

# Hierarchical Clustering

TEAM YOLO

Zeqing Jin

Xianlin Shao

Yifei Zhang

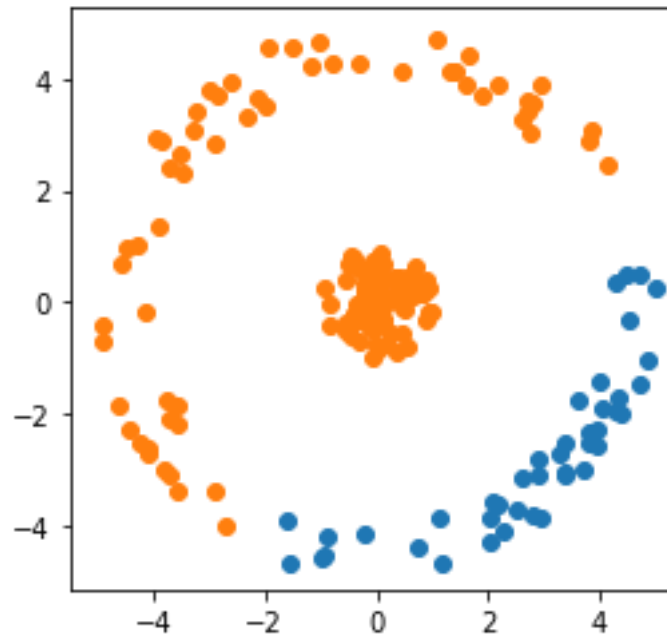
Zilan Zhang

# Introduction

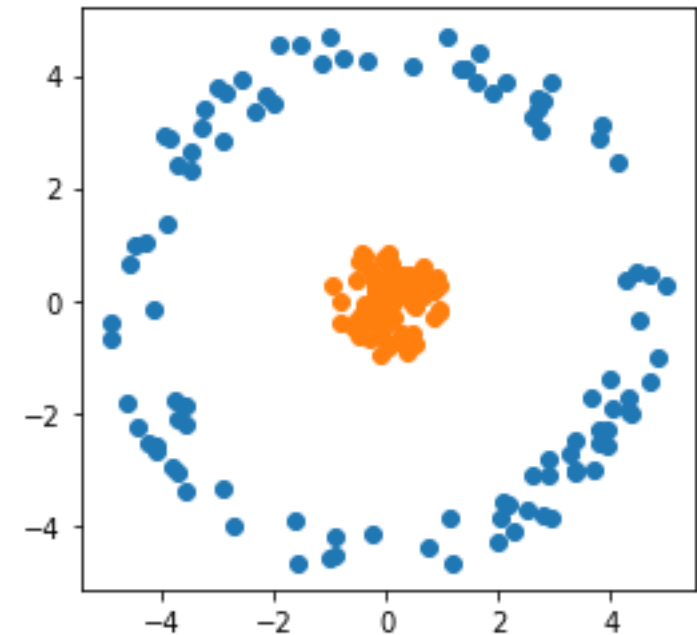
- Hierarchical clustering is a general family of clustering algorithms that build clusters by merging or splitting them successively. <sup>[2]</sup>
- Two common hierarchy algorithms:
  - Agglomerative clustering
  - Divisive clustering

# Limitations of K-means Clustering

- Non-spherical data points
- Prior assumption of similar number of data points in each cluster



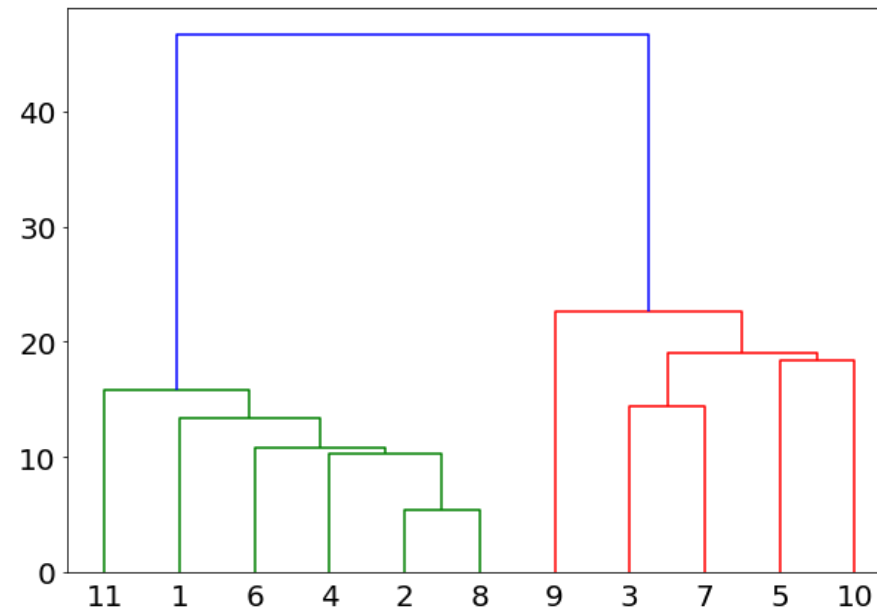
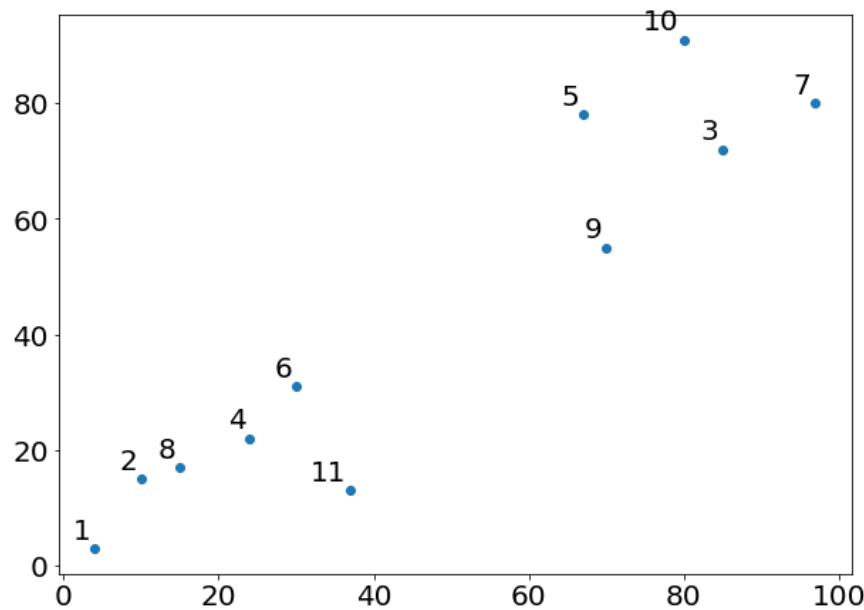
K-means



Hierarchical

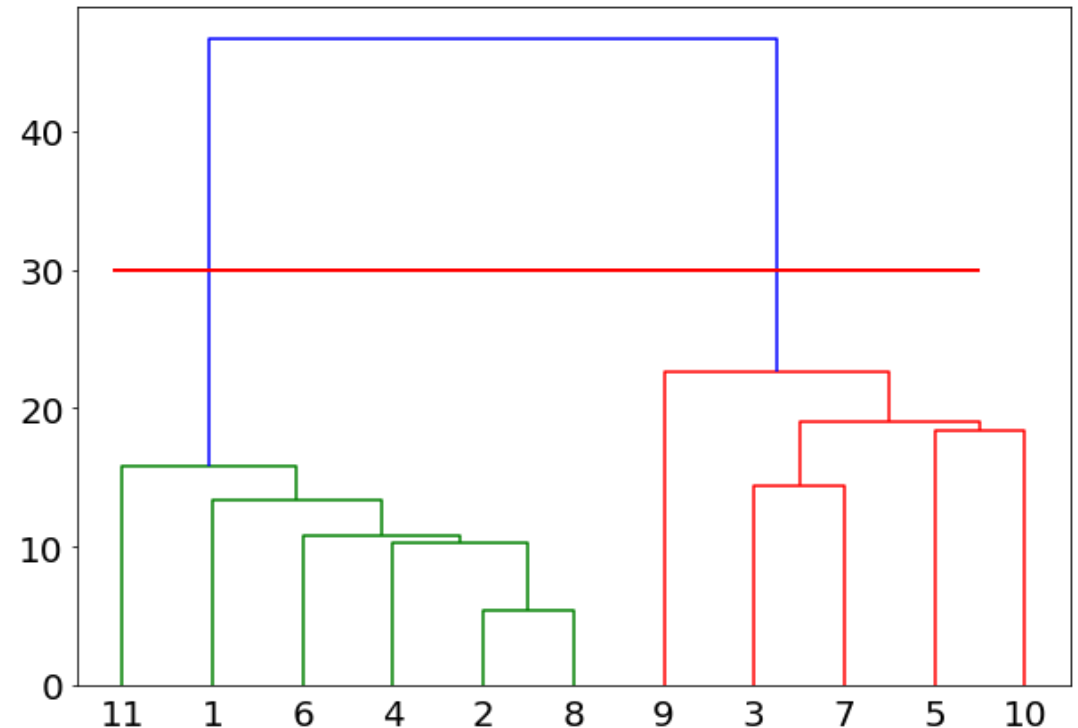
# Dendrogram

- Dendrogram is a tree-like hierarchy which shows the relationship between objects.



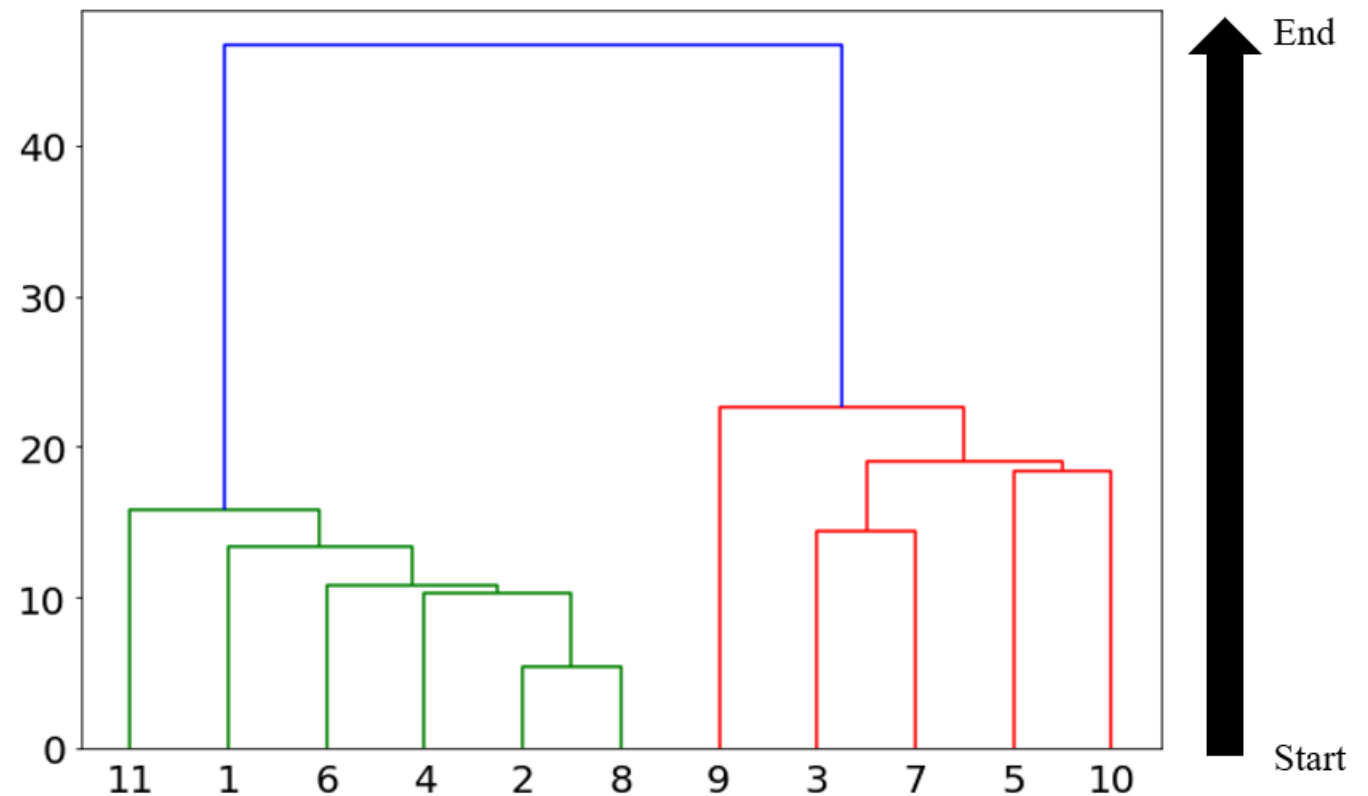
# Dendrogram

- Dendrogram implicitly contains all possible values of the number of clusters
- Shows relative relations between clusters (points)
- Fails to show all the absolute distances between points



# Agglomerative Clustering

- Start with  $n$  clusters containing one single point.
- End up with one cluster containing  $n$  objects.



# Agglomerative Clustering <sup>[1]</sup>

---

**Algorithm 1:** Agglomerative Hierarchical Clustering

---

**Input:**  $n$  data points

**Output:** final clustering result over  $n$  data points

- 1 Initialize  $n$  clusters  $\mathbf{c}_i, i = 1, \dots, n$ ;
  - 2 Initialize the dissimilarity matrix;
  - 3 **for** *the number of clusters  $k$  decreases from  $n$  to 1* **do**
  - 4     Merge the two clusters  $\mathbf{c}_i, \mathbf{c}_j$  with the smallest dissimilarity according to dissimilarity matrix;
  - 5     Update the dissimilarity matrix;
  - 6 **end for**
-

# Agglomerative Clustering

- Euclidean distance between points:

$$d(i, j) = \sqrt{\sum_{p=1}^q (x_{ip} - x_{jp})^2}$$

where  $q$  is dimension of the point

- Dissimilarity matrix:

$$\mathbf{S} = \begin{bmatrix} d(1,1) & \cdots & d(1,n) \\ \vdots & \ddots & \vdots \\ d(n,1) & \cdots & d(n,n) \end{bmatrix}$$



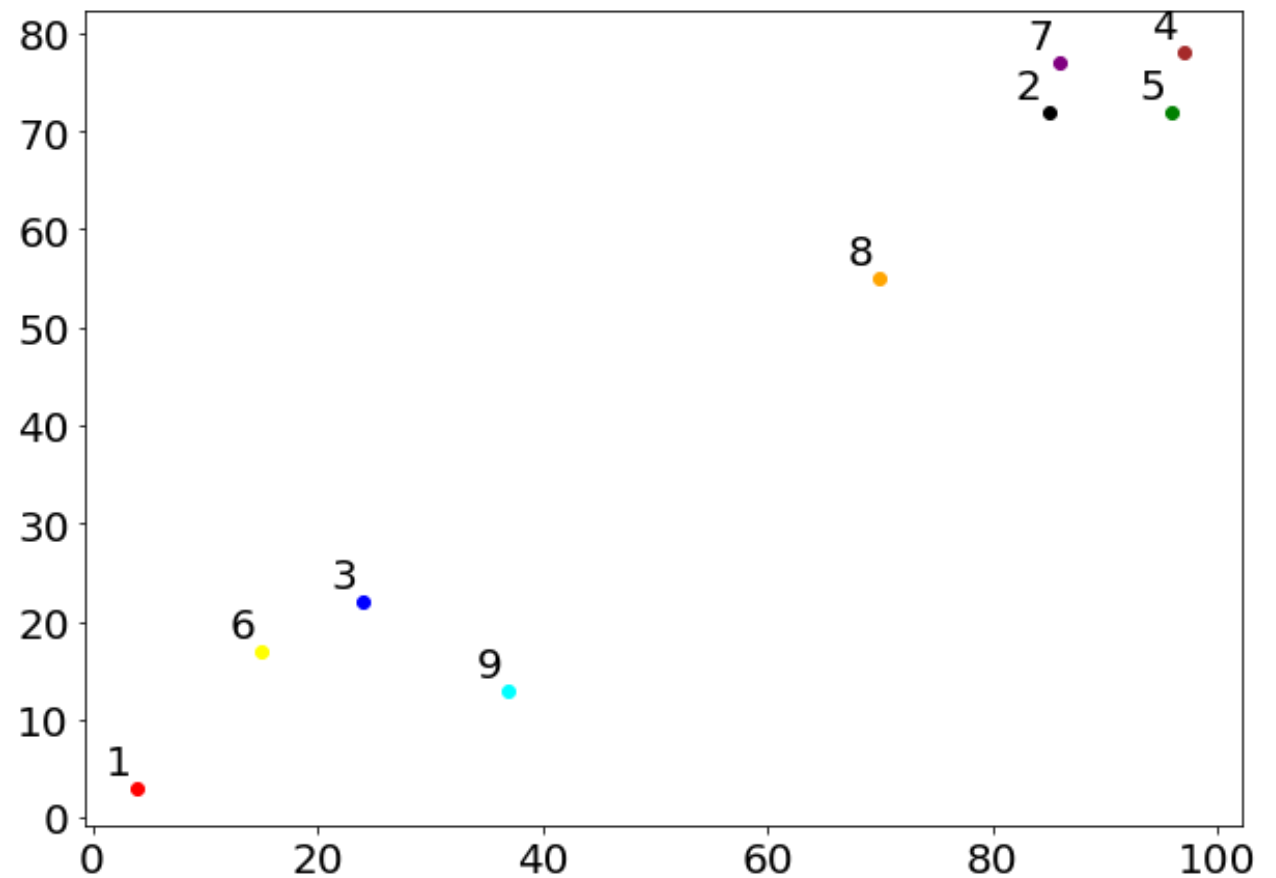
# Distance between clusters

- **Complete linkage:**
  - Maximum distance between clusters
- **Single linkage:**
  - Minimum distance between clusters
- **Average linkage:**
  - Average distance between clusters
- **Centroid linkage:**
  - Distance between centroids of clusters
- **Ward's linkage:**
  - Increase in sum of squares if two clusters are merged

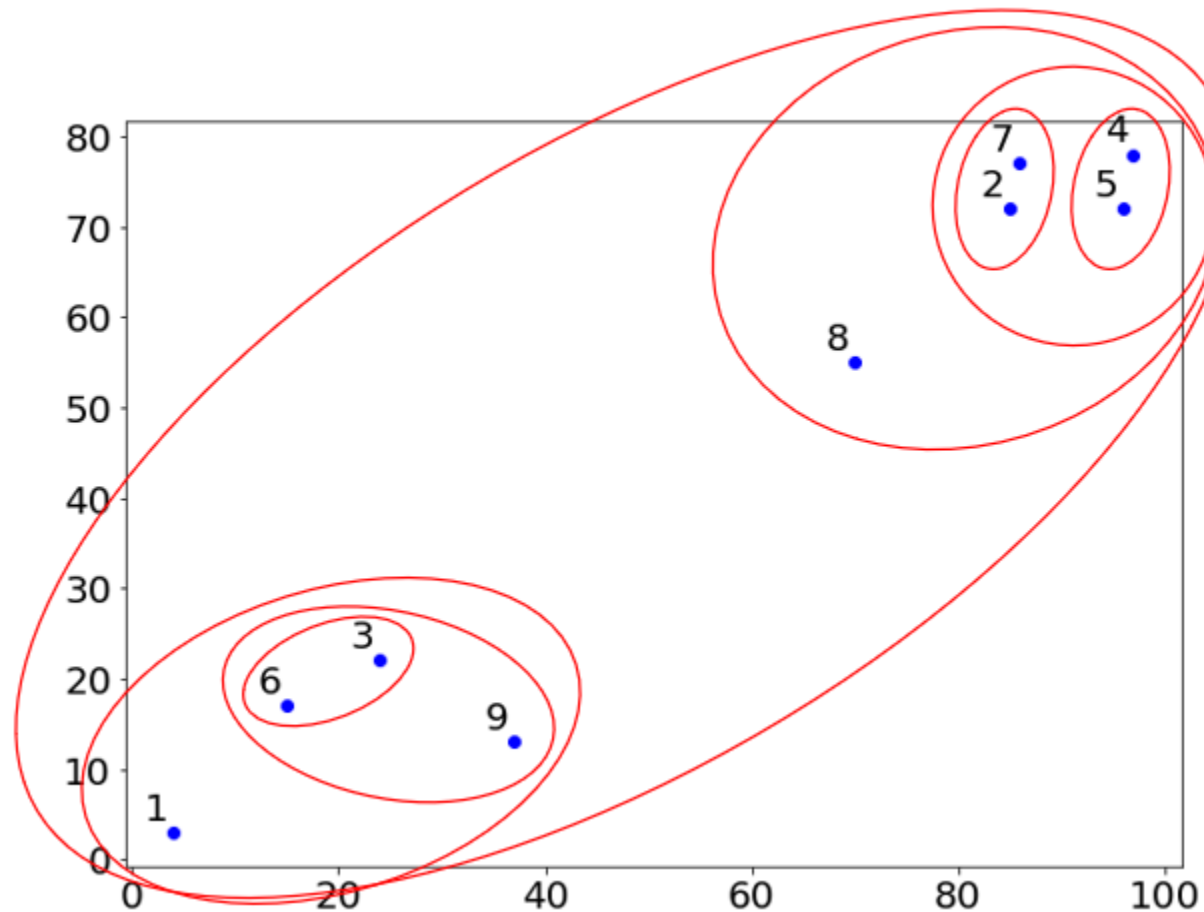
# Agglomerative Clustering Example

- Start from 9 clusters
- Complete linkage
- $9 \times 9$  dissimilarity matrix

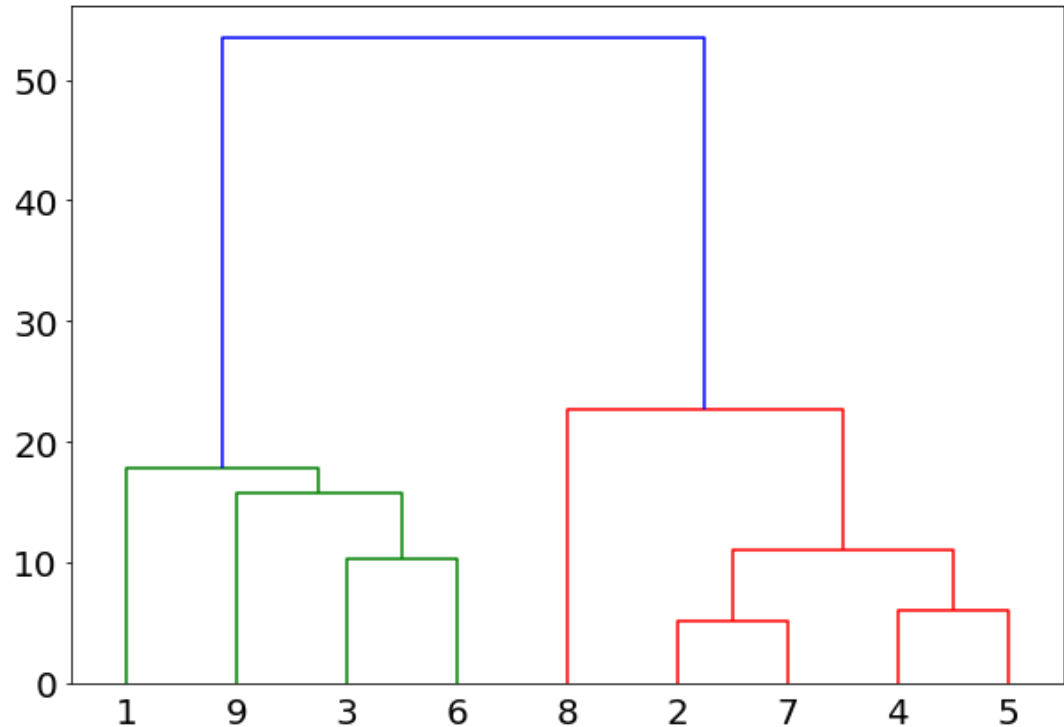
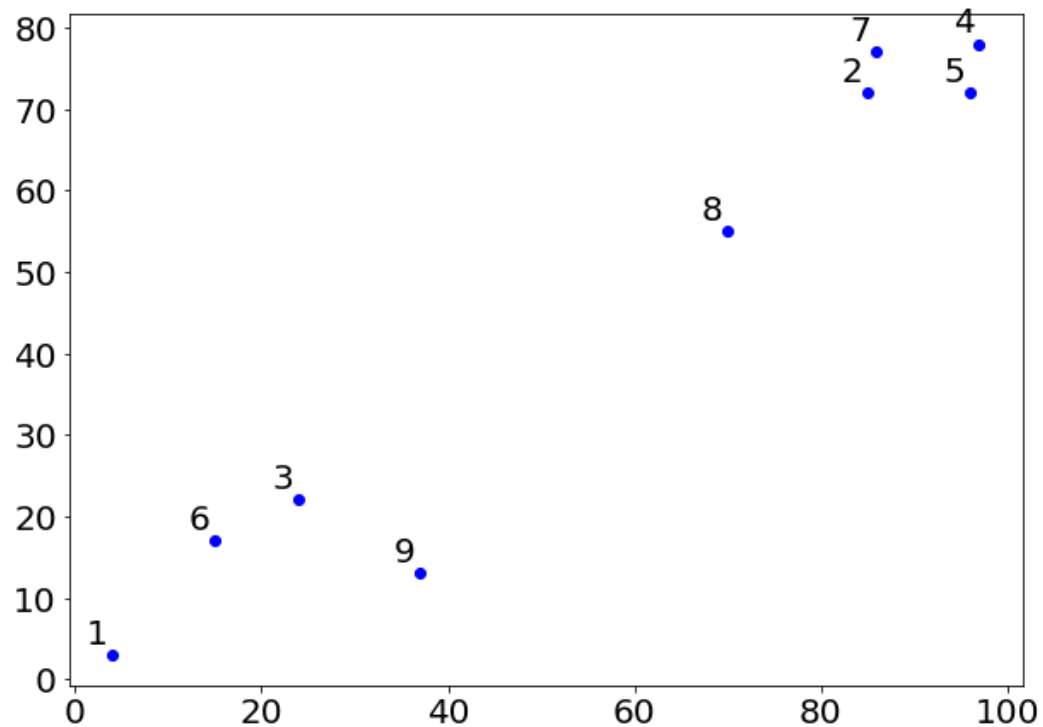
How many distances do we need to calculate? 81?



# Agglomerative Clustering Example

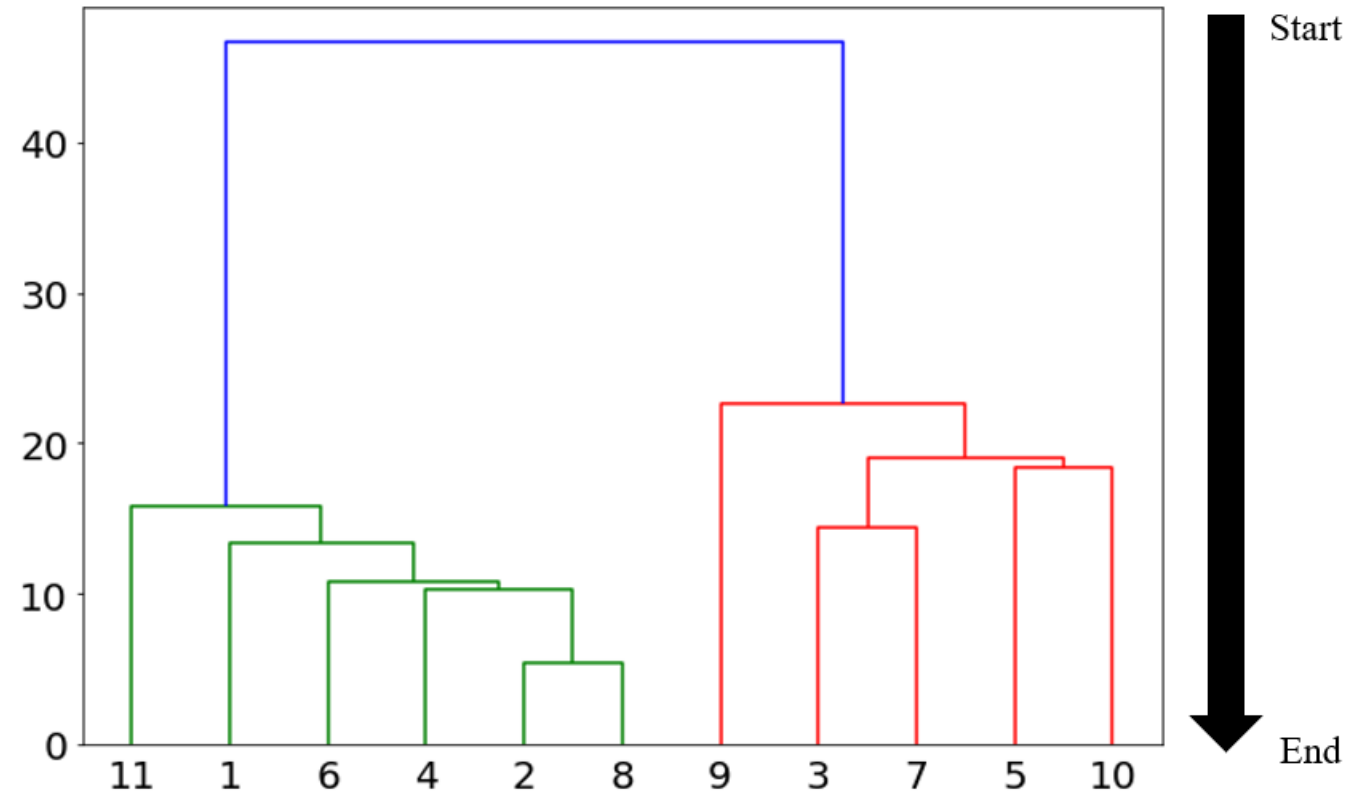


# Agglomerative Clustering Example



# Divisive Clustering (DIANA)

- Start with one cluster containing all  $n$  points.
- End up with  $n$  clusters containing one object.



# Divisive Clustering (DIANA) <sup>[1]</sup>

---

**Algorithm 2:** Divisive Analysis Clustering (DIANA)

---

**Input:**  $n$  data points

**Output:** final clustering result

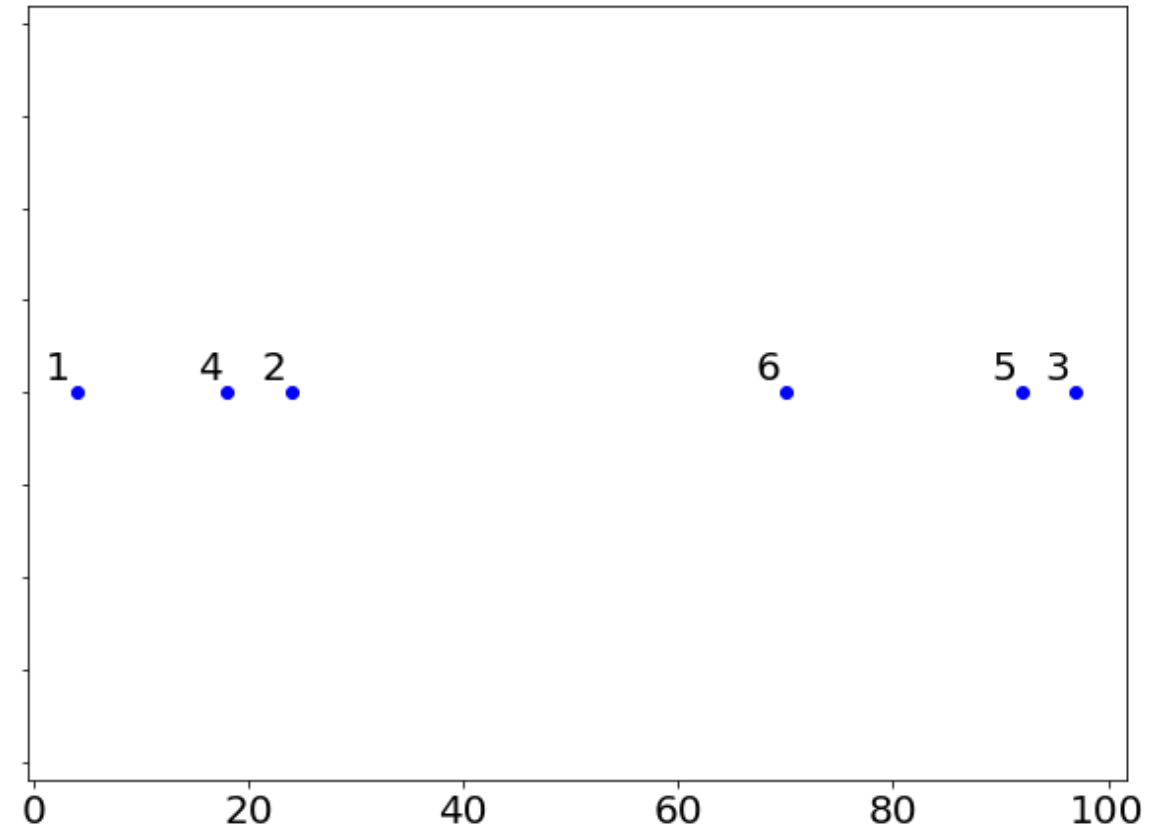
```
1 Initialize one cluster with all objects  $\mathbf{c}_1$ ;  
2 for the number of clusters  $k$  increases from 1 to  $n$  do  
3   Choose the cluster  $\mathbf{C}_i$  with the largest diameter value;  
4   Within  $\mathbf{C}_i$ , choose the object that has the maximum distance with the other objects as one  
   cluster and split this object as a splinter cluster;  
5   Update  $\mathbf{C}_i$ ;  
6   while True do  
7     for each data point  $j$  in  $\mathbf{C}_i$  do  
8       Calculate the distance  $d_1$  between the data  $j$  and the other objects in  $\mathbf{C}_i$  as one  
       cluster;  
9       Calculate the distance  $d_2$  between the data  $j$  and the splinter cluster;  
10      Calculate the difference  $\delta d_j = d_1 - d_2$ ;  
11    end for  
12    if  $\max \delta d_j$  is positive then  
13      Move the the data  $j$  with positive  $\delta d_j$  to the splinter cluster and update  $\mathbf{C}_i$ ;  
14    else  
15      break;  
16    end if  
17  end while  
18 end for
```

---

# Divisive Clustering Example

- Start from one cluster
- Complete linkage
- $6 \times 6$  dissimilarity matrix

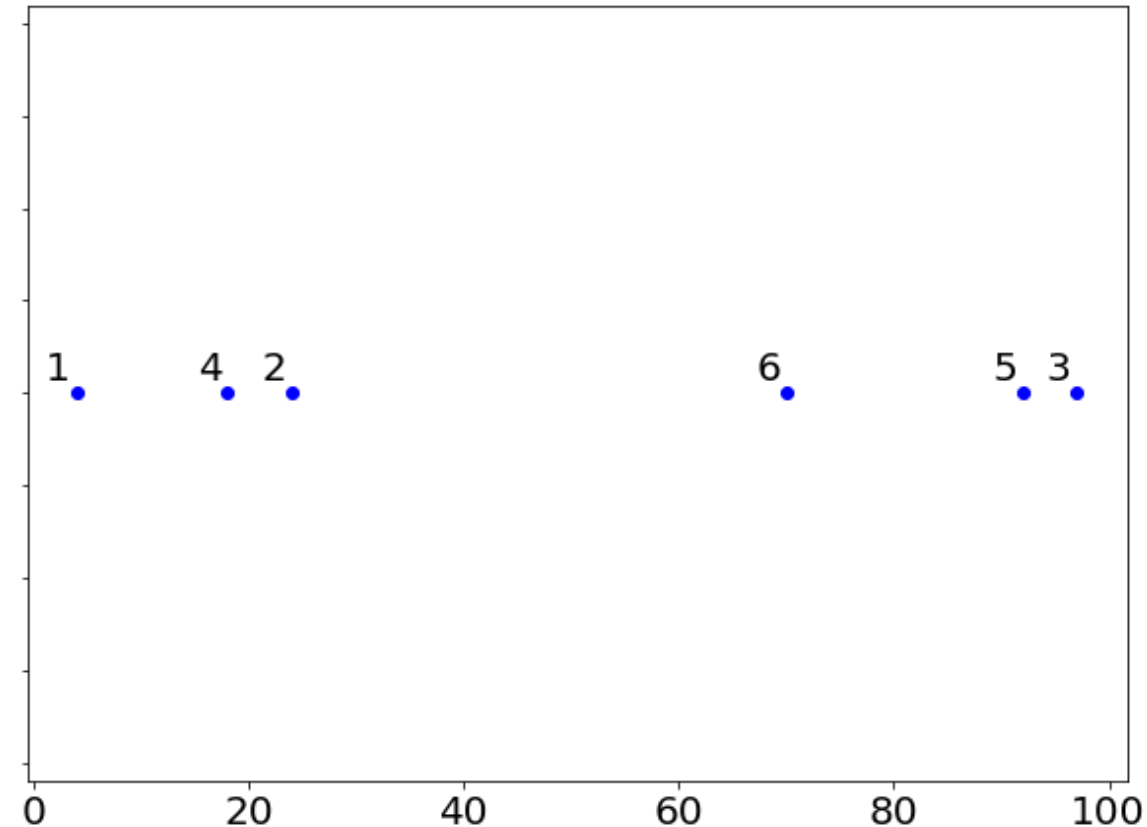
$$S = \begin{bmatrix} 0 & 20 & 93 & 14 & 88 & 66 \\ 20 & 0 & 73 & 6 & 68 & 46 \\ 93 & 73 & 0 & 79 & 5 & 27 \\ 14 & 6 & 79 & 0 & 74 & 52 \\ 88 & 68 & 5 & 74 & 0 & 22 \\ 66 & 46 & 27 & 52 & 22 & 0 \end{bmatrix}$$



# Divisive Clustering Example

- Calculate the distance between each point and the other objects

Point	Distance to other points
1	93
2	73
3	93
4	79
5	88
6	66

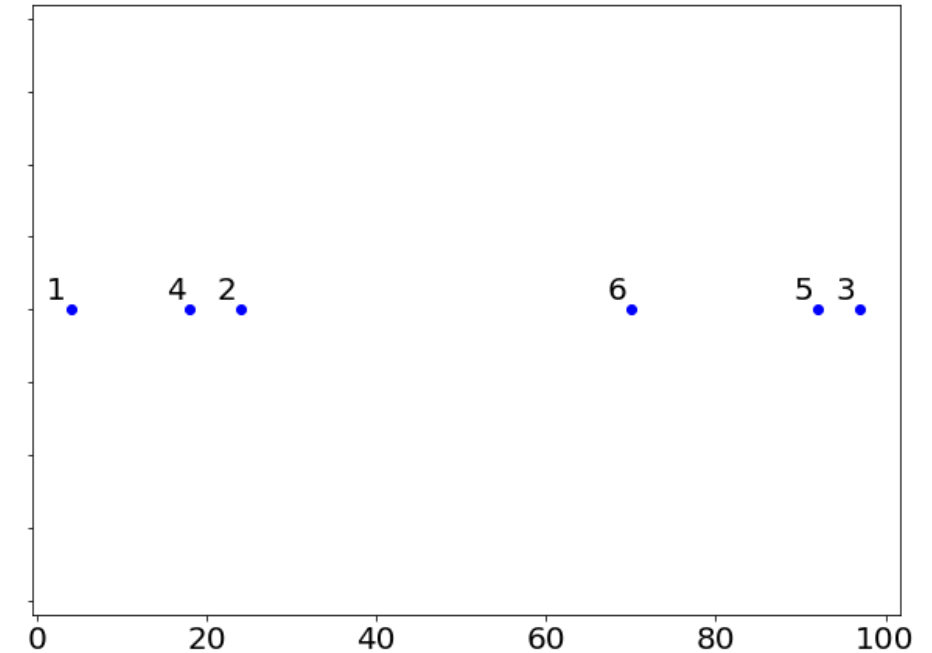


- Splinter cluster {1}



# Divisive Clustering Example

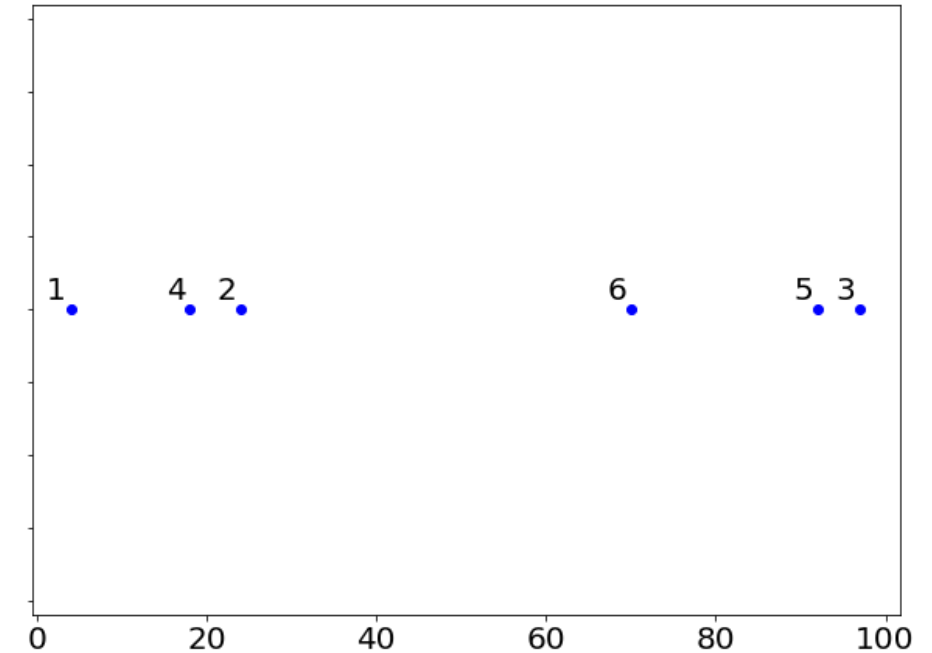
- Calculate the distance between each remaining point and the other objects
- Also the distance to the splinter cluster
- Splinter cluster {1,4}



Point	Distance to other points	Distance to the splinter cluster	Difference
2	73	20	53
3	79	93	-14
4	79	14	65
5	74	88	-14
6	52	66	-14

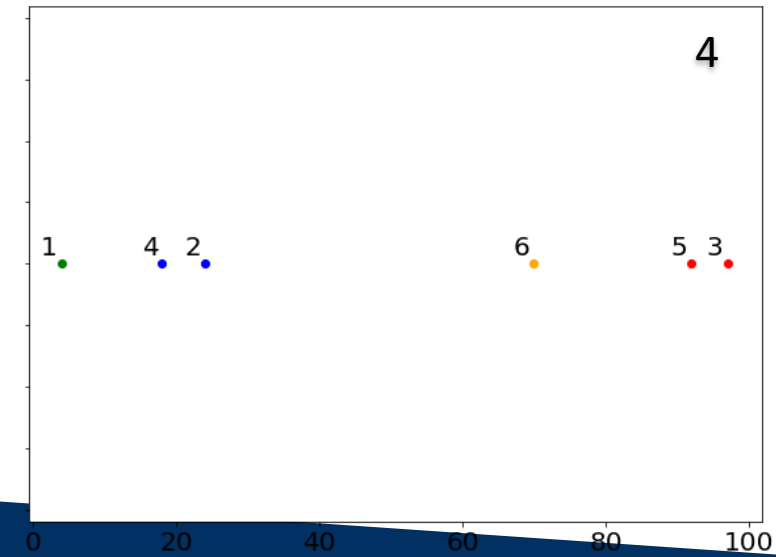
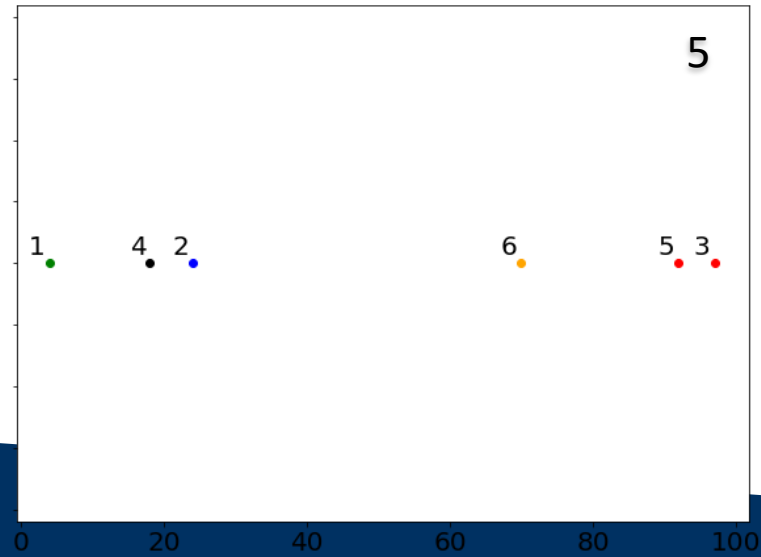
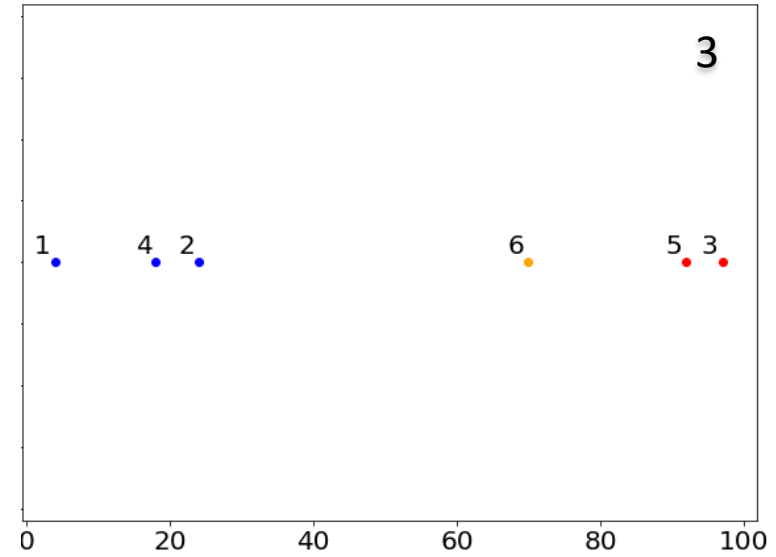
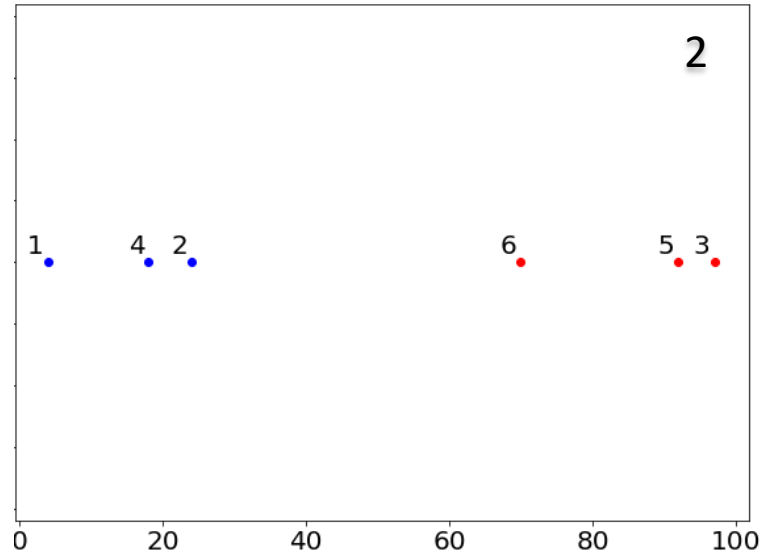
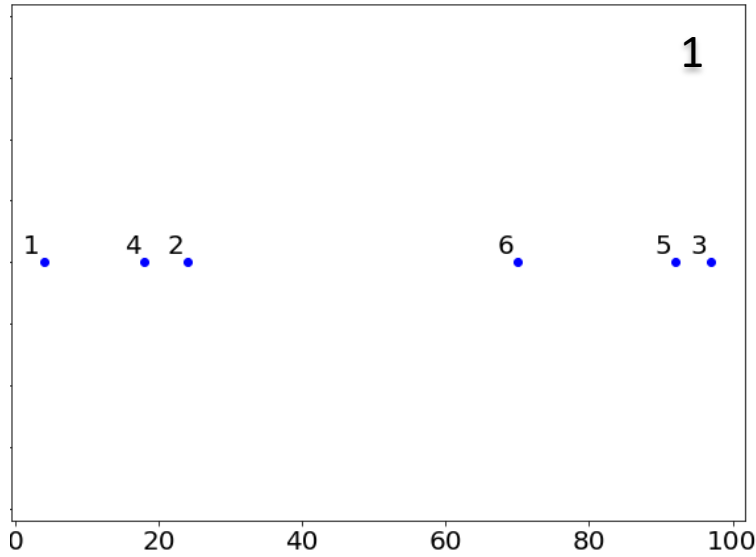
# Divisive Clustering Example

- Repeat the previous step
- Splinter cluster {1, 4, 2}
- {1,2,3,4,5,6} into {1,2,4} and {3,5,6}



Point	Distance to other points	Distance to the splinter cluster	Difference
2	73	20	53
3	73	93	-20
5	68	88	-20
6	46	66	-20

# Divisive Clustering (DIANA)

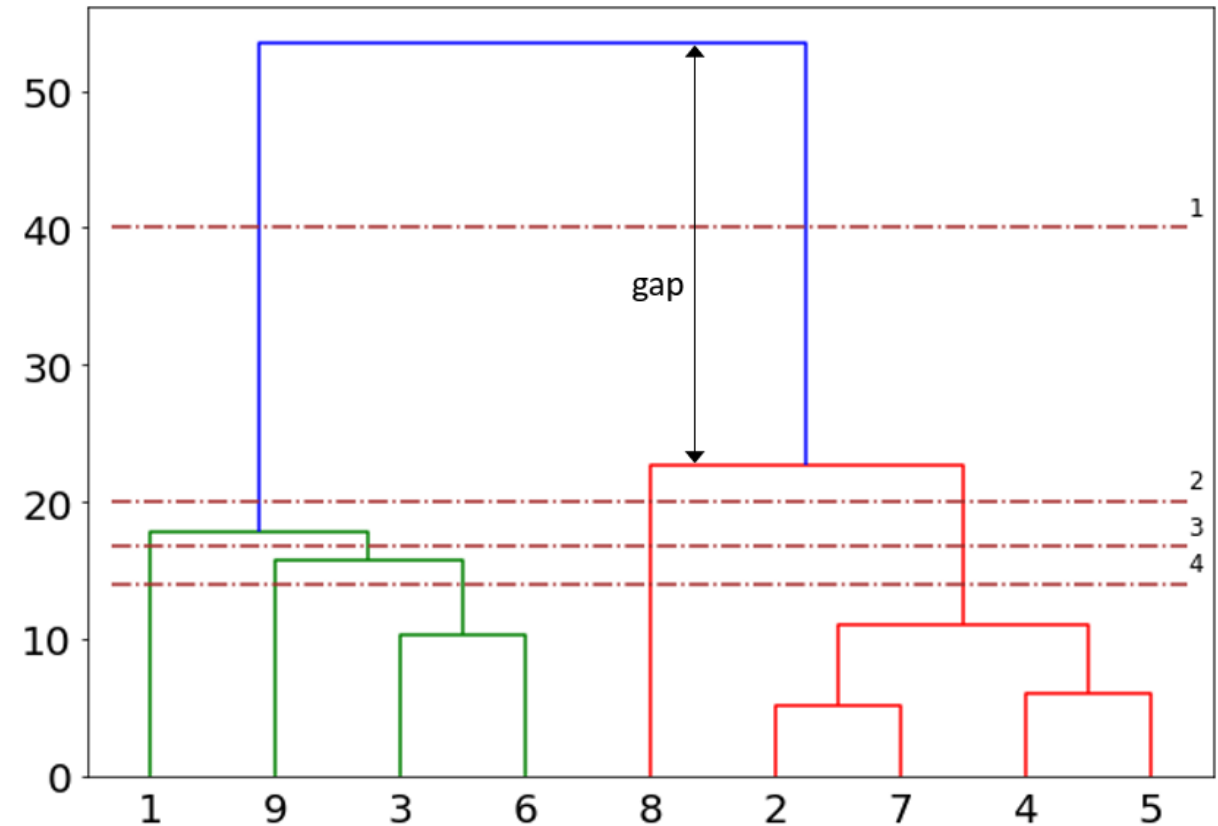


# Determination of $k$ (General)

- Elbow method <sup>[2]</sup>
  - Total within-cluster sum of square (WSS)
- Average silhouette method <sup>[3]</sup>
  - Average silhouette
- Gap statistic method
  - Gap statistic

# Determination of $k$ (from Dendrograom)

- Cut at different dissimilarity levels gives multiple values of  $k$
- Cut at the largest dissimilarity gap gives a roughly reasonable  $k$
- Affected by the linkage type since dissimilarity may change after each iteration.



# Specific Hierarchical Algorithms

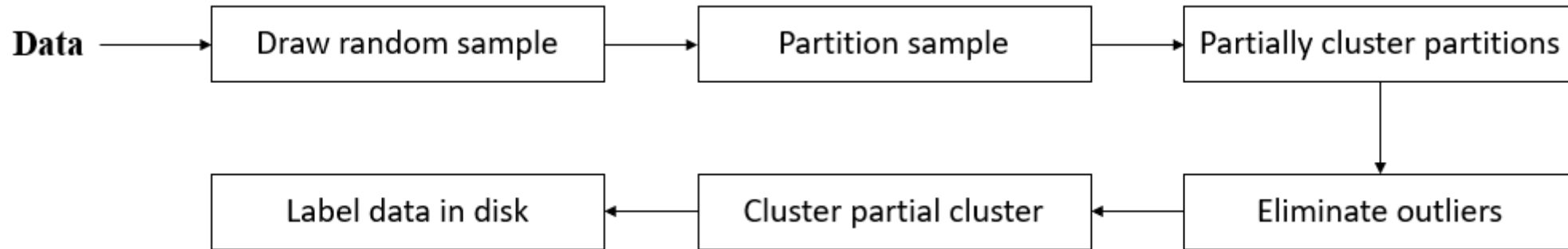
- Linkage algorithm
  - Single linkage, average linkage, complete linkage
- CURE (Clustering Using REpresentatives)
- BIRCH (Balanced Iterative Reducing and Clustering using Hierarchies) (Optional)

# Linkage algorithm [4]

- Single linkage:
  - Time complexity  $O[n^3]$  (simplest implementation)
  - Sensitive to outliers
- Complete linkage
  - Time complexity can be reduced to  $O[n^2 \log n]$
  - Cluster similar objects
- Average linkage
  - Compromise between single and complete
  - Often fails in complicated cluster shapes

# CURE (Clustering Using REpresentatives) [4]

- A hierarchical based clustering technique



- Representative points and shrinking factor
- Apply to outliers



# Reference

- [1]. Leonard Kaufman and Peter J Rousseeuw. *Finding groups in data: an introduction to cluster analysis*. Vol. 344. John Wiley & Sons, 2009.
- [2]. Bradley Boehmke Brandon Greenwell. *Hands-On Machine Learning with R*. Feb. 2020. URL: <https://bradleyboehmke.github.io/HOML/hierarchical.html#fig:dendrogram2>.
- [3]. Godfrey and Kate. *Determining The Optimal Number Of Clusters: 3 Must Know Methods*. Feb. 2020. URL: <https://bradleyboehmke.github.io/HOML/kmeans.html#eq:tot-within-ss>.
- [4]. M Kuchaki Rafsanjani, Z Asghari Varzaneh, and N Emami Chukanlo. “A survey of hierarchical clustering algorithms”. In: *The Journal of Mathematics and Computer Science* 5.3 (2012), pp. 229–240.