# DMG ASSIGNMENT - 2 CLUSTERING

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Jatin Tyagi: Contributed in developing the visualisation part in this assignment and also contributed in Question 1,2,3.

Pendyala Ritvik: Contributed in developing the visualisation part in this assignment and also contributed in Question 3,5.

Pratyush Jain: Made the Readme file for this assignment, he helped in choosing the datasets for the assignment and helped in Question 2,3,5.

Vatsal Lakhmani: Made the PPT for this assignment and made contributions in visualisation part and Question 2,5.

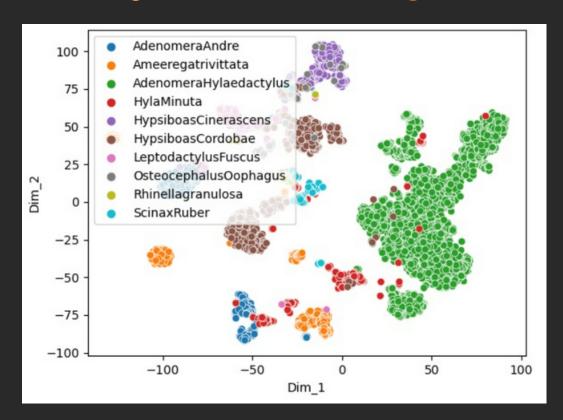
Yatish Garg: Made the Report file for this assignment and also made contributions in visualisation part and Question 2,4.



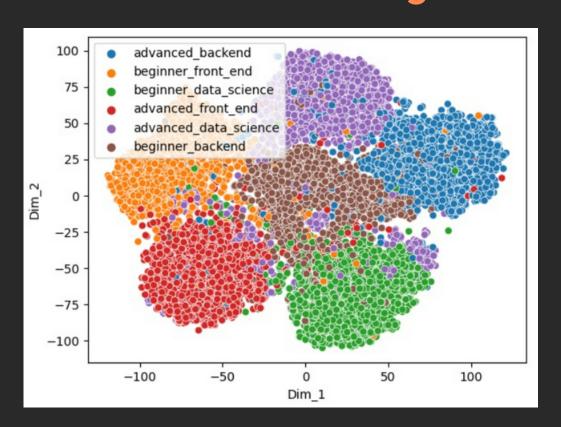
### **Density based clustering**

### Dataset 1

7195 rows22 featuresTarget: Species



### **Hierarchical clustering**



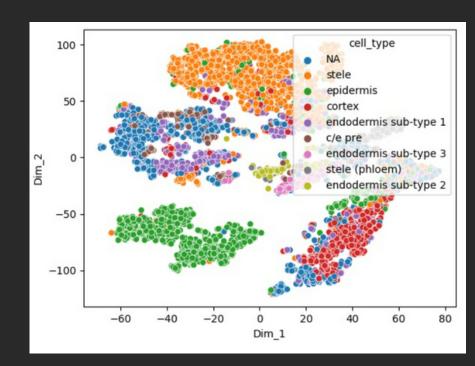
### Dataset 2

1000 rows12 featuresTarget: Profile

### **Prototype clustering**

### Dataset 3

5283rows17 featuresTarget: Profile





### Q) 2b Advantages

- Automatically discover arbitrarily shaped clusters when the algorithm is run.
- Find clusters completely surrounded by different clusters.
- It's Robust nature for the outlier detection makes it advantageous
- Require just 2 points, which are very insensitive to ordering the points in the database.
- It can automatically identify the noise data while clustering

### **Disadvantages**

- This is not partitionable at multiprocessor systems
- It becomes tricky with datasets of alternating densities.
- Sensitive towards minPoints and EVS
- Fails to identify clusters if densities vary and if the data is too sparse.
- Sampling affects density measures.
- When the dataset is of neck type it fails.

### **Q) 2c**

Density based clustering algorithm chosen is: **OPTICS** 

### **Advantages:**

- It does not require density parameters.
- The clustering order is useful for extracting the basic clustering information.
- It operates on variable epsilon.

# Q) 2d

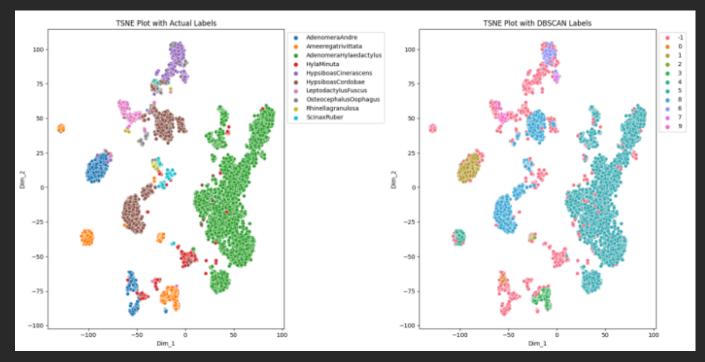
### **Dataset 1**

# 

epsilon: 1.9

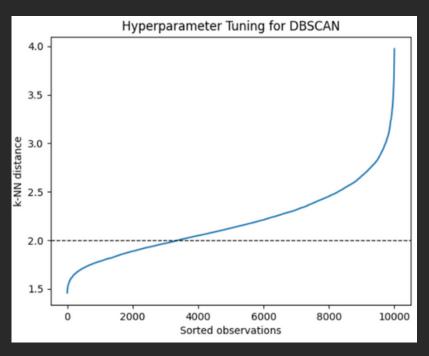
### **DBSCAN Implementation**

Silhouette Coefficient: 0.234
Davies Bouldin Score: 2.185
Adjusted Rand Index: 0.802
Adjusted Mutual Information: 0.698



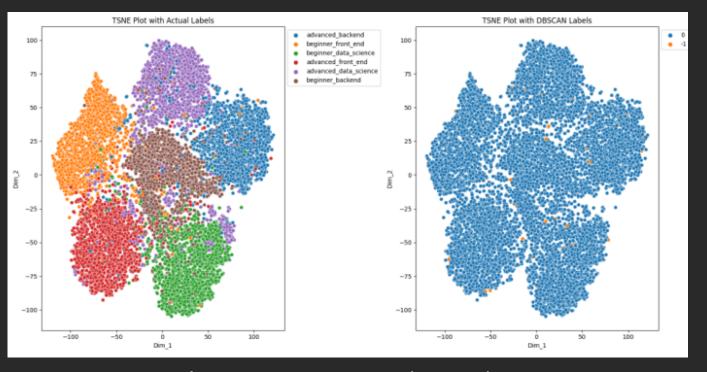
Est. Clusters: 10 Est. Noice Points: 1467

### **Dataset 2**



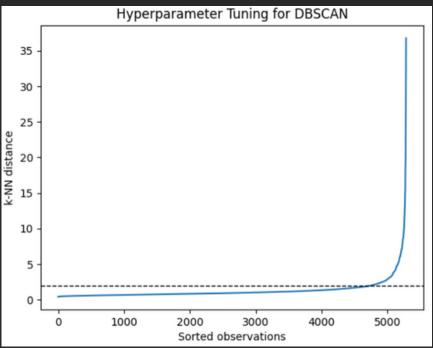
epsilon: 2.6

Silhouette Coefficient: 0.263
Davies Bouldin Score: 7.402
Adjusted Rand Index: 0.000
Adjusted Mutual Information: 0.000



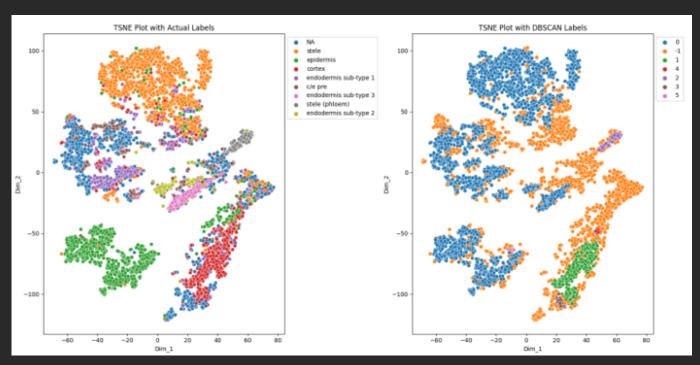
Est. Clusters: 1 Est. Noice Points: 51

### **Dataset 3**



epsilon: 0.7

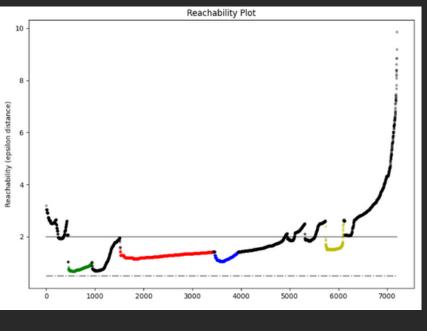
Silhouette Coefficient: -0.081 Davies Bouldin Score: 1.669 Adjusted Rand Index: 0.119 Adjusted Mutual Information: 0.206

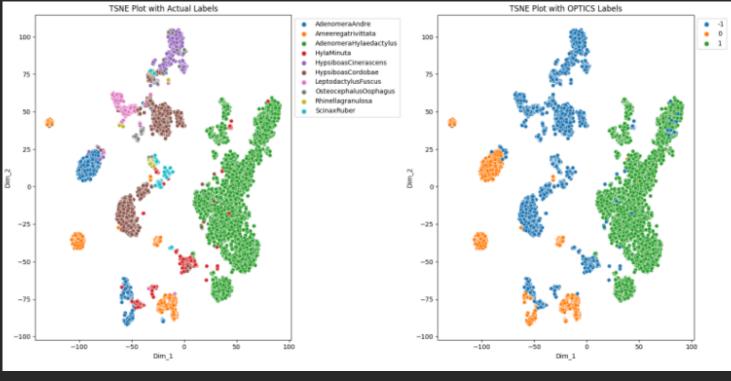


Est. Clusters: 6 Est. Noice Points: 1871

### **OPTICS Implementation**

### **Dataset 1**



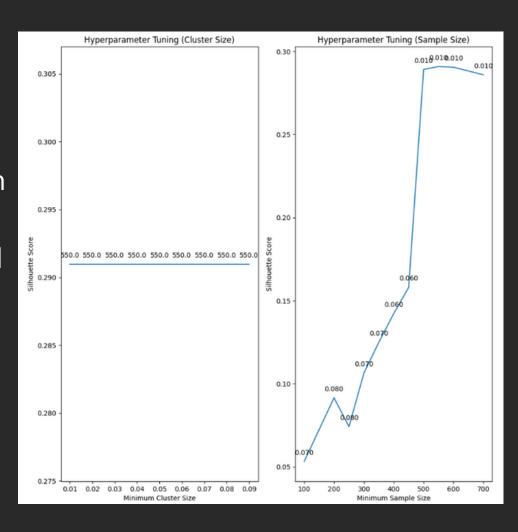


Reachability Plot

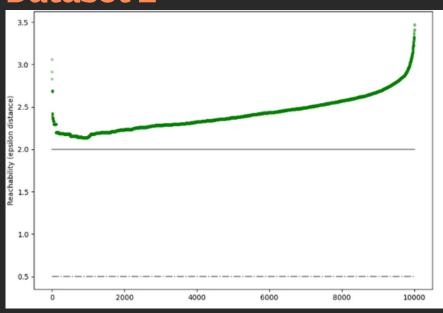
Optics Clustering Plot

Silhouette Coefficient: 0.2910 Davies Bouldin Score: 2.3854 Adjusted Rand Index: 0.7252 Adjusted Mutual Information:

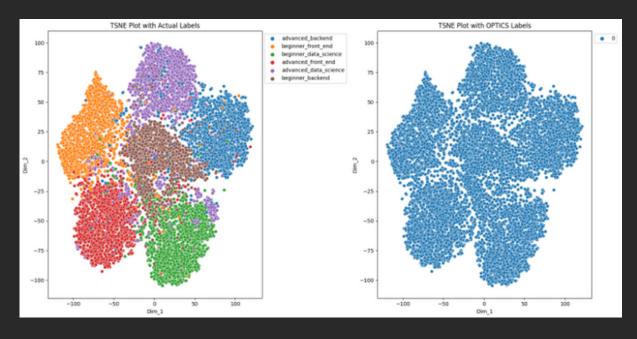
0.6879



### **Dataset 2**

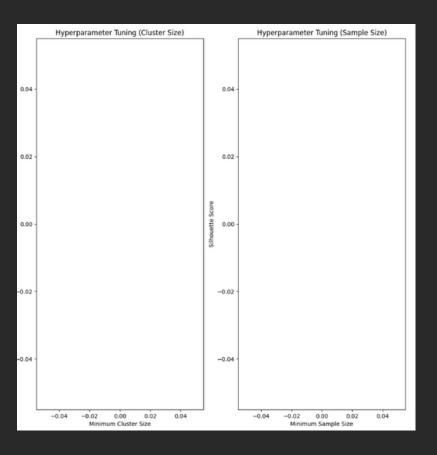


Reachability Plot

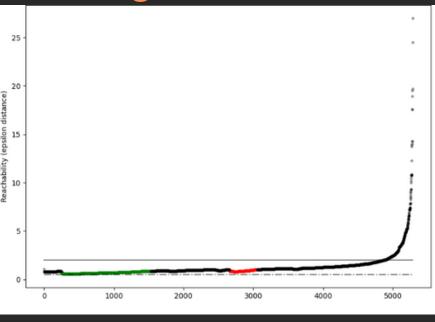


Optics Clustering Plot

Silhouette
Coefficient: -inf
Davies Bouldin
Score: inf
Adjusted Rand
Index: 0.0000
Adjusted Mutual
Information:
0.0000



**Dataset 3** 

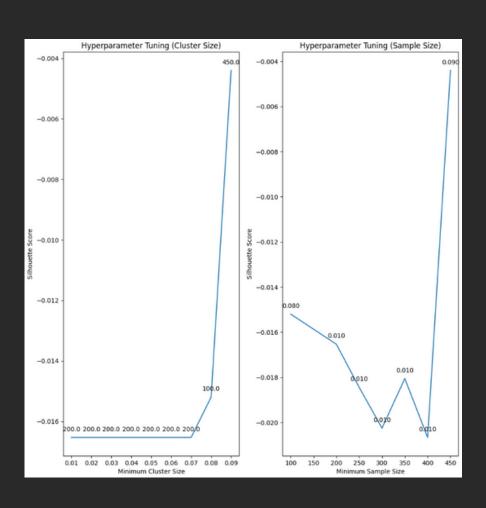


stelle spidermis sub-type 1 comex endodermis sub-type 2 sendodermis sub-type 3 sendodermis sub-type 2 sendodermis sub-type 3 sendodermis

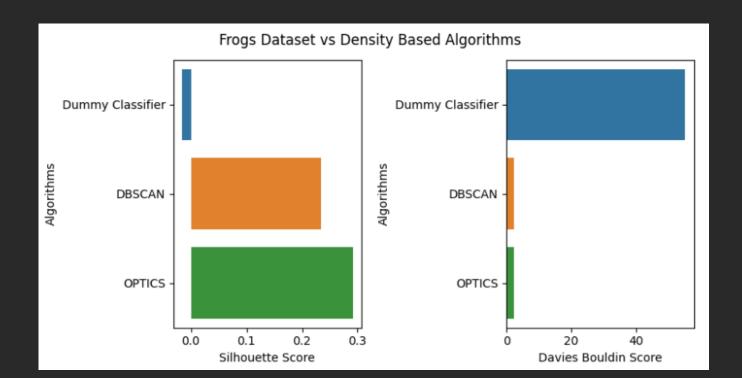
Reachability Plot

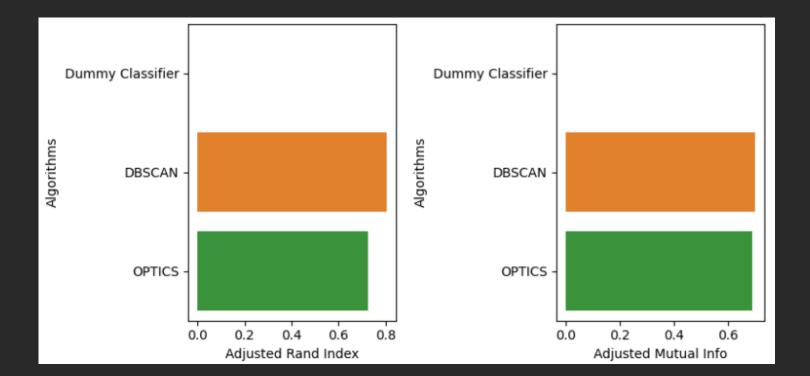
Optics Clustering Plot

Silhouette
Coefficient:
-0.0152
Davies Bouldin
Score: 39.68
Adjusted Rand
Index: -0.0024
Adjusted Mutual
Information:
0.0002

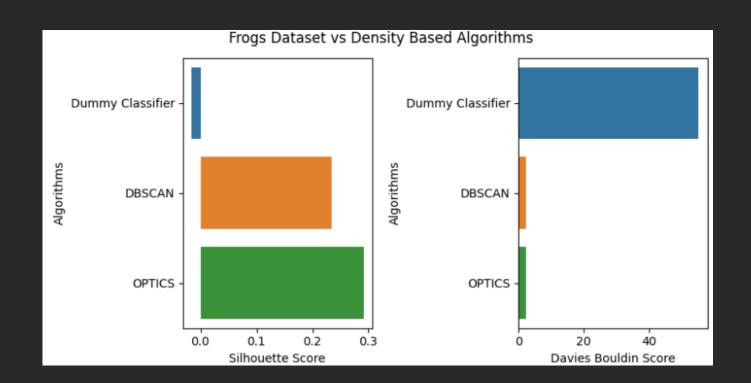


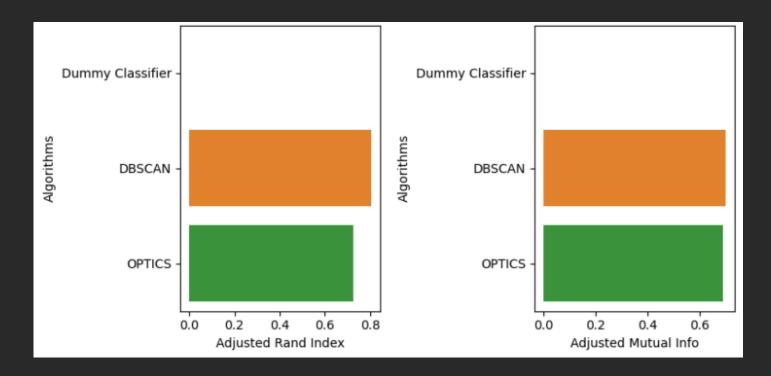
Q) 2e



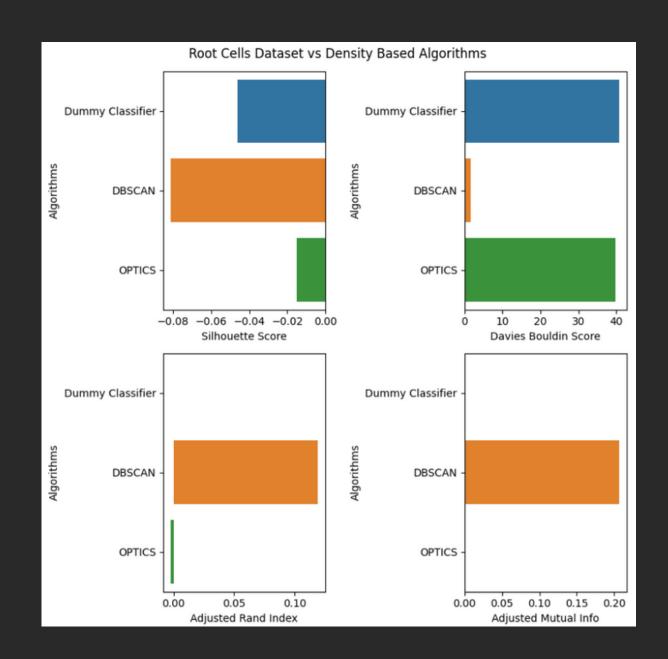


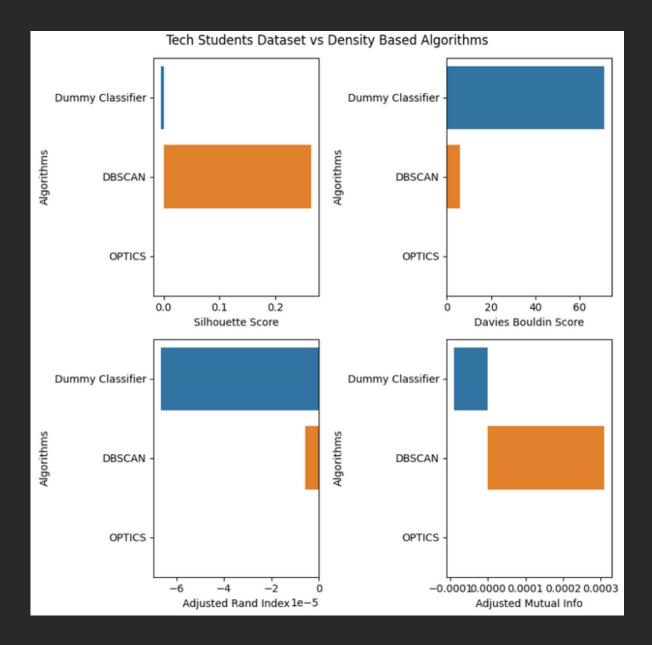
**Q) 2f** 



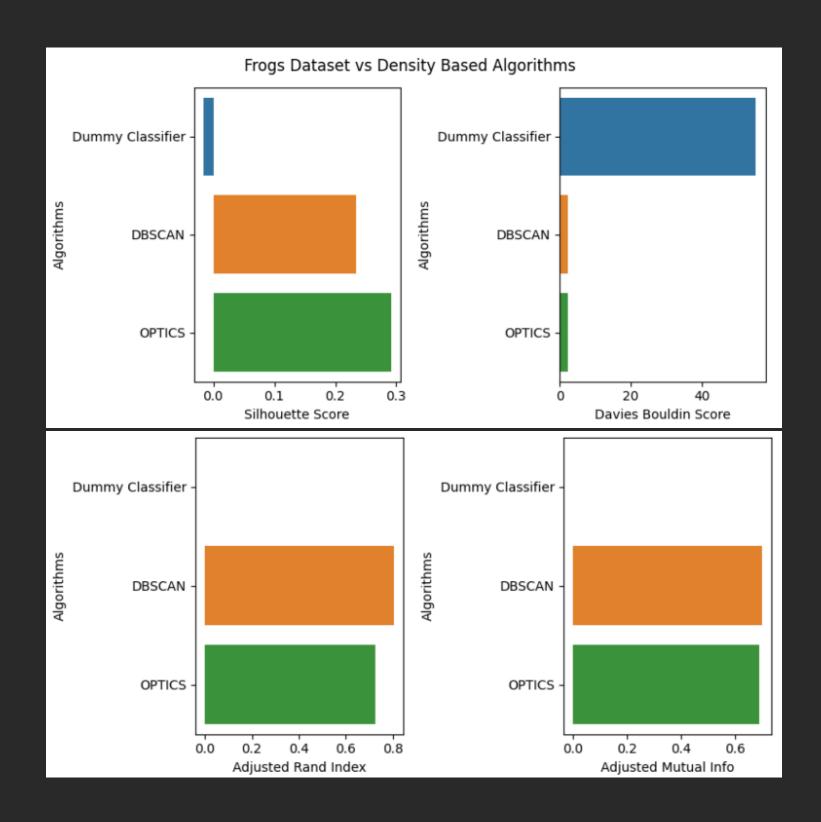


# **Q) 2g**





### **RESULTS**



Dataset 1 was the best fitting dataset in Q2.

According to the plotted graphs(As shown on the left side) DBSCAN showed slightly better results than Optics

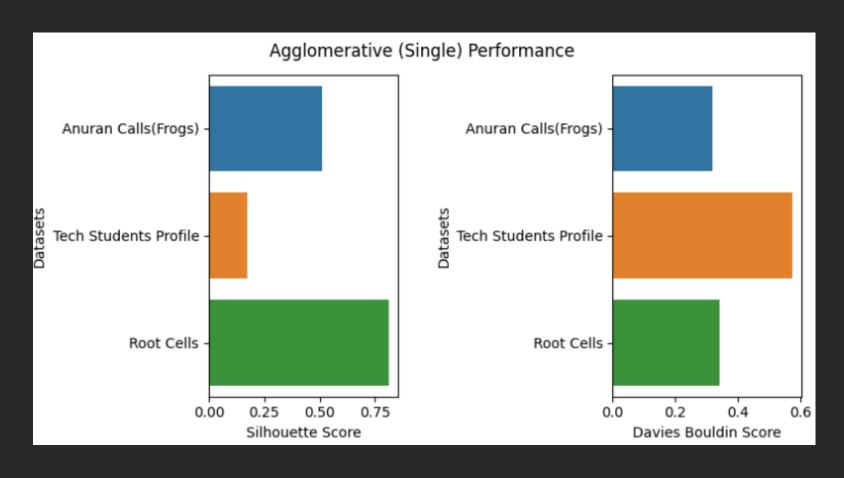
# Q) 3 Hierarchical clustering

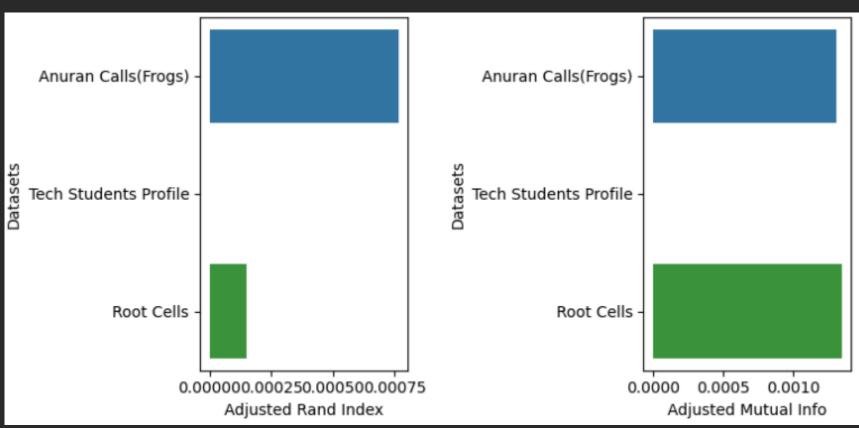
Two approaches: top-down(Agglomerative clustering) and bottom-up (Divisive Clustering)

### **Advantages**

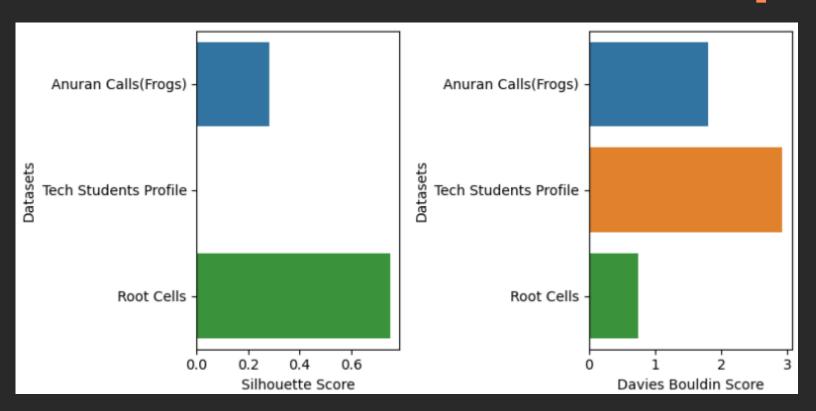
- There is no need to give any prior information about the number of clusters
- This algorithm is easy to implement as it gives best result.
- It works form the dissimilarities between the objects to be grouped together.

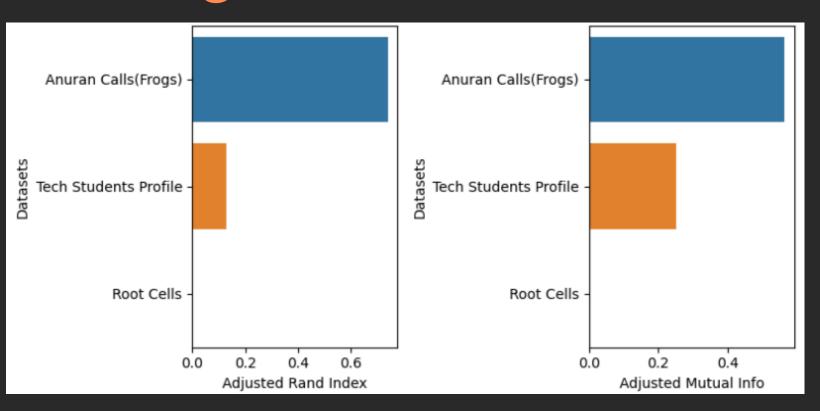
# Single linkage



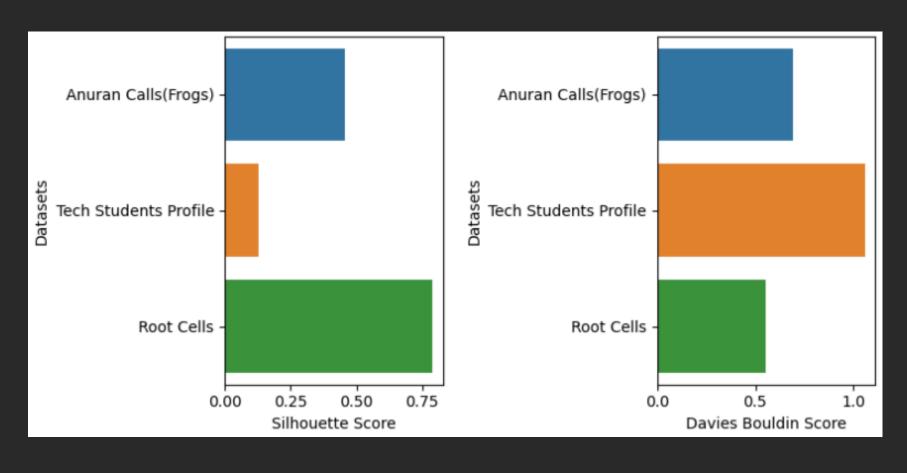


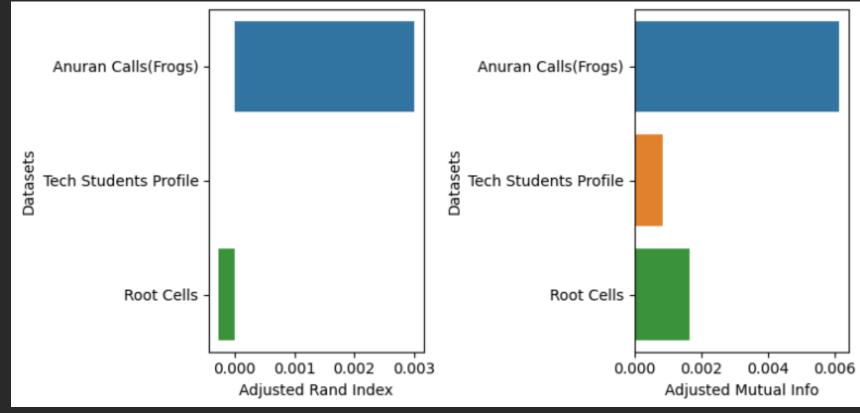
# Complete linkage



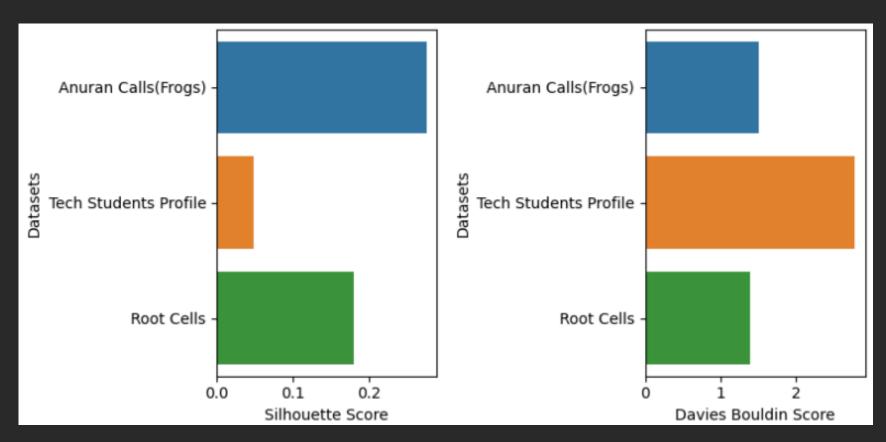


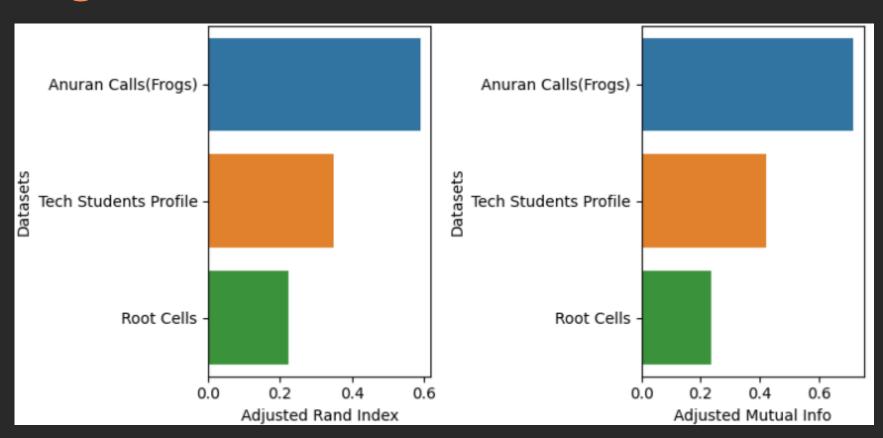
# Average linkage





# Ward linkage





# RESULTS

The best shown results were from the complete linkages and the wards linkage. Though wards linkage showed the best results in this case.

The best results were shown by the frog dataset, though tech students dataset showed appreciable results in it too.

# Q) 4 Prototype based clustering

Limitations of k-means:

- Clustering outliers
- Scaling with number of dimensions

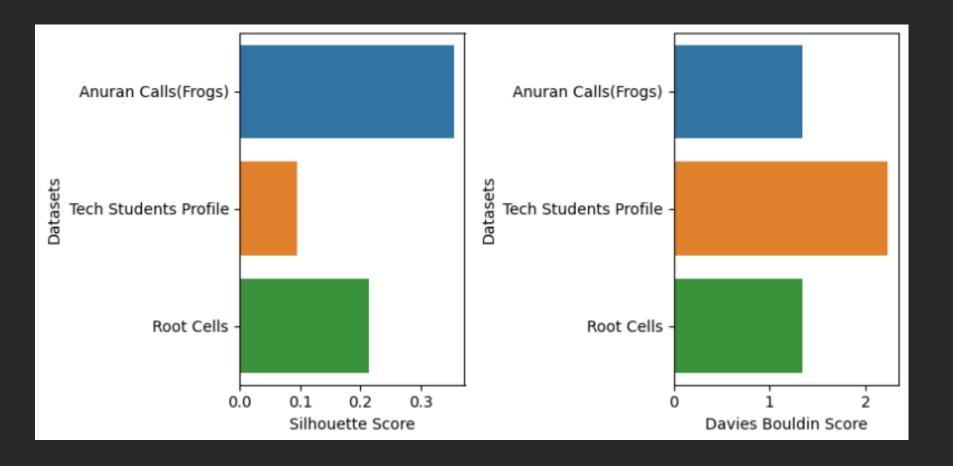
To encounter Clustering outliers limitation we use:

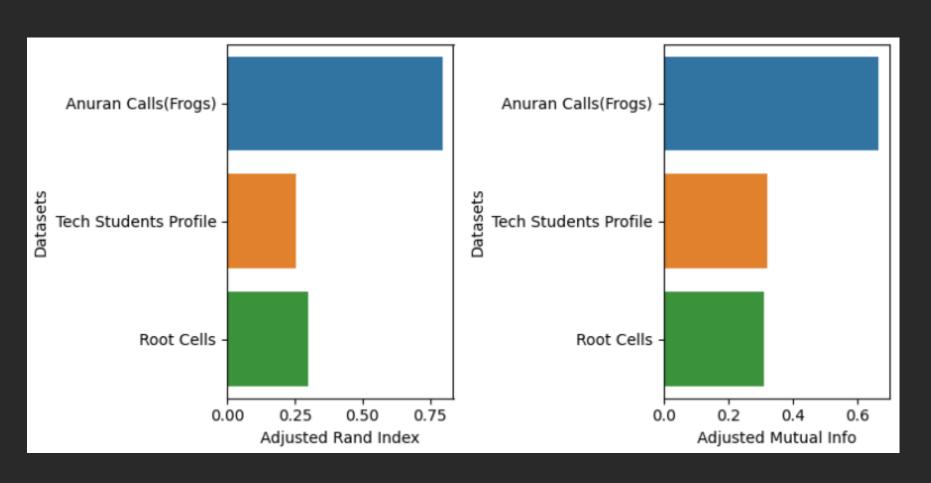
**Clustering Outliers: K- medoids clustering** 

To encounter scaling limitation we can use:

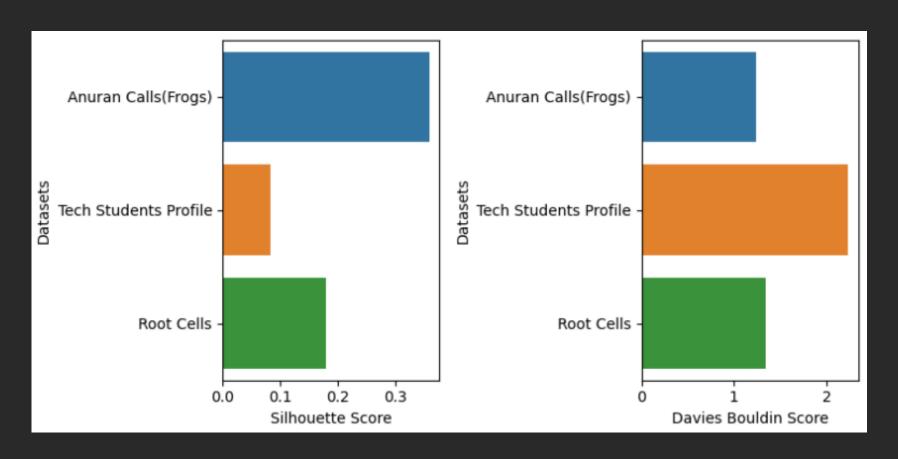
Number of Dimensions: Spectral Clustering

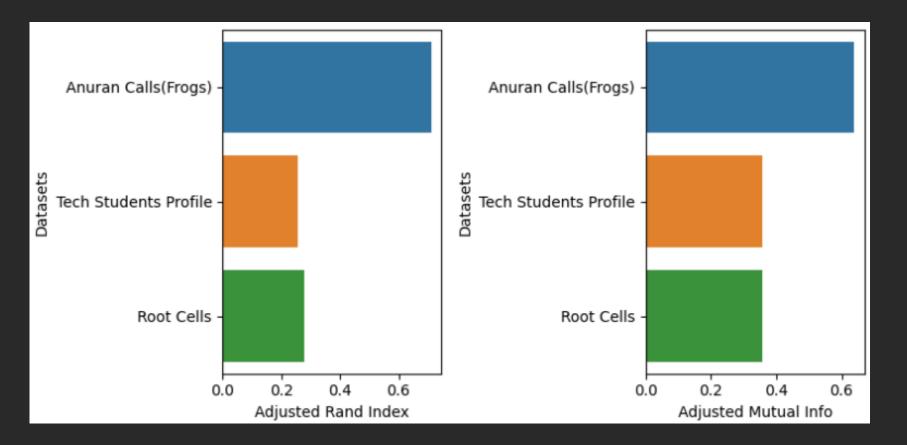
# **K means Performance**



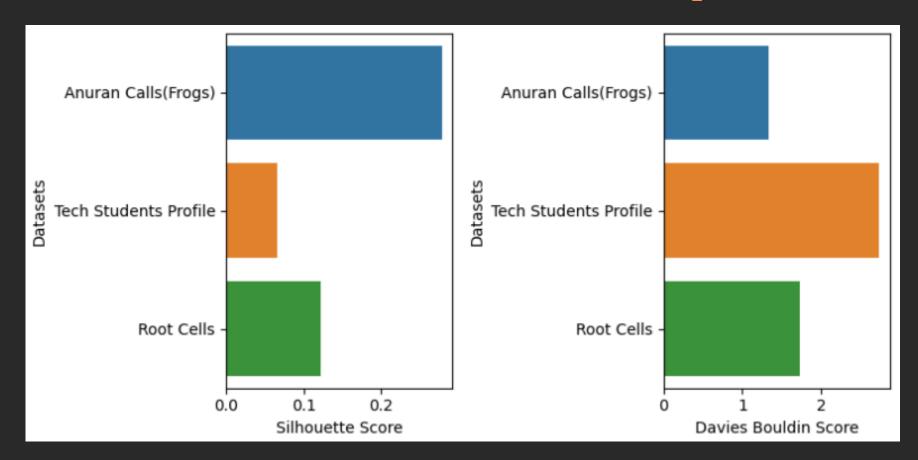


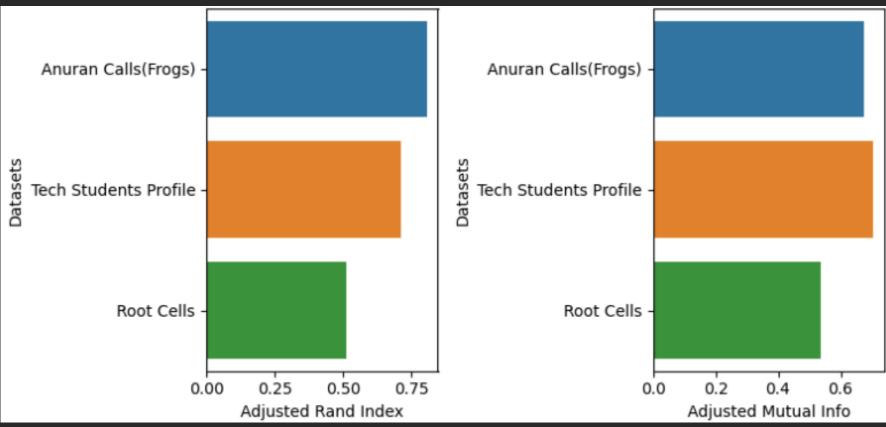
# **K medoid Performance**





# **Spectral Performance**





# RESULTS

The best shown results in this case was by the third dataset though other two datasets showed good enough results too.

# Thank You