

# Clustering

..

# Supervised v/s unsupervised learning

Supervised Learning

$$y = f(x_1, x_2, x_3, x_4, \dots)$$

unsupervised Learning

$$\cancel{y} = f(x_1, x_2, x_3, x_4, \dots)$$

↓

$$(x_1, x_2, x_3, x_4, \dots)$$

# Unsupervised Learning

- Since there is no **Y** (response variable)
  - There is no possibility of **prediction**
- Goal in unsupervised learning
  - Discover aspects about the data and the variables
  - Mathematically **visualize** data
  - Are there sub-groups among the variables
- Type of unsupervised learning methods
  - PCA (Principal Component Analysis)
  - Clustering

# Challenges in unsupervised learning

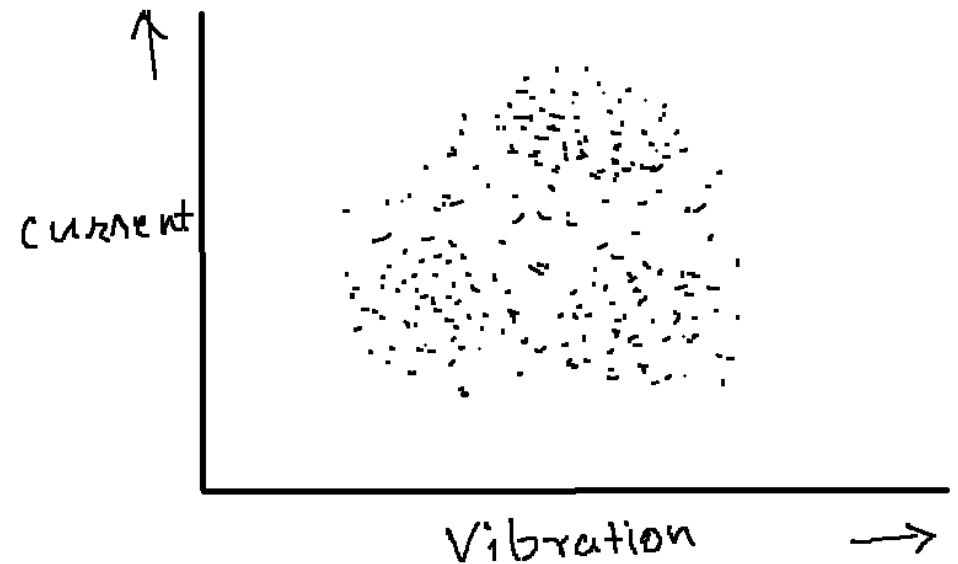
- It is a subjective process
- There is no criteria to objectively evaluate the results
  - No cross-validation methods
  - No possibility of validating results on independent data
- Consequently – unsupervised learning is performed as part of:
  - Exploratory Data Analysis
  - In order to discover **structure** in the data

# Examples of unsupervised learning

Measure the following

- Vibration
- Current

During metal cutting



# Use of unsupervised learning

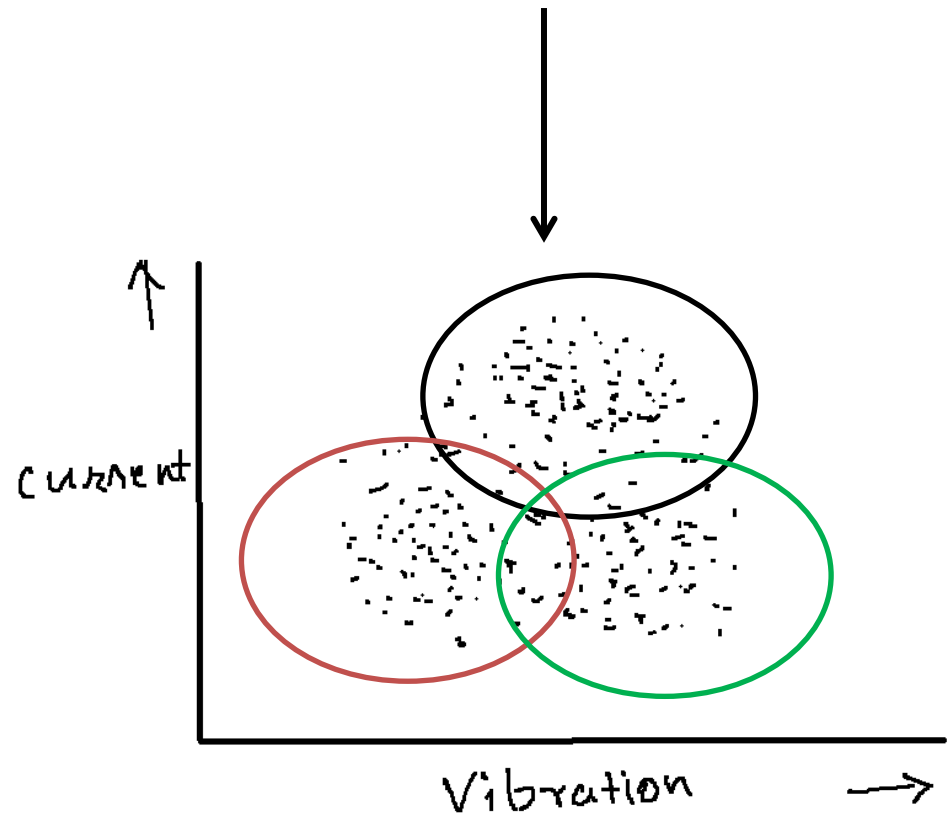
Measure the following

- Vibration
- Current

During metal cutting

The groups observed possibly indicate a behavioural pattern of the machine tool + cutting tool system

Is it possible to "discover" inherent groups?



# Clustering

- Clustering: Unsupervised learning method
- Used to “discover” inherent grouping of data points
- Popular clustering methods
  - K-means clustering
    - Partition the observations into a pre-specified number of clusters
  - Hierarchical clustering
    - Number of possible clusters is an outcome of the clustering process

# K-means Clustering

- Requires us to decide and specify, beforehand, the **desired number** of clusters
  - How do we decide?
  - Typically based on an understanding of the domain
- The method then results in assigning every point to one of these clusters
- Note:
  - Every data point will end up belonging to one and only one cluster



# K-means Clustering

- Approach
  - Specify the number of desired clusters
  - Algorithm begins by assigning input data points randomly to these clusters
  - The algorithm then iterates:
    - Derives the centroid of every cluster
    - Finds distances of all input points from the centroids of all the clusters
    - Reassigns every data point to the “nearest” cluster
    - This process is continued until there are no more reassignments

# K-means Clustering in Action

- Assume the following :
  - Input point set: [1 2 3 5 7 8 10 13 16]
    - Points along X axis ...
  - Desired number of clusters : 3
- Initial cluster assignment ... random

POINTS	[ 1	2	3	5	7	8	10	13	16 ]	
INIT ASSIGN	[1]	[2]	[3]	[1]	[2]	[3]	[1]	[2]	[3]	Initial random assignment of points to required number of clusters
CLUSTER CENTROIDS	c1	$(1+5+10)/3 = 5.33$								
	c2	$(2+7+13)/3 = 7.33$								
	c3	$(3+8+16)/3 = 9$								

# K-means Clustering in Action

- Distance calculation & cluster re-assignment

POINTS [ 1 2 3 5 7 8 10 13 16 ]

INIT  
ASSIGN [1] [2] [3] [1] [2] [3] [1] [2] [3]

CLUSTER  
CENTROIDS  
C1  $(1+5+10)/3 = 5.33$   
C2  $(2+7+13)/3 = 7.33$   
C3  $(3+8+16)/3 = 9$

DISTANCES FROM INITIAL CLUSTERS

PT	1	2	3	5	7	8	10	13	16
C1	<u>4.33</u>	<u>3.33</u>	<u>2.33</u>	<u>0.33</u>	1.67	2.67	4.67	7.67	10.67
C2	6.33	5.33	4.33	2.33	<u>0.33</u>	<u>0.67</u>	2.67	5.67	8.67
C3	8.00	7.00	6.00	4.00	2.00	1.00	<u>1.00</u>	<u>4.00</u>	<u>7.00</u>
NEW CLUST	1	1	1	1	2	2	3	3	3

# K-means Clustering in Action

- Cluster re-assignment ... step 2

PT	1	2	3	5	7	8	10	13	16
C1	<u>4.33</u>	<u>3.33</u>	<u>2.33</u>	<u>0.33</u>	1.67	2.67	4.67	7.67	10.67
C2	6.33	5.33	4.33	2.33	<u>0.33</u>	<u>0.67</u>	2.67	5.67	8.67
C3	8.00	7.00	6.00	4.00	2.00	1.00	<u>1.00</u>	<u>4.00</u>	<u>7.00</u>
NEW CLUST	1	1	1	1	2	2	3	3	3

CENTROIDS:  $C1 = (1+2+3+5)/4 = 2.75$

$C2 = (7+8)/2 = 7.50$

$C3 = (10+13+16)/3 = 13.00$

DISTANCES FROM NEW CLUSTERS.

PT	1	2	3	5	7	8	10	13	16
C1	<u>1.75</u>	<u>0.75</u>	<u>0.25</u>	<u>2.25</u>	4.25	5.25	7.25	10.25	13.25
C2	6.50	5.50	4.50	2.50	<u>0.5</u>	<u>0.50</u>	<u>2.50</u>	5.5	8.50
C3	12.00	11.00	10.00	8.00	6.0	5.0	3.0	<u>0.0</u>	<u>3.0</u>

# K-means Clustering in Action

- Cluster re-assignment ... Step 3

PT	1	2	3	5	7	8	10	13	16
C1	<u>1.75</u>	<u>0.75</u>	<u>0.25</u>	<u>2.25</u>	4.25	5.25	7.25	10.25	13.25
C2	6.50	5.50	4.50	2.50	<u>0.5</u>	<u>0.50</u>	<u>2.50</u>	5.5	8.50
C3	12.00	11.00	10.00	8.00	6.0	5.0	3.0	<u>0.0</u>	<u>3.0</u>

CENTROIDS

$$C1 = (1+2+3+5)/4 = 2.75$$
$$C2 = (7+8+10)/3 = 8.33$$
$$C3 = (13+16)/2 = 14.5$$

No change in  
clusters  
assignments.  
Algorithm stops.

DISTANCES FROM NEW CLUSTERS.

PT	1	2	3	5	7	8	10	13	16
C1	<u>1.75</u>	<u>0.75</u>	<u>0.25</u>	<u>2.25</u>	4.25	5.25	7.25	10.25	13.25
C2	7.33	6.33	5.33	3.33	<u>1.33</u>	<u>0.33</u>	<u>1.67</u>	4.67	7.67
C3	13.5	12.5	11.5	9.50	7.50	6.5	4.50	<u>1.50</u>	<u>1.50</u>



# K-means Clustering: Final Clusters

- Input points
  - [1 2 3 5 7 8 10 13 16]
- Clusters
  - [1 2 3 5] [7 8 10] [13 16]
- Note:
  - In K-means clustering, the initial cluster assignments are random
  - Therefore, the resulting clusters could represent only a local minima
  - Given a data set, it is therefore important to run K-means clustering algorithm multiple times – each starting from a distinct initial random cluster assignment

# Formally ...

The  $K$ -means clustering procedure results from a simple and intuitive mathematical problem. We begin by defining some notation. Let  $C_1, \dots, C_K$  denote sets containing the indices of the observations in each cluster. These sets satisfy two properties:

1.  $C_1 \cup C_2 \cup \dots \cup C_K = \{1, \dots, n\}$ . In other words, each observation belongs to at least one of the  $K$  clusters.
2.  $C_k \cap C_{k'} = \emptyset$  for all  $k \neq k'$ . In other words, the clusters are non-overlapping: no observation belongs to more than one cluster.

Good clustering is one for which the **within cluster** variation is as less as possible

Source: *Introduction to Statistical Learning*; Auth: James, Witten, Hastie, Tibshirani; Springer

# Formally ...

Using Euclidean distance, the total within cluster variation can be expressed as:

$$W(C_k) = \frac{1}{|C_k|} \sum_{i, i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2$$

$|C_k|$  denotes the number of observations in the  $k$ th cluster

The clustering problem therefore reduces to:

$$\text{minimize}_{C_1, \dots, C_K} \left\{ \sum_{k=1}^K \frac{1}{|C_k|} \sum_{i, i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 \right\}$$

The k-means algorithm is based on this problem formulation  $\frac{1}{|C_k|} \sum_{i, i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 = 2 \sum_{i \in C_k} \sum_{j=1}^p (x_{ij} - \bar{x}_{kj})^2$

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# Formally ...

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# Hierarchical Clustering

- Hierarchical clustering does not require specify the number of clusters
- It results in a hierarchical, tree-like, representation of the observations
  - Known as a ***Dendrogram***
  - (*Based on **bottom-up** or **agglomerative** clustering*)

# Hierarchical Clustering: Method

- Given a point set, the Hierarchical Clustering algorithm proceeds as follows:
  - Every data point is assigned to its own unique cluster
  - The algorithm then iterates as follows:
    - A distance metric (Euclidean – by default) is used to find out pair-wise distance between all the clusters (**see next slide**)
    - A pair of clusters are found such that the distance between them is the smallest of all pair-wise distances
    - The two clusters are merged
    - The process is repeated till only one cluster remains
  - The above process results in a Dendrogram which, when cut at a particular level, results in a specific number of clusters
    - The required number of clusters can thus be obtained

# Hierarchical Clustering: Distance between Clusters

- Distance between clusters is evaluated using one of the following “linkages”: of these, **Average and Complete are used the most**

<i>Linkage</i>	<i>Description</i>
Complete	Maximal intercluster dissimilarity. Compute all pairwise dissimilarities between the observations in cluster A and the observations in cluster B, and record the <i>largest</i> of these dissimilarities.
Single	Minimal intercluster dissimilarity. Compute all pairwise dissimilarities between the observations in cluster A and the observations in cluster B, and record the <i>smallest</i> of these dissimilarities. Single linkage can result in extended, trailing clusters in which single observations are fused one-at-a-time.
Average	Mean intercluster dissimilarity. Compute all pairwise dissimilarities between the observations in cluster A and the observations in cluster B, and record the <i>average</i> of these dissimilarities.
Centroid	Dissimilarity between the centroid for cluster A (a mean vector of length $p$ ) and the centroid for cluster B. Centroid linkage can result in undesirable <i>inversions</i> .

# Hierarchical Clustering : Example

- Input point set: [1 2 3 5 7 8 10 13 16]
- Initial cluster: [1] [2] [3] [5] [7] [8] [10] [13] [16]
- Inter-cluster distance: Using “Complete Linkage”

	1	2	3	5	7	8	10	13	16
1	-	1	2	4	6	7	9	12	15
2	-	-	1	3	5	6	8	11	14
3	-	-	-	2	3	5	7	10	13
5	-	-	-	-	2	3	5	8	11
7	-	-	-	-	-	1	3	6	9
8	-	-	-	-	-	-	2	5	8
10	-	-	-	-	-	-	-	3	6
13	-	-	-	-	-	-	-	-	3
16	-	-	-	-	-	-	-	-	-

# Hierarchical Clustering: Example

- Step 1

	1	2	3	5	7	8	10	13	16
1	-	1	2	4	6	7	9	12	15
2	-	-	1	3	5	6	8	11	14
3	-	-	-	2	3	5	7	10	13
5	-	-	-	-	2	3	5	8	11
7	-	-	-	-	-	1	3	6	9
8	-	-	-	-	-	-	2	5	8
10	-	-	-	-	-	-	-	3	6
13	-	-	-	-	-	-	-	-	3
16	-	-	-	-	-	-	-	-	-

# Hierarchical Clustering: Example

- Step 2

	[1 2]	3	5	7	8	10	13	16
[1 2]	-	2	4	6	7	9	12	15
3	-	-	2	4	5	7	10	13
5	-	-	-	2	3	5	8	11
7	-	-	-	-	1	3	6	9
8	-	-	-	-	-	2	5	8
10	-	-	-	-	-	-	3	6
13	-	-	-	-	-	-	-	3
16	-	-	-	-	-	-	-	-

# Hierarchical Clustering

- Step 3

	[1 2]	3	5	[7 8]	10	13	16
[1 2]	-	2	4	7	9	12	15
3	-	-	2	5	7	10	13
5	-	-	-	3	5	8	11
[7 8]	-	-	-	-	3	6	9
10	-	-	-	-	-	3	6
13	-	-	-	-	-	-	3
16	-	-	-	-	-	-	-



# Hierarchical Clustering

- Step 3

	[1 2 3]	5	[7 8]	10	13	16
[1 2 3]	-	4	7	9	12	15
5	-	-	3	5	8	11
[7 8]	-	-	-	3	6	9
10	-	-	-	-	3	6
13	-	-	-	-	-	3
16	-	-	-	-	-	-

# Hierarchical Clustering

- Step 4

	[1 2 3]	[5 7 8]	10	13	16
[1 2 3]	-	7	9	12	15
[5 7 8]	-	-	5	8	11
10	-	-	-	3	6
13	-	-	-	-	3
16	-	-	-	-	-

- Step 5

	[1 2 3]	[5 7 8]	[10 13]	16
[1 2 3]	-	7	12	15
[5 7 8]	-	-	8	11
[10 13]	-	-	-	6
16	-	-	-	-

# Hierarchical Clustering

- Step 6

	[1 2 3]	[5 7 8]	[10 13]	16
[1 2 3]	-	7	12	15
[5 7 8]	-	-	8	11
[10 13]	-	-	-	6
16	-	-	-	-

- Step 7

	[1 2 3]	[5 7 8]	[10 13 16]
[1 2 3]	-	7	15
[5 7 8]	-	-	11
[10 13 16]	-	-	-

# Hierarchical Clustering

- Step 8

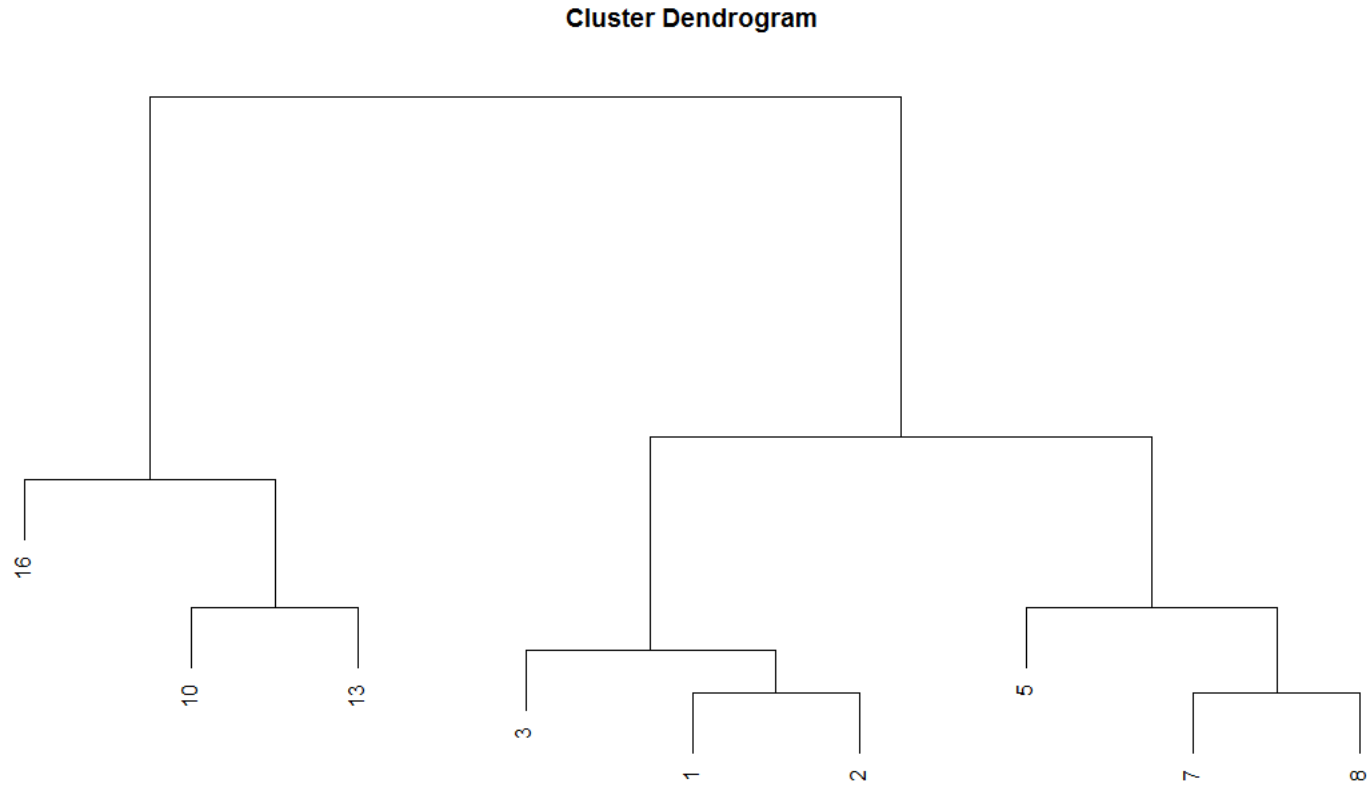
	[1 2 3 5 7 8]	[10 13 16]
[1 2 3 5 7 8]	-	15
[10 13 16]	-	-

- Step 9: Final

	[1 2 3 5 7 8 10 13 16]
[1 2 3 5 7 8 10 13 16]	-

# Hierarchical Clustering

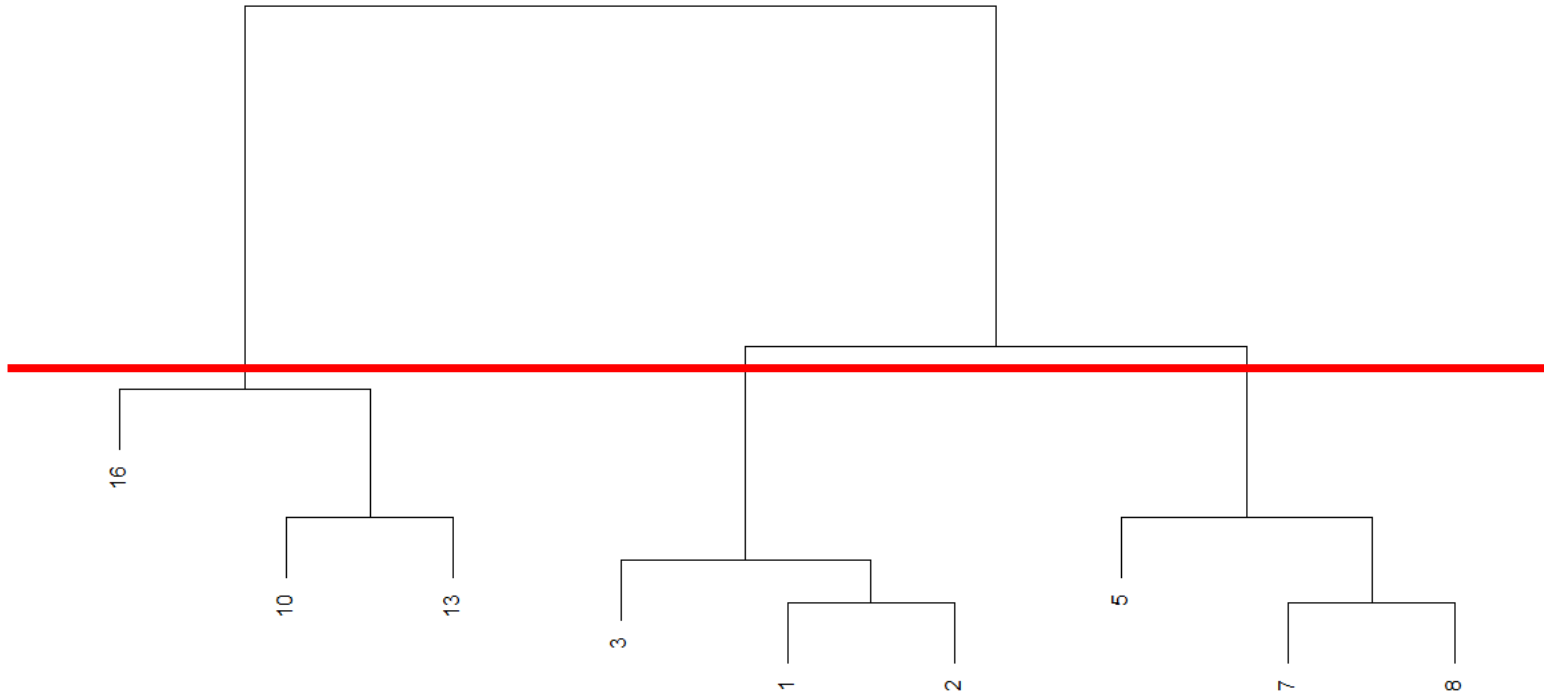
- Resulting Dendrogram



# Hierarchical Clustering

- Cutting the dendrogram to get clusters

Cluster Dendrogram

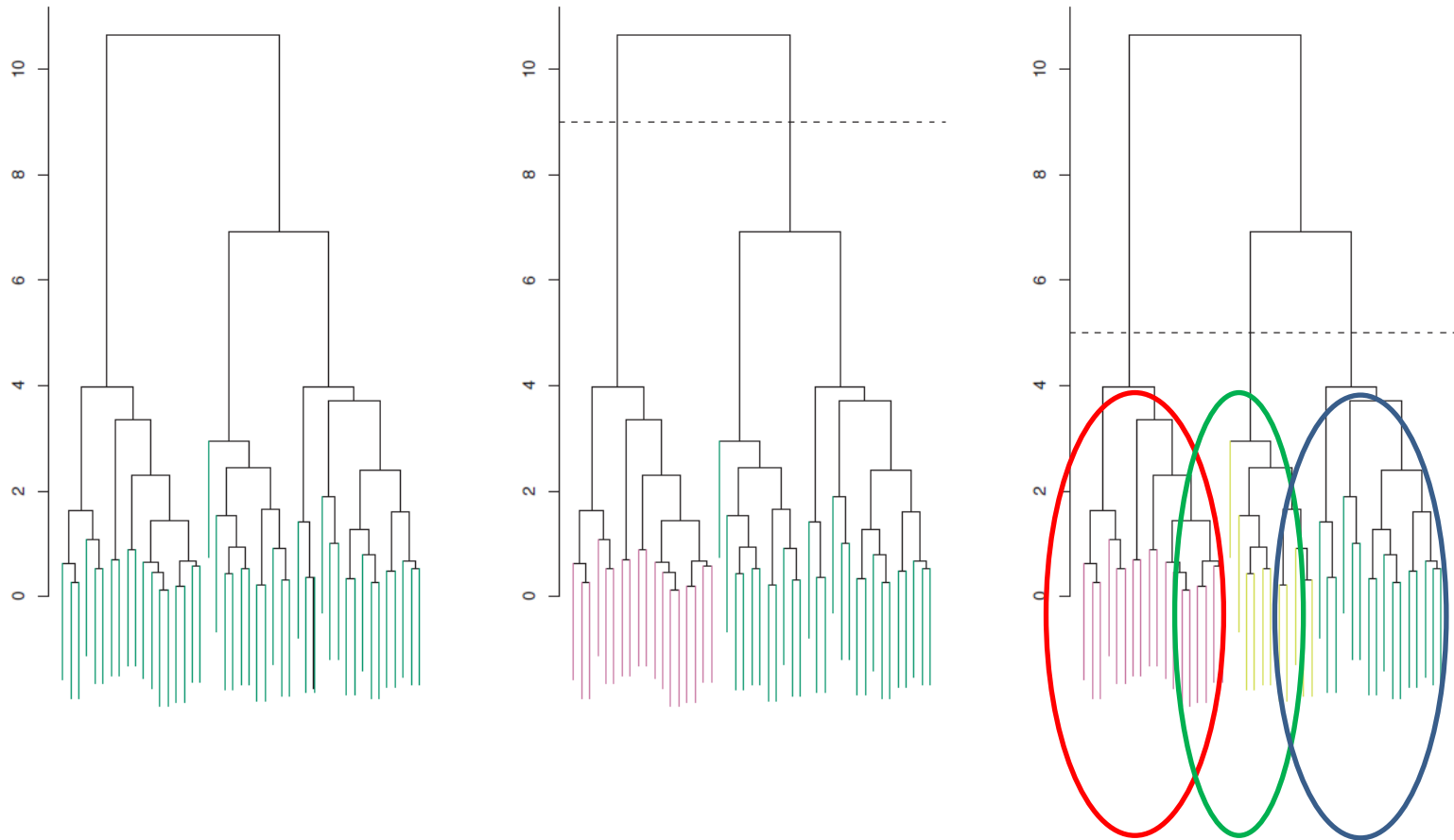


- Resulting clusters: [1 2 3] [5 7 8] [10 13 16]

# Interpreting the Dendrogram

- No conclusions can be drawn
  - Based on the proximity of branches along the horizontal axis
- Observations that fuse lower in the tree
  - Close to each other
- Observations that fuse higher in the tree
  - Are dissimilar
- Height of the fusion
  - Indicate how different the branches / leaves are
- Identifying distinct number of groups
  - Based on the leg lengths

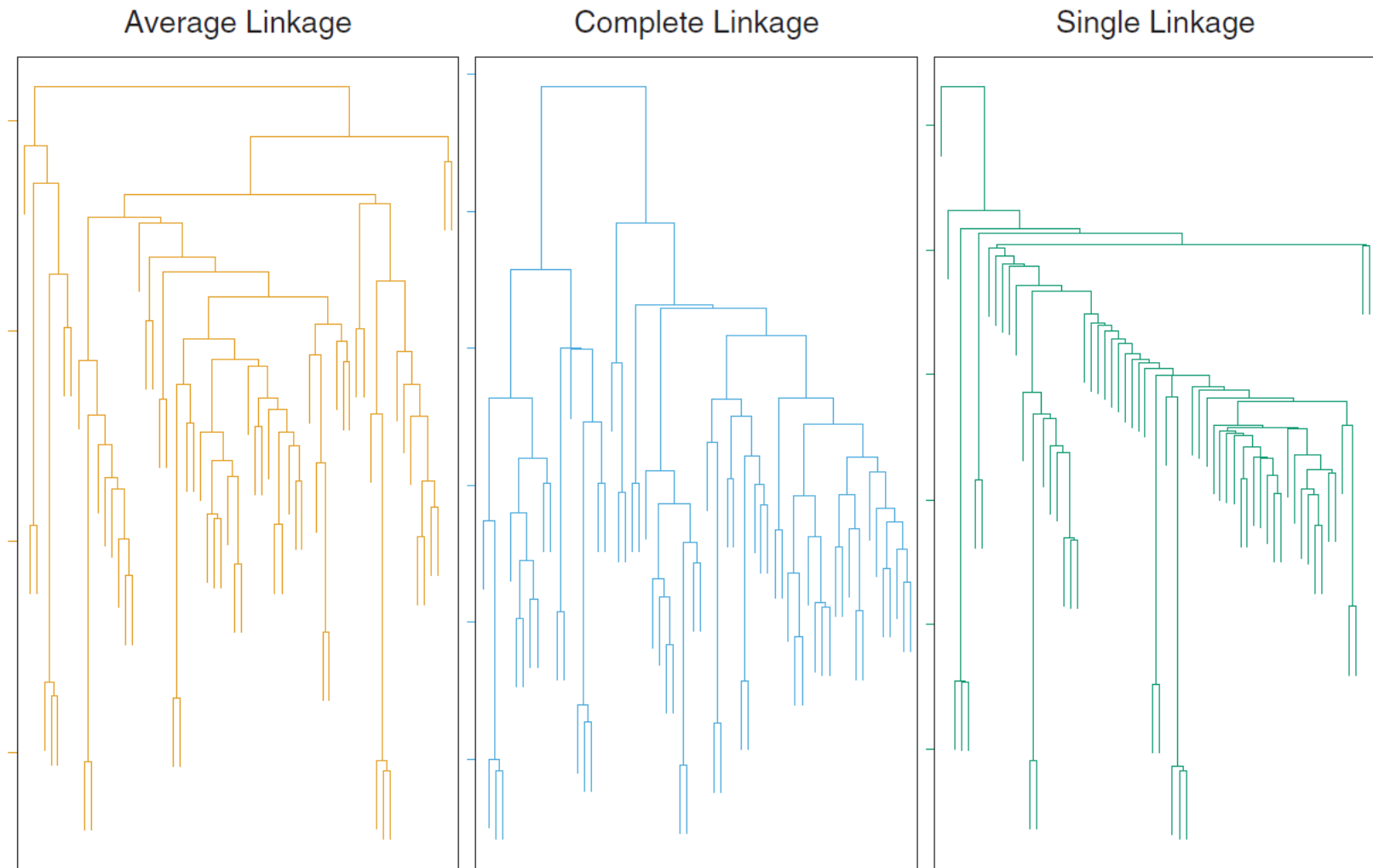
# Interpreting the Dendrogram



Source: *Introduction to Statistical Learning*; Auth: James, Witten, Hastie, Tibshirani; Springer



# Comparison between linkages



Source: *Introduction to Statistical Learning*; Auth: James, Witten, Hastie, Tibshirani; Springer

# Points to be considered in clustering

- Which dissimilarity measure (linkage)?
- How many clusters?
- Presence of outliers
  - Since every point is forced into a cluster
  - This may lead to distortions
- Observations at different scales
  - How to handle such observations?
  - Should they be centered and scaled?

# Clustering performance

