lib\site-packages\sklearn\cluster\\_kmeans.p
y:1382: UserWarning: KMeans is known to have a memory leak on Windows
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 warnings.warn(



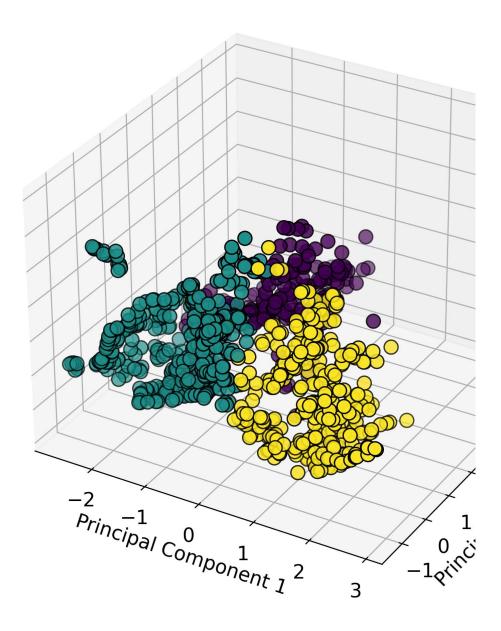


```
In [32]: # Visualizing clusters in 3D using PCA components
fig = plt.figure(figsize=(10, 8))
ax = fig.add_subplot(111, projection='3d')

scatter = ax.scatter(X_pca[:, 0], X_pca[:, 1], X_pca[:, 2], c=cluster_
ax.set_title(f'Clusters of Penguins (PCA-reduced Data, 3 clusters)')
ax.set_xlabel('Principal Component 1')
ax.set_ylabel('Principal Component 2')
ax.set_zlabel('Principal Component 3')
ax.legend(*scatter.legend_elements(), title='Cluster')
plt.show()
```

## Figure 6

# Clusters of Penguins (PCA-reduced Data,





In [ ]:	M	