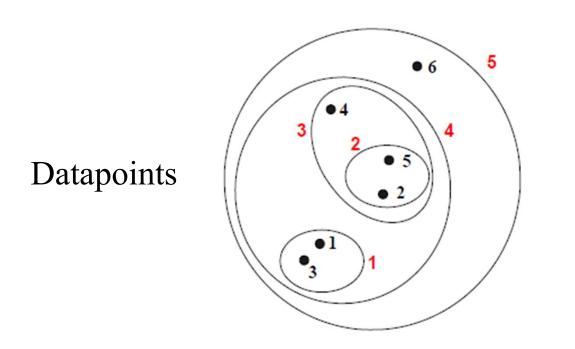
# HIERARCHICAL CLUSTERING

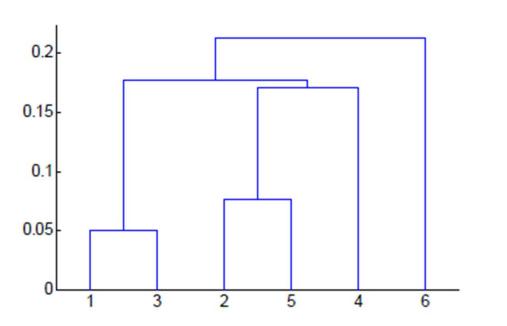
Sourav Karmakar

souravkarmakar29@gmail.com

## HIERARCHICAL CLUSTERING

- Produces a set of nested clusters organized as a hierarchical tree
- Can be visualized as a dendrogram: A tree like diagram that records the sequences of merges or splits.





Dendrograms

- Strengths of Hierarchical clustering:
- 1. Do not have to assume any particular number of clusters: Any desired number of clusters can be obtained by 'cutting' the dendrogram at the proper level.
- 2. They may correspond to meaningful taxonomies. Example: book cataloguing, biological sciences (e.g. animal kingdom)

### HIERARCHICAL CLUSTERING

#### Two main types of Hierarchical Clustering

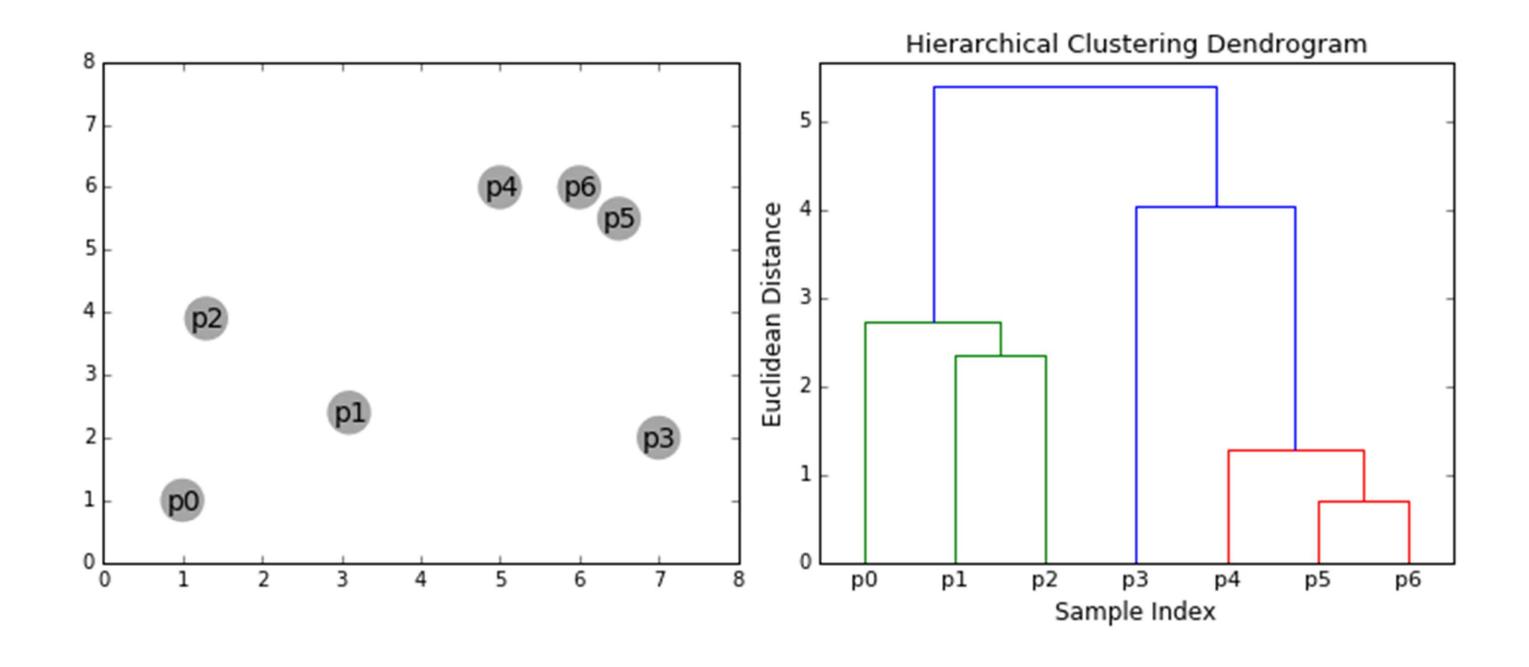
#### 1. Agglomerative: (Bottom Up Approach)

- Start with the points as individual clusters.
- At each step, merge the closest pair of clusters until you are with only one cluster (or k clusters).

#### 2. Divisive: (Top Down Approach)

- Start with one, all-inclusive cluster
- At each step, split a cluster until each cluster contains a point (or there are k clusters)

Though both the approaches are perfectly alright in producing quality clusters, people prefer to use Agglomerative approach more than Divisive approach. This is because it is less computationally expensive to deal with smaller clusters as compared to the large all inclusive cluster.



- Traditional Hierarchical Clustering algorithms use a similarity or distance matrix to split or merge the cluster. This matrix is known as **Proximity Matrix**.
- The basic Agglomerative Clustering algorithm is as following.
  - 1. Let each data point be a Cluster.
  - 2. Compute the proximity matrix.
  - 3. Repeat
  - 4. Merge the two closest clusters.
  - 5. Update the proximity matrix.
  - 6. Until only a single (or K many) cluster remains. (K is user provided number of clusters)
- Different approaches on defining the distance / proximity between two clusters distinguishes different algorithms. However the basic algorithm as shown above is followed everywhere in Agglomerative clustering.

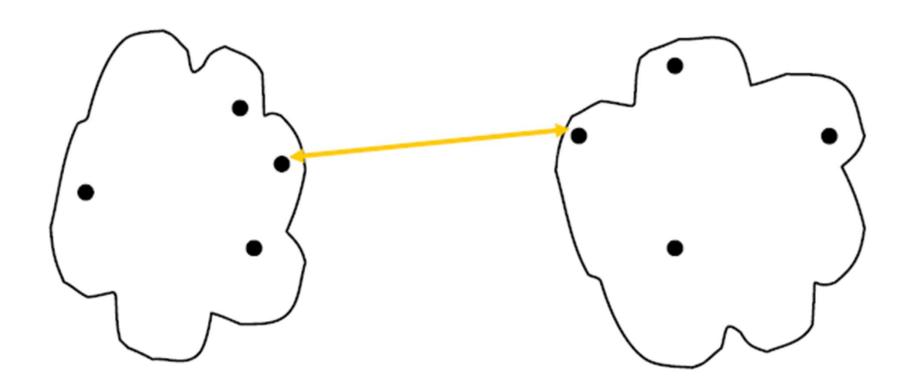
- How to define inter-cluster Distance / Proximity ?
- Following are the different ways to define the inter-cluster distance / proximity.
  - MIN or Single Linkage
  - MAX or Complete Linkage
  - Group Average
  - Distance Between Centroids

We shall now briefly discuss these methods.

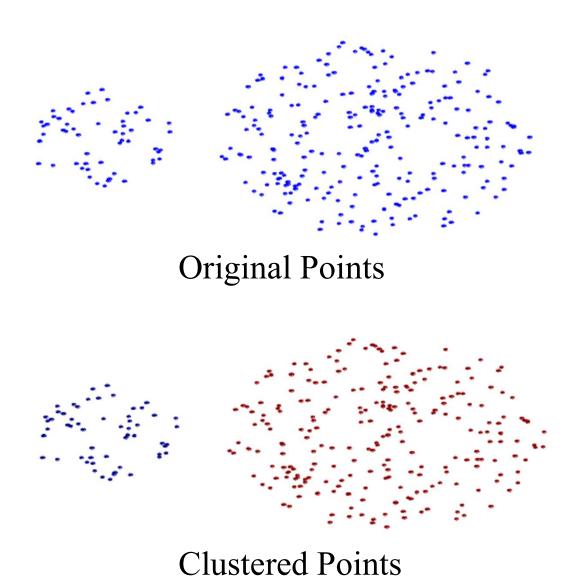
#### MIN or Single Linkage:

Here *minimum* of the all inter-cluster distances is considered for proximity calculation.

$$d_{C_i,C_j} = minimum \left\{ dist(\vec{x}, \vec{y}) : \vec{x} \in C_i \text{ and } \vec{y} \in C_j \right\}$$

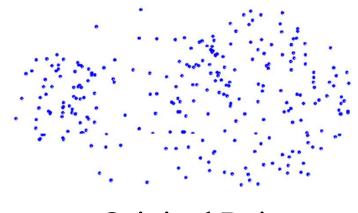


#### **Strength of MIN or Single Linkage**

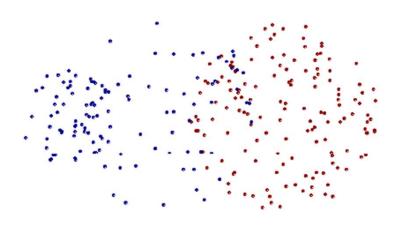


Single Linkage can handle shapes of different sizes

#### **Limitation of MIN or Single Linkage**



**Original Points** 



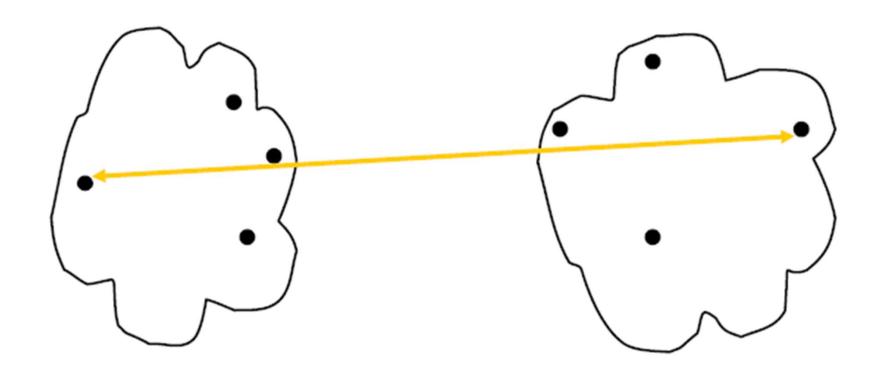
**Clustered Points** 

Single Linkage is susceptible to noise and outliers

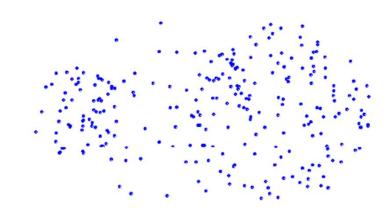
#### **MAX or Complete Linkage:**

Here *maximum* of the all inter-cluster distances is considered for proximity calculation.

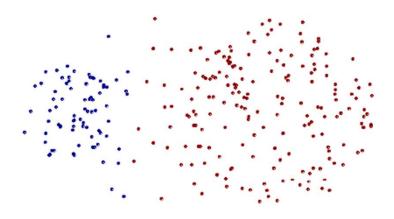
$$d_{C_i,C_j} = maximum \left\{ dist(\vec{x}, \vec{y}) : \vec{x} \in C_i \ and \ \vec{y} \in C_j \right\}$$



#### **Strength of MAX or Complete Linkage**



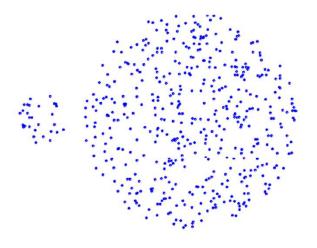
**Original Points** 



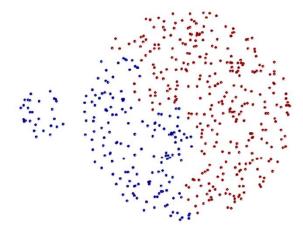
**Clustered Points** 

Complete Linkage is less susceptible to noise

#### **Limitation of MAX or Complete Linkage**



**Original Points** 



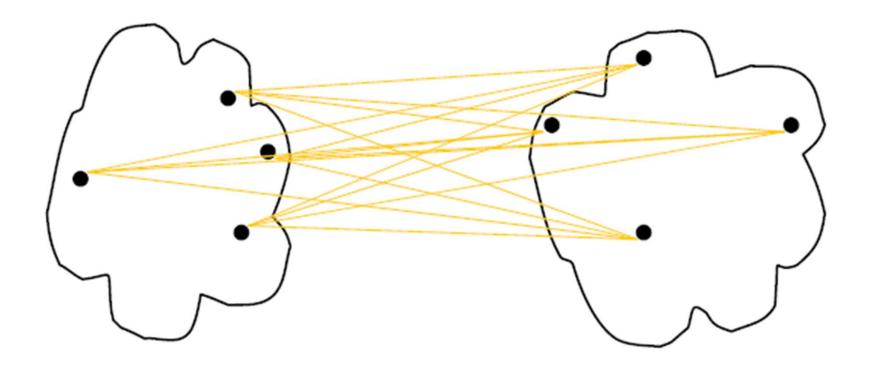
**Clustered Points** 

Complete Linkage tends to break large clusters

#### **Group Average:**

Here Average of the all inter-cluster distances is considered for proximity calculation.

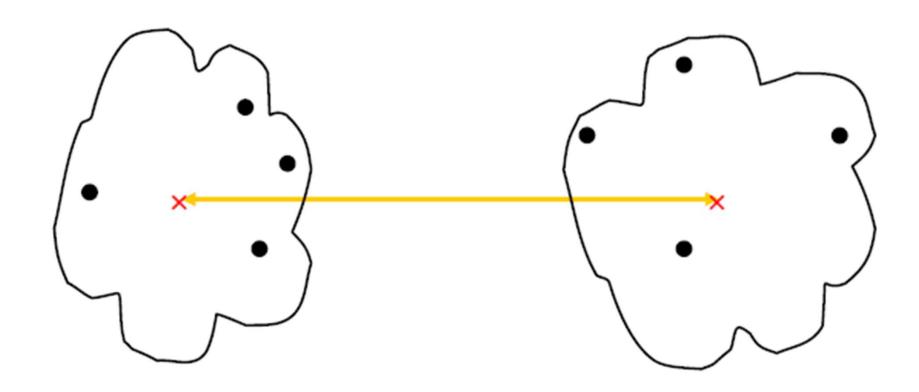
$$d_{C_i,C_j} = Average\left\{dist(\vec{x},\vec{y}) : \vec{x} \in C_i \text{ and } \vec{y} \in C_j\right\} = \frac{1}{|C_i||C_j|} \sum_{\vec{x} \in C_i, \vec{y} \in C_i} dist(\vec{x},\vec{y})$$



#### **Distance Between Centroids:**

Centroid of cluster  $C_i$  is computed as  $\vec{\mu}_i = \frac{1}{|C_i|} \sum_{\vec{x} \in C_i} \vec{x}$ , Similarly we compute centroid  $\vec{\mu}_j$  of cluster  $C_j$ .

$$d_{C_i,C_j} = dist\left(\overrightarrow{\boldsymbol{\mu}}_i,\overrightarrow{\boldsymbol{\mu}}_j\right)$$



## Thank You