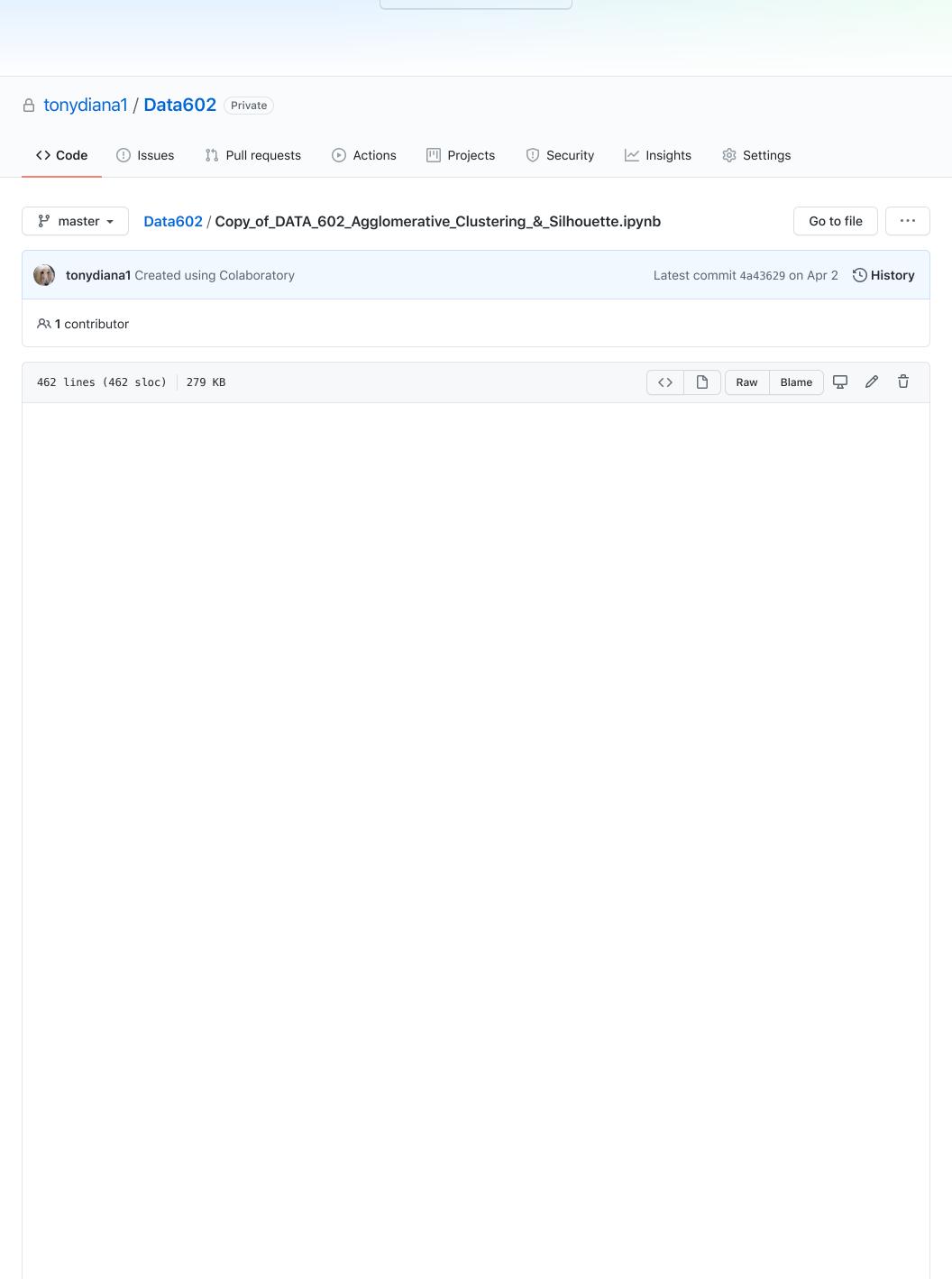


Learn Git and GitHub without any code!

Using the Hello World guide, you'll start a branch, write comments, and open a pull request.

Read the guide



EXERCISE 1: AGGLOMERATIVE CLUSTERING

```
In [0]: import pandas as pd
    import numpy as np
    import matplotlib.pyplot as plt
    from sklearn.decomposition import PCA
    from sklearn.cluster import AgglomerativeClustering
    from sklearn.preprocessing import StandardScaler, normalize
    from sklearn.metrics import silhouette_score
    import scipy.cluster.hierarchy as shc
In [0]: # Load the cc_general.csv file

X = pd.read_csv('/content/cc_general.csv')
```

```
In [0]: # Scaling the data so that all the features become comparable
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)

# Normalizing the data so that the data approximately
# follows a Gaussian distribution
X_normalized = normalize(X_scaled)

# Converting the numpy array into a pandas DataFrame
X normalized = pd.DataFrame(X normalized)
```

Use Principal Component Analysis with two components

Dropping the CUST ID column from the data

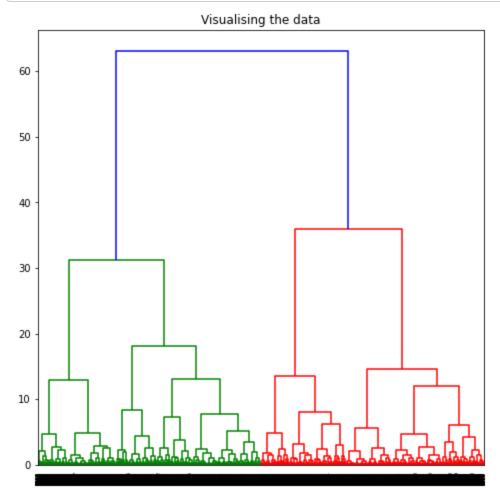
X = X.drop('CUST_ID', axis = 1)

Handling the missing values

```
In [0]: pca = PCA(n_components = 2)
X_principal = pca.fit_transform(X_normalized)
X_principal = pd.DataFrame(X_principal)
X_principal.columns = ['P1', 'P2']
```

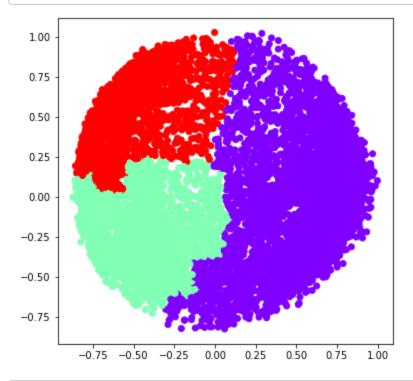
Plot a dendogram

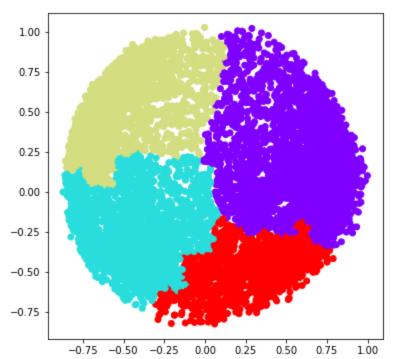
```
In [9]: plt.figure(figsize =(8, 8))
   plt.title('Visualising the data')
   Dendrogram = shc.dendrogram((shc.linkage(X_principal, method ='ward')))
```



Use Agglomerative Clustering with 2, 3, and 4 clusters

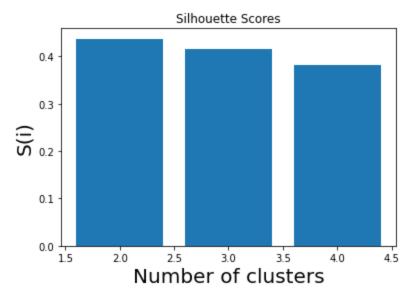
```
1.00 -
0.75 -
0.50 -
0.00 -
-0.25 -
-0.75 -0.50 -0.25 0.00 0.25 0.50 0.75 1.00
```





Compute the silhouette scores with K = 2, 3, and 4 The silhouette ranges from -1 to +1, where a high value indicates that the object is well matched to its own cluster and poorly matched to neighboring clusters. A value of 0 indicates that the sample is on or very close to the decision boundary between two neighboring clusters and negative values indicate that those samples might have been assigned to the wrong cluster. The Silhouette Coefficient is calculated using the mean intra-cluster distance (a) and the mean nearest-cluster distance (b) for each sample. The Silhouette Coefficient for a sample is (b - a) / max(a, b).

```
In [14]: # Plotting a bar graph to compare the results
   plt.bar(k, silhouette_scores)
   plt.title('Silhouette Scores')
   plt.xlabel('Number of clusters', fontsize = 20)
   plt.ylabel('S(i)', fontsize = 20)
   plt.show()
```

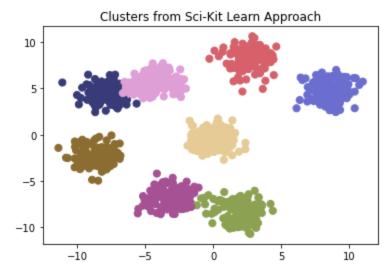


EXERCISE 2: AGGLOMERATIVE CLUSTERING

Illustration on how to use linkage and affinity, as well as sklearn and SciPy approach

```
In [0]: from sklearn.cluster import AgglomerativeClustering
    from sklearn.datasets import make_blobs
    import matplotlib.pyplot as plt
    from scipy.cluster.hierarchy import linkage, dendrogram, fcluster
    ac = AgglomerativeClustering(n_clusters = 8, affinity="euclidean", linkage="average")
    X, y = make_blobs(n_samples=1000, centers=8, n_features=2, random_state=800)
    distances = linkage(X, method="centroid", metric="euclidean")
    sklearn_clusters = ac.fit_predict(X)
    scipy_clusters = fcluster(distances, 3, criterion="distance")
```

```
In [16]: plt.figure(figsize=(6,4))
   plt.title("Clusters from Sci-Kit Learn Approach")
   plt.scatter(X[:, 0], X[:, 1], c = sklearn_clusters ,s=50, cmap='tab20b')
   plt.show()
```



```
In [3]: plt.figure(figsize=(6,4))
    plt.title("Clusters from SciPy Approach")
    plt.scatter(X[:, 0], X[:, 1], c = scipy_clusters ,s=50, cmap='tab20b')
    plt.show()
```

