# Artificial Intelligence: Semi-supervised Learning

#### Do you remember this slide? Naïve Bayes Example 3

$$P(H_{1} | E_{2}) = \frac{P(H_{1}) \times P(E_{2} | H_{1})}{P(E_{2})} = \frac{.2x.2}{.31} = .129$$

$$P(H_{2} | E_{2}) = \frac{P(H_{2}) \times P(E_{2} | H_{2})}{P(E_{2})} = \frac{.5x.3}{.31} = .484$$

$$P(H_{3} | E_{2}) = \frac{P(H_{3}) \times P(E_{2} | H_{3})}{P(E_{2})} = \frac{.3x.4}{.31} = .387$$

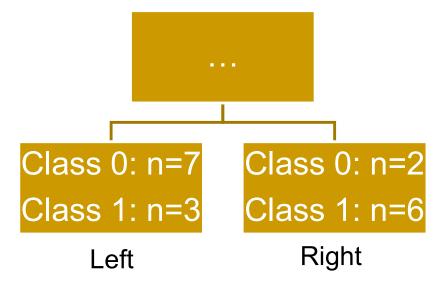
 $\mathbb{R}H_2$  is the most likely hypothesis, given the evidence  $P(H_2 \mid E_2)$  is the highest

Tomorrow the weather will be bad

$$H_{NB} = \underset{H_i}{\operatorname{argmax}} \frac{P(H_i) \times P(E|H_i)}{P(E)}$$

# How about this slide? (Decision Tree classification error)

What would be the classification error in these leaves?



- In the left leaf, the prediction (predicted label) is class 0
- The predicted probability of class 1 is 0.3
- We are 30% confident that label should be class 1 (thus 70% confident of being class 0)
- In the right leaf, we are 75% confident of predicting class 1

#### Prediction Probability

- Most machine learning classifiers not only returns (predicts) the class label but also a probability indicating the confidence
- Usually, the class prediction is based on this probability applying a cut point
- Cut point is by default 0.5 for binary classification, but is adjusted based on the importance of performance metrics (recall, precision, specificity, ...)

```
Probability that this observation belongs to class 1 \equiv p (y=1) p (y=1) > 0.5 => Classifier predicts class 1 p (y=1) <= 0.5 => Classifier predicts class 0
```

#### Why semi-supervised learning?

- Unlabeled data is cheap and available
- Labeled data can be hard to get
- Labelling (data annotation) can be very expensive
  - tedious task and time-consuming
  - may need experts' intervention
  - error-prone
- Motivation: Using both labeled and unlabeled data for learning

(\$17-\$36/hr) from companies with openings that are

HIRING) May 2023

niting now. Find job postings near you and 1-click applyt

#### Semi-supervised learning: Self training

- n<sub>I</sub> labelled samples (x, f(x))
- n<sub>n</sub> unlabeled samples (x,)
- Usually, n<sub>I</sub> << n<sub>n</sub>
- A classifier (learner) x -> f(x)

**Assumption**: Classifier's high confidence predictions are correct.

- Train on labelled samples
- Predict on unlabeled samples
- Add  $(x, \hat{y})$ 's with high confidence to the labelled samples
- Repeat

#### Self training semi-supervised

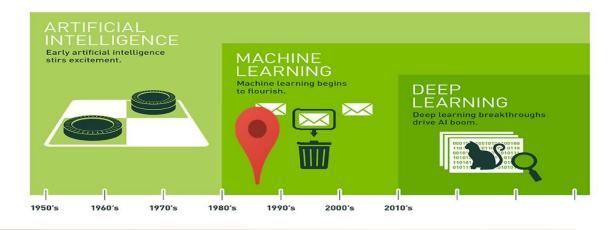
- A simple semi-supervised learning method.
- Iteration using existing classifiers
- Mistakes can be costly especially in the first iterations
  - Add only very confident samples to the labelled set
  - Add samples using the confidence as weight
  - Unable a sample if the confidence drops

• ...

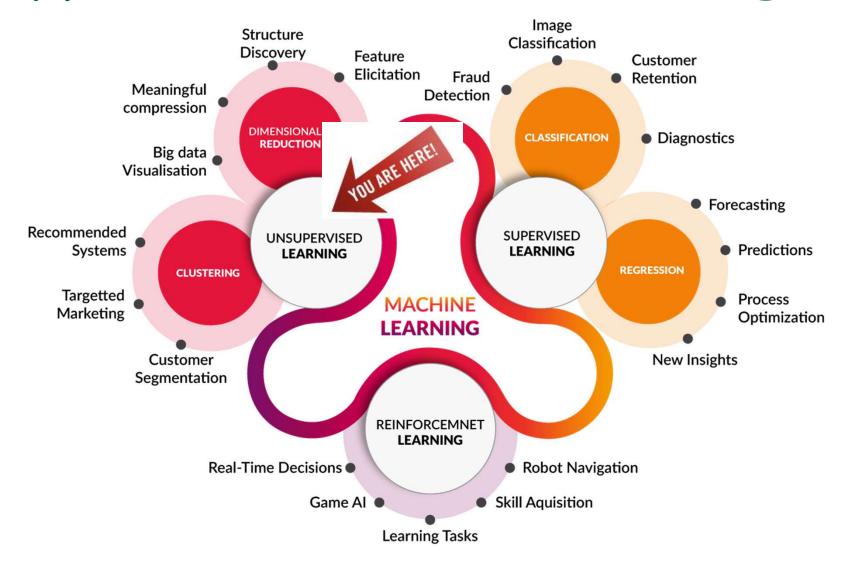
# Artificial Intelligence: Unsupervised Learning

# Today

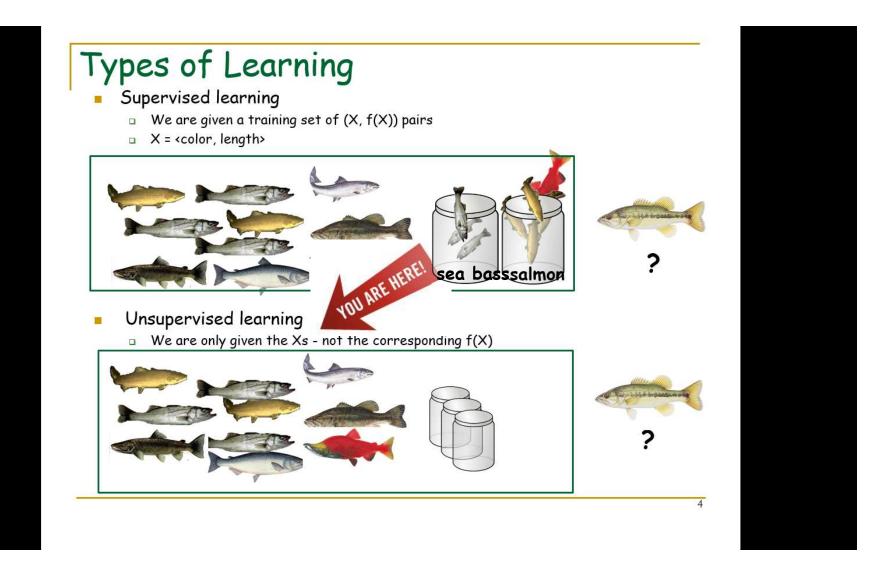
- YOU ARE HERE!
- Unsupervised Learning
- 2. k-means Clustering
- 3. Hierarchical Clustering



# Types of Machine Learning



#### Remember this slide?



### Unsupervised Learning



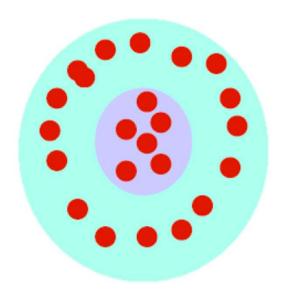
- Learn without labeled examples
  - □ i.e. X is given, but not f(X)

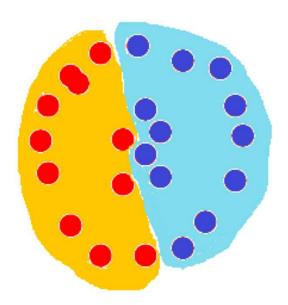
	small nose	big teeth	small eyes	moustache	f(X) = ?
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- Without a f(X), you can't really identify/label a test instance
- But you can:
  - Cluster/group the features of the test data into a number of groups
  - Discriminate between these groups without actually labeling them

# What is Clustering

- The organization of unlabeled data into similarity groups called clusters.
- A cluster is a collection of data items which are "similar" between them, and "dissimilar" to data items in other clusters.





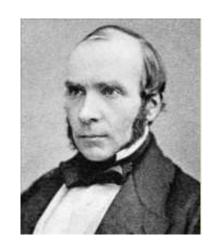
# Applications of Clustering

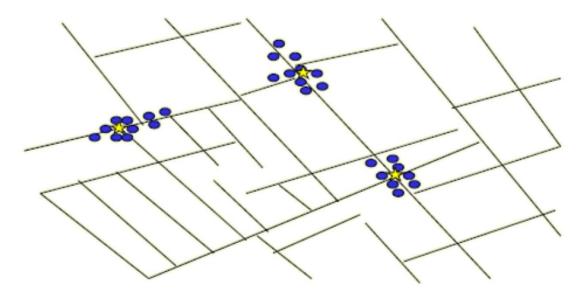
- Exploratory data analysis (EDA)
- Customer segmentation in marketing to identify similar groups of customers based on their purchase behavior
- Image segmentation in computer vision to group pixels with similar attributes for object recognition
- Document clustering in natural language processing (NLP)

...

# Historic Application of Clustering

- John Snow, a London physician plotted the location of cholera on a map during an outbreak in the 1850s.
- The locations indicated that cases were clustered arounds certain intersections where there were polluted wells - thus exposing both the problem and the solution.





FROM: Nina Mishra HP Labs

# Clustering

Represent each instance as a vector  $\langle a_1, a_2, a_3, ..., a_n \rangle$ 

Each vector can be visually represented in an n-dimensional

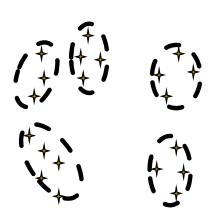
space

_	$X_5$		
	) X <sub>2</sub>		
		X4 ×	
	-,,,,- -,,,,,,,,,-	-	

	$a_1$	a <sub>2</sub>	$a_3$	Output
$X_1$	1	0	0	?
X <sub>2</sub>	1	6	0	?
<b>X</b> <sub>3</sub>	8	0	1	3
X <sub>4</sub>	6	1	0	?
X <sub>5</sub>	1	7	1	?

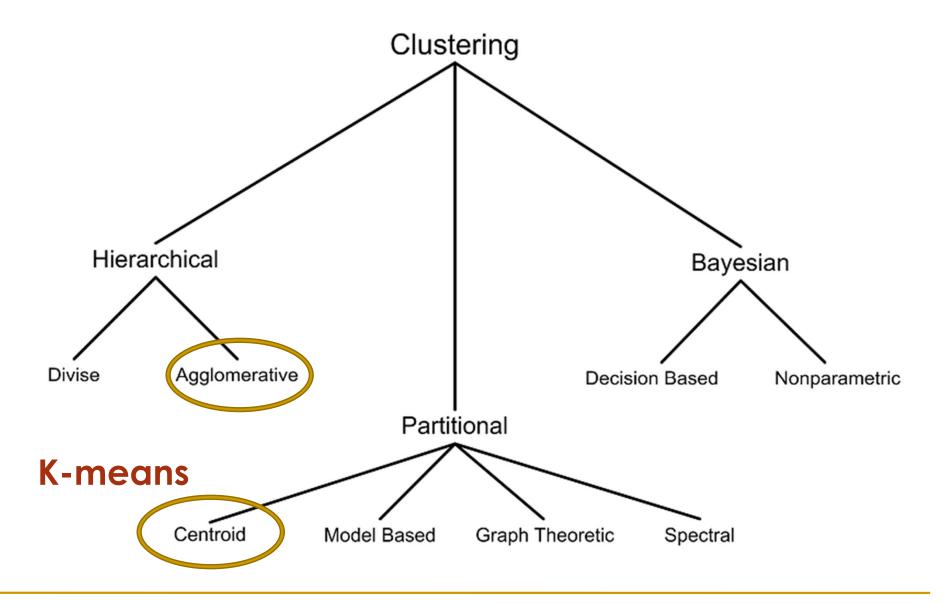
# Clustering

Clustering algorithm



- Represent test instances on a n dimensional space
- Partition them into regions of high density
  - How? ... many algorithms (ex. k-means)
- Compute the centroid of each region as the average of data points in the cluster

# Clustering Techniques



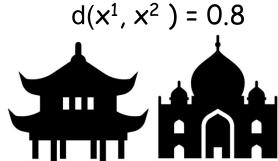
# k-means Clustering

- User selects how many clusters they want... (the value of k)
- 1. Place k points into the space (ex. at random). These points represent initial group centroids.
- 2. Assign each data point  $x_n$  to the nearest centroïd.
- 3. When all data points have been assigned, recalculate the positions of the K centroïds as the average of the cluster (new centroïds)
- 4. Repeat Steps 2 and 3 until none of the data instances change group.

# Nearest centroid!! But how to define similarity (distance)?

1. For two objects  $x^1$ ,  $x^2$  distance  $d(x^1, x^2)$  is a numerical representation of their dissimilarity.





- 2. Several ways to define it in Machine Learning.
  - Euclidean distance
  - Manhattan distance

#### Euclidean Distance

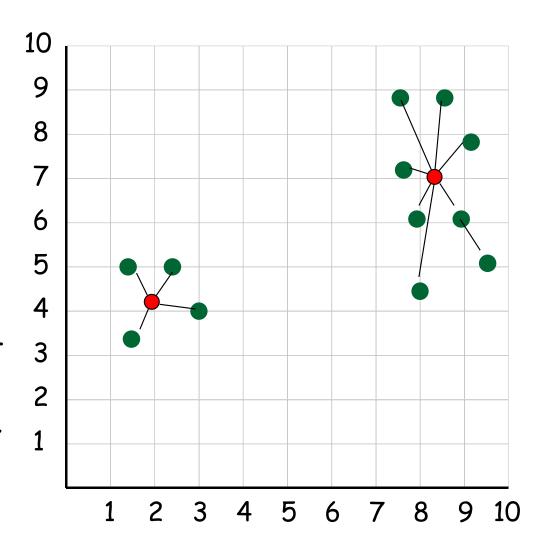
- To find the nearest centroid...
- a possible metric is the Euclidean distance
- distance between 2 pts

$$p = (p_1, p_2, ...., p_n)$$

$$q = (q_1, q_2, ...., q_n)$$

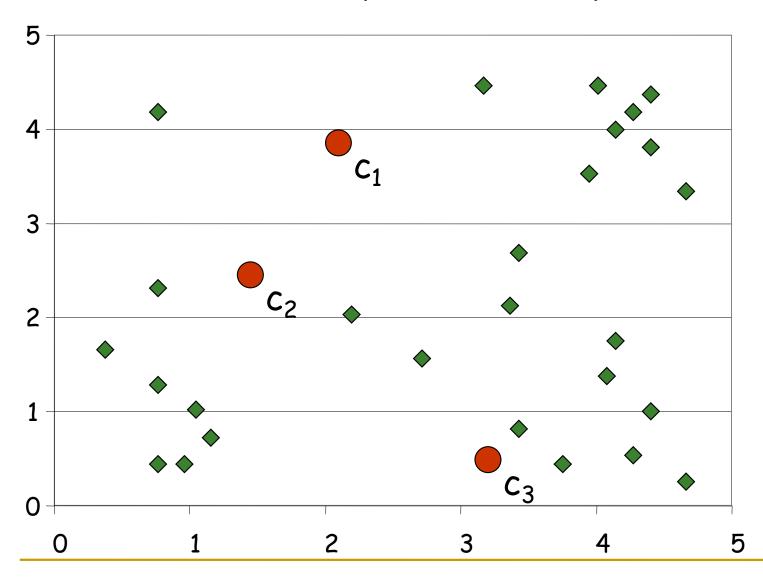
$$d = \sqrt{\sum_{i=1}^{n} (p_i - q_i)^2}$$

- where to assign a data point x?
- For all k clusters, chose the one where x has the smallest distance



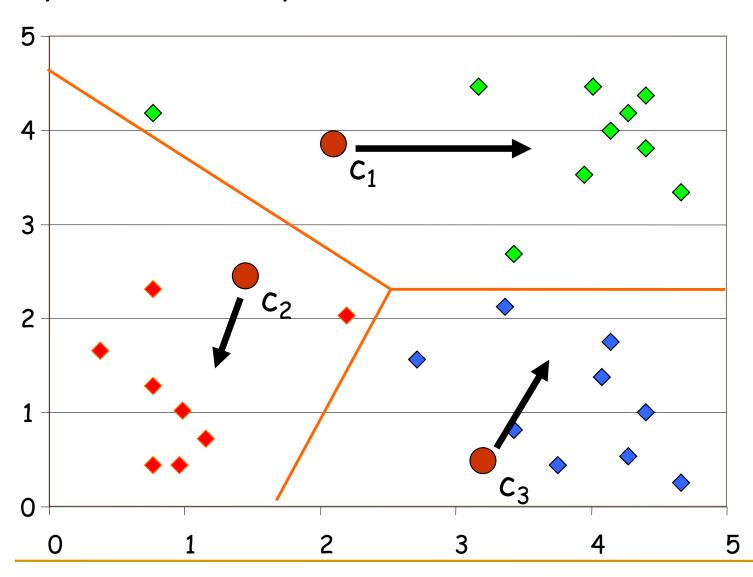
# Example (in 2-D... i.e. 2 features)

initial 3 centroïds (ex. at random)

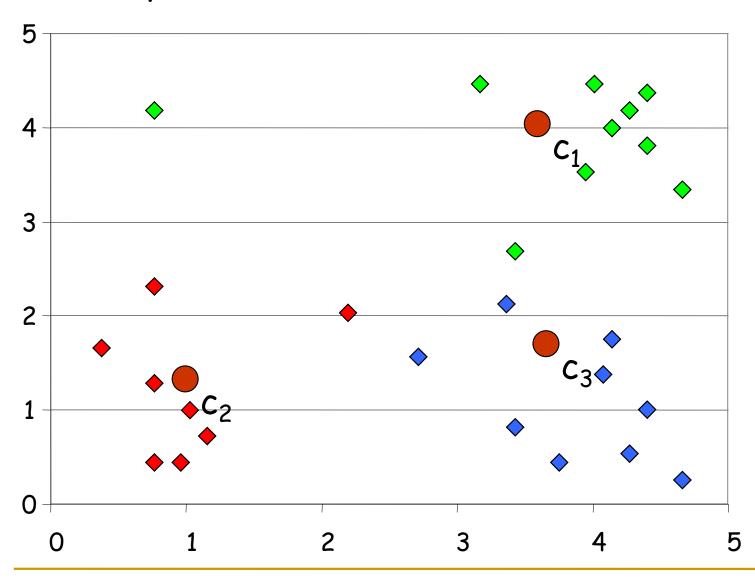


# Example

partition data points to closest centroïd

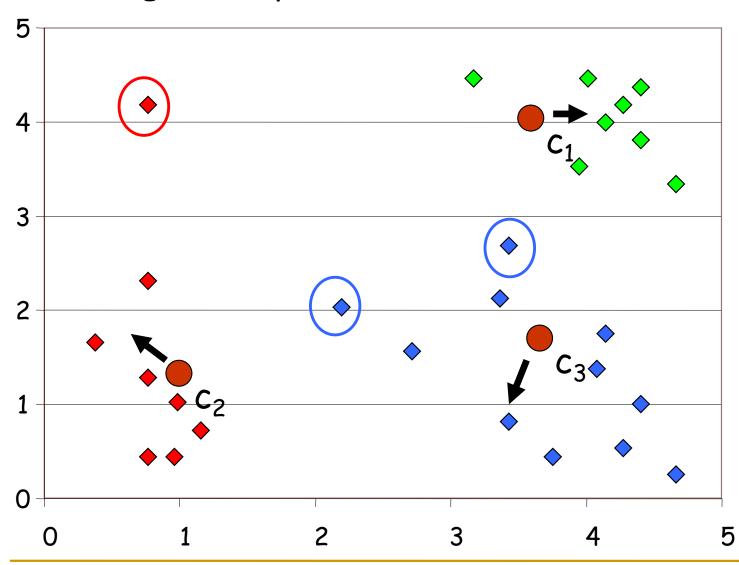


# Example re-compute new centroïds

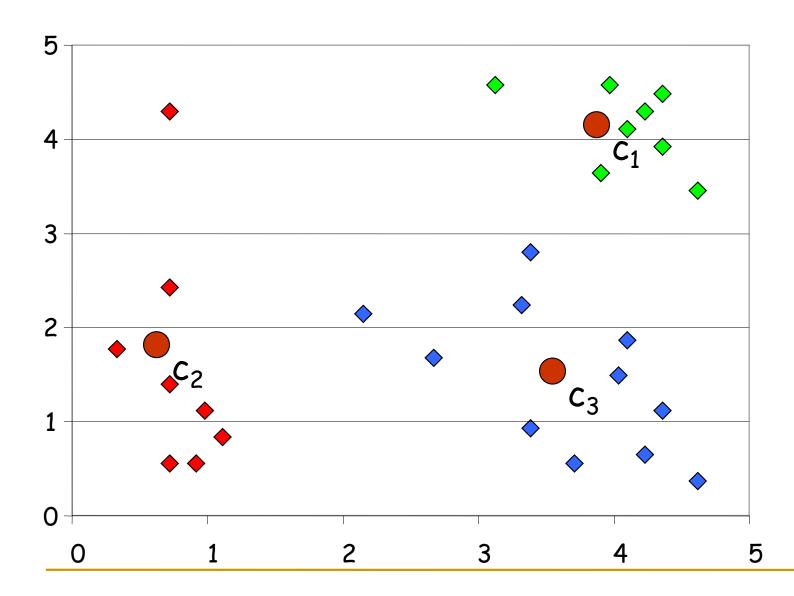


# Example

re-assign data points to new closest centroids



# Example



#### Notes on k-means

- converges very fast!
- BUT:
  - very sensitive to initial choice of centroids
    - many find useless clusters...
  - user must set initial k
    - not easy to do...
- many other clustering algorithms...

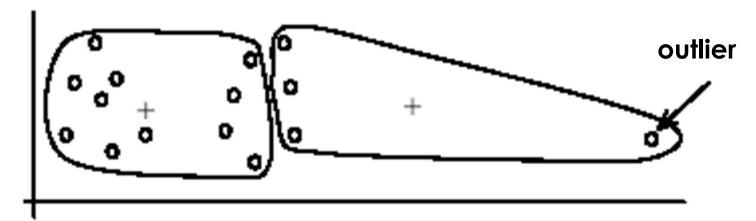
### Why use k-means?

- Strengths:
  - Simple
    - Easy to understand and implement
  - Efficient: Time complexity O(t·k·n)
    - n number of data points
    - k number of clusters
    - t number of iterations
  - With small k and t, linear performance on practical problems

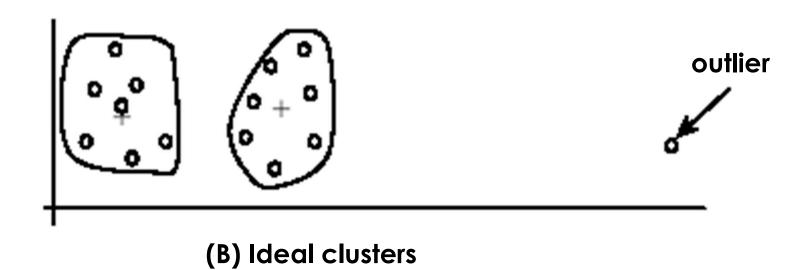
#### Weakness of k-means

- User needs to specify k
- Algorithm is sensitive to outliers
  - i.e., data points that are far away from others
  - Could be errors in the data or special data points with very different characteristics

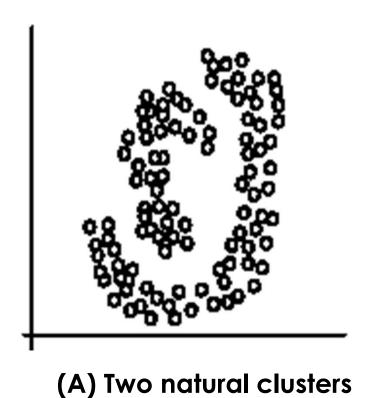
#### Outliers



(A) Undesirable clusters

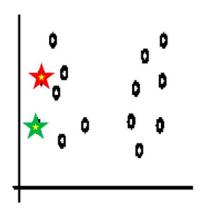


# Special data structures

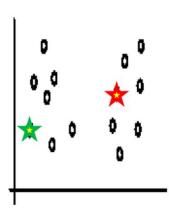


(B) k-means clusters

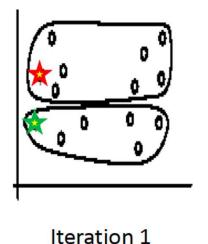
# Sensitivity to initial seeds

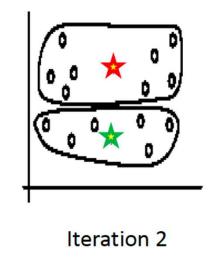


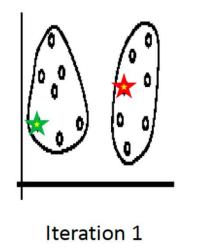
Random selection of seeds (centroids)

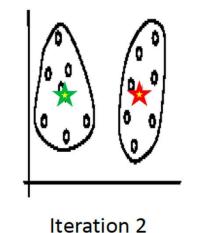


Random selection of seeds (centroids)









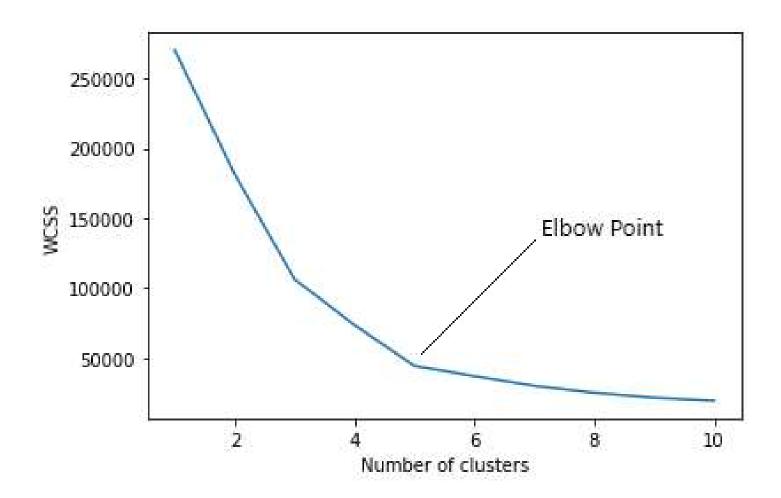
# How to decide the number of clusters (K)

- In some contexts, it may be given
  - Classification of clients into Platinum,
     Gold, and Silver
- Finding the "Optimal" K
  - Trying a few plausible values
  - Using the elbow method

#### Elbow Method

- Define a clustering performance metric
  - Within cluster sum of square distances (WCSS)
- Calculate the performance for a few K
  - Most performance metrics are expected to decrease when K increasing
  - Take the K at the elbow

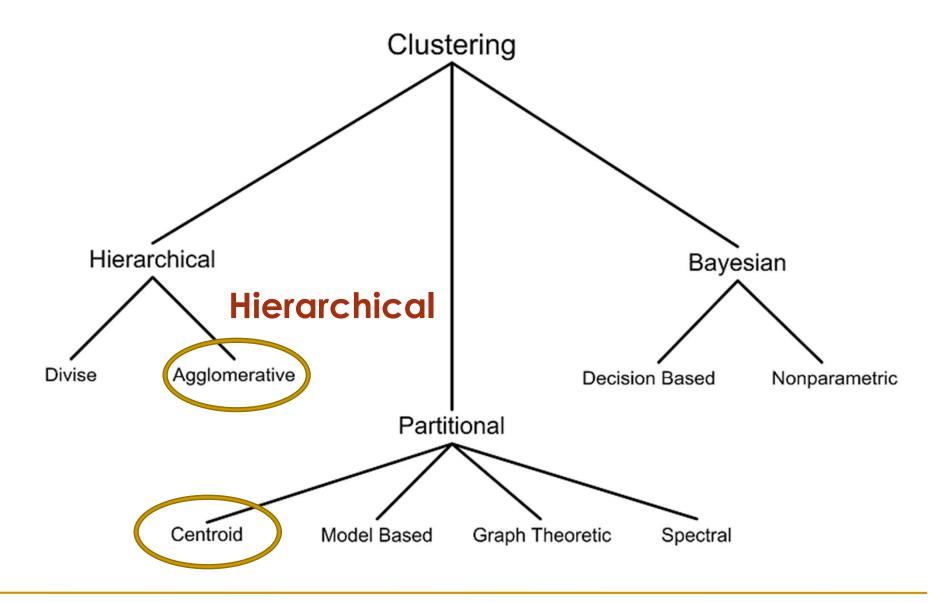
#### Elbow Method



### K-means: Summary

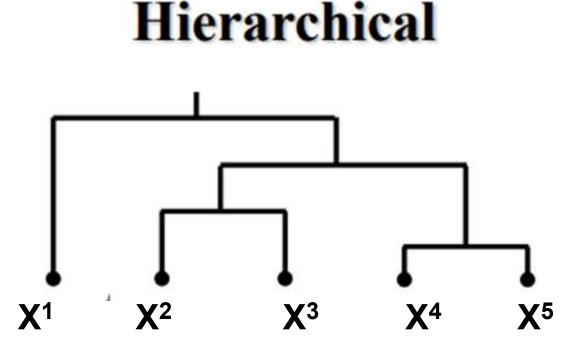
- Despite weaknesses, k-means is still one of the most popular algorithms, due to its simplicity and efficiency
- No clear evidence that any other clustering algorithm performs better in general
- Comparing different clustering algorithms is a difficult task.
  - No one knows the correct clusters!

# Clustering Techniques



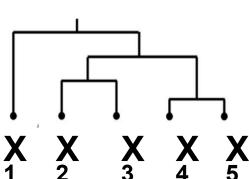
# Hierarchical Clustering

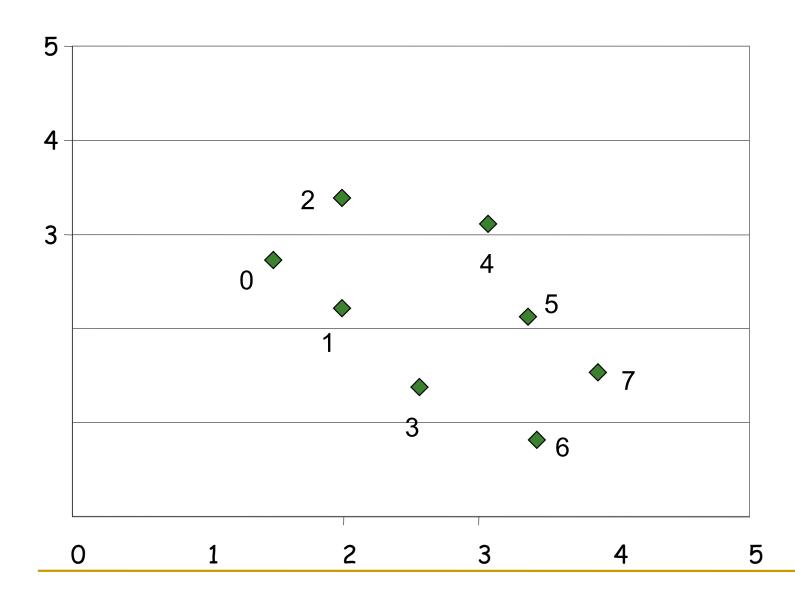
- A hierarchical decomposition of the observations using distance-based criteria
- Bottom-Up
   (Agglomerative)
   Clustering is the
   most popular

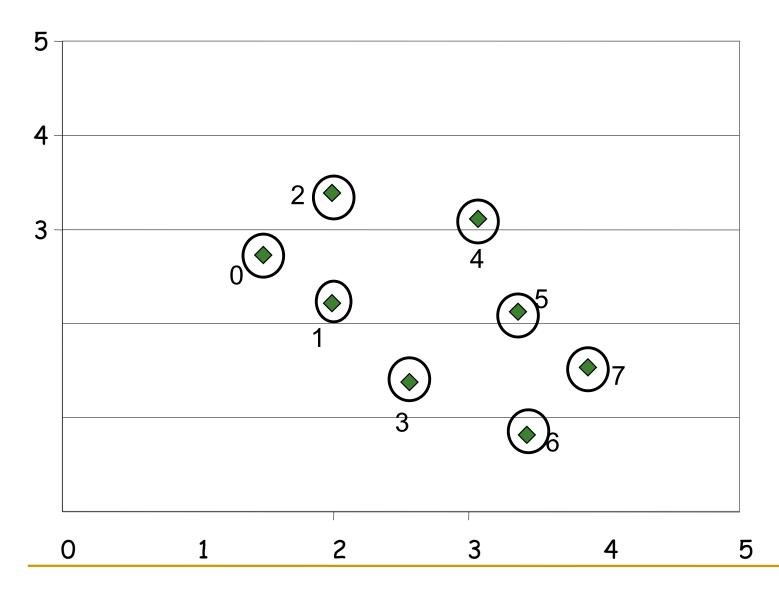


# Bottom-Up (Agglomerative) Clustering Hierarchical

- 1) Start with every observation as a cluster
- 2) Find the best two clusters to merge to become a new cluster
- Repeat Step 2 until all clusters are merged as a single cluster with all n observation.
- Analyze the cluster formation (or the dendrograms) and select the optimal number of clusters

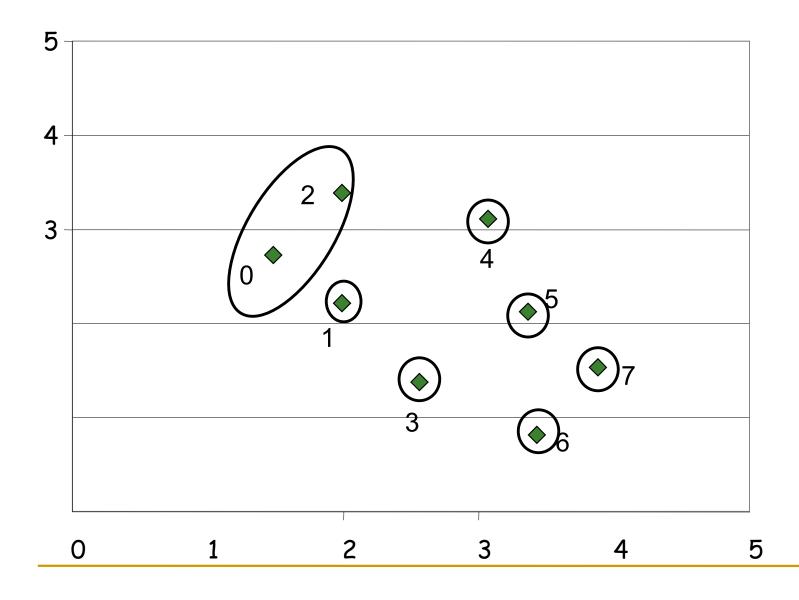


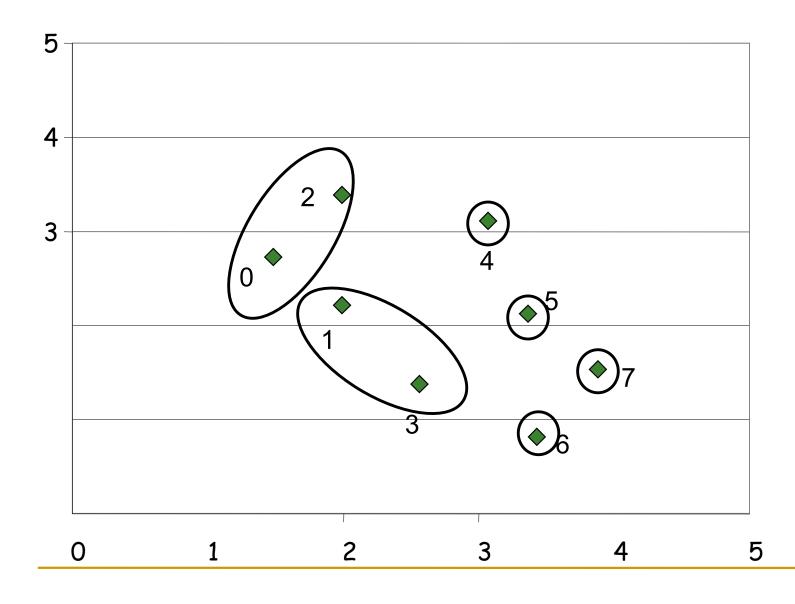


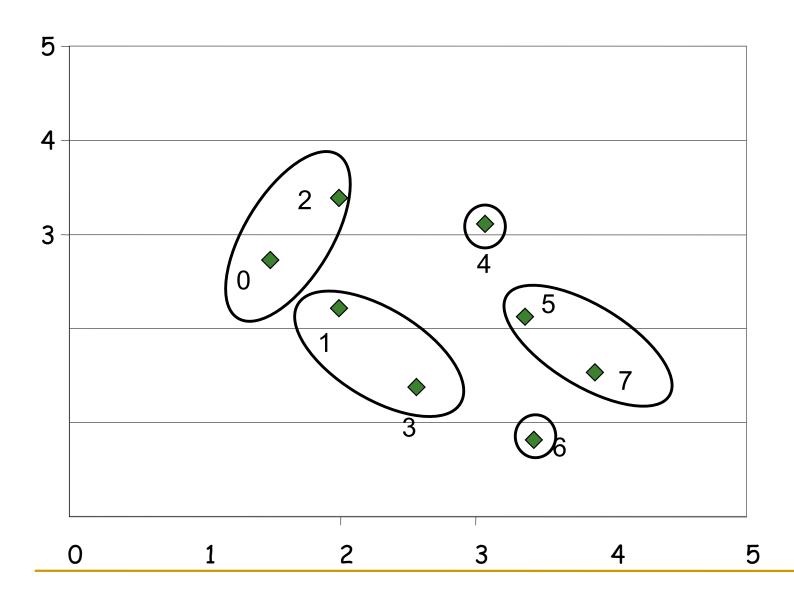


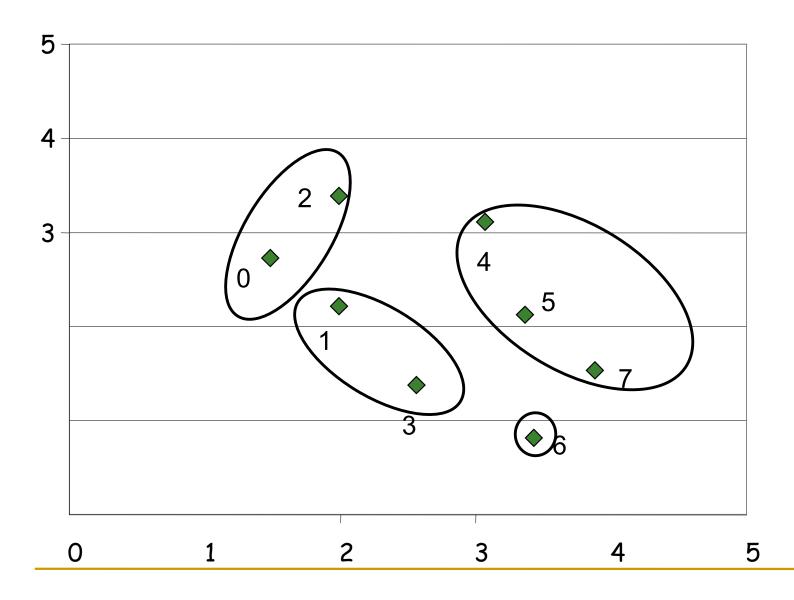
Every observation as a cluster

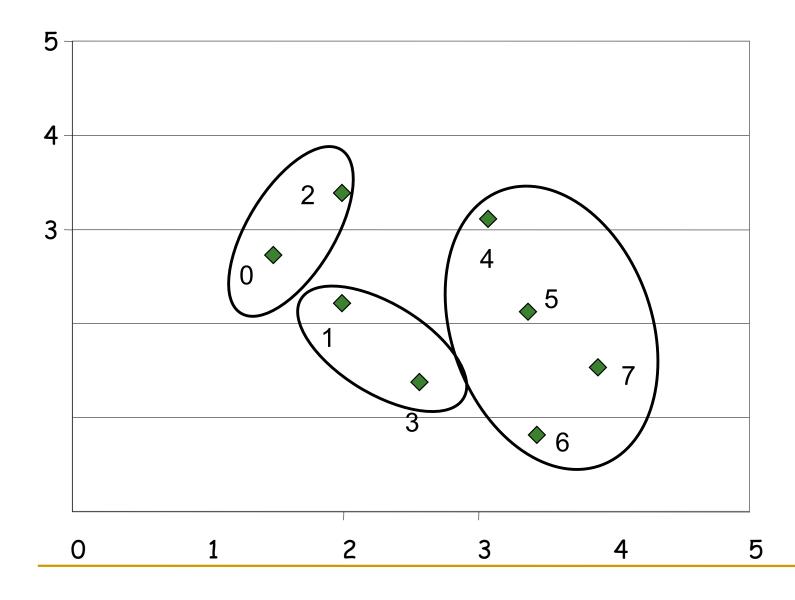
Nc = m = 8

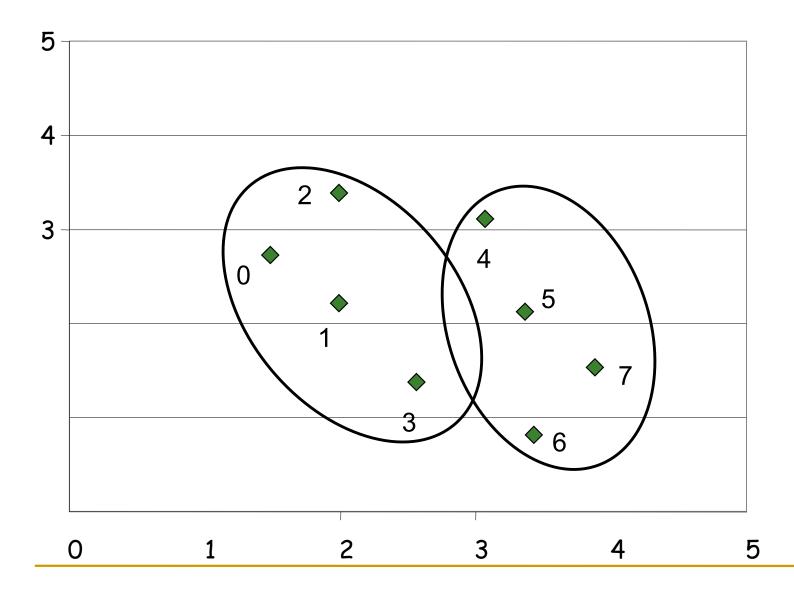


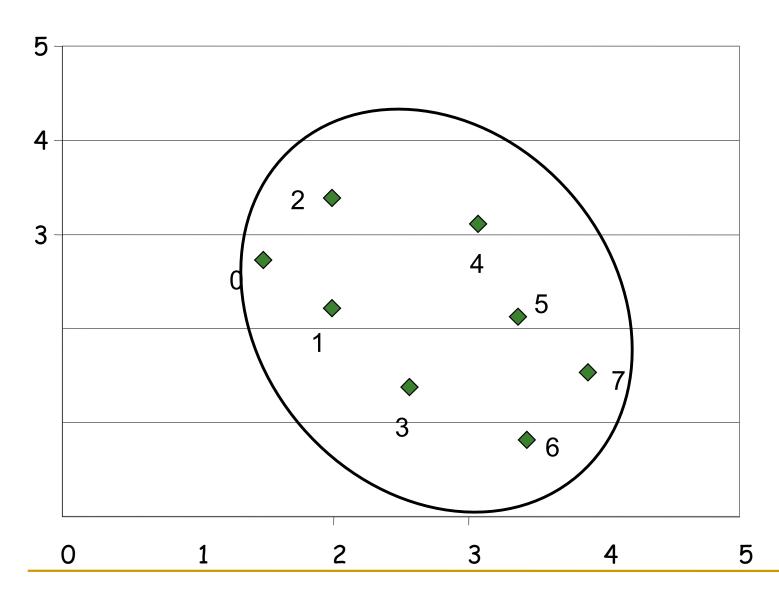










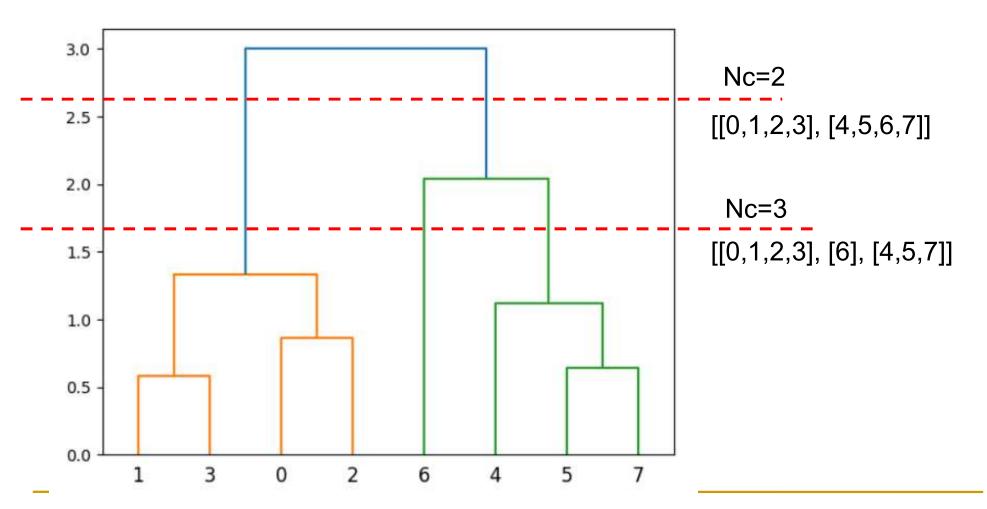


One cluster with all m=8 observation

## Optimal number of clusters

A big jump in distance (A high cost to merge two clusters) indicates that these two clusters are not similar, and the merge is not a proper decision.

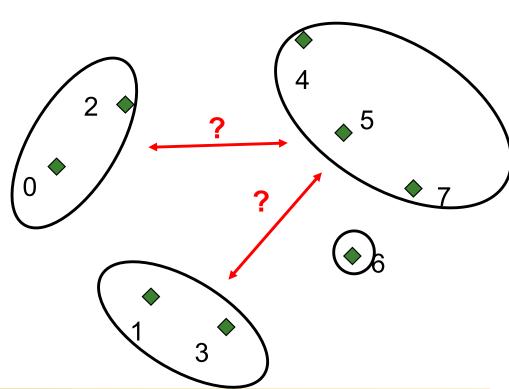
A horizontal line cutting such "improper" moves gives the optimal number of clusters



#### Distance of two clusters?

- We know how to find the distance of two observations. But, what about two cluster?
- Different approaches
  - Closest members
  - Farthest members
  - Average of all members
  - Distance of centroids

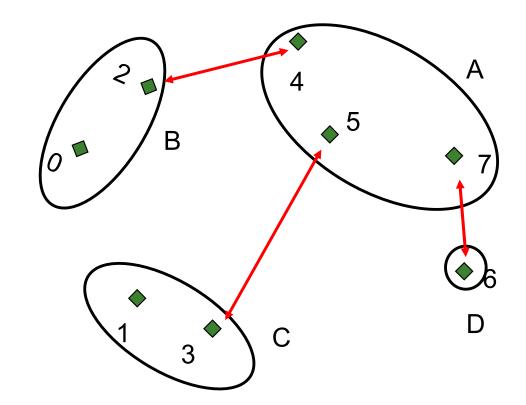
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#### Distance of two clusters: Single Link

- Distance of two closest members
- Potentially results in skinny long clusters

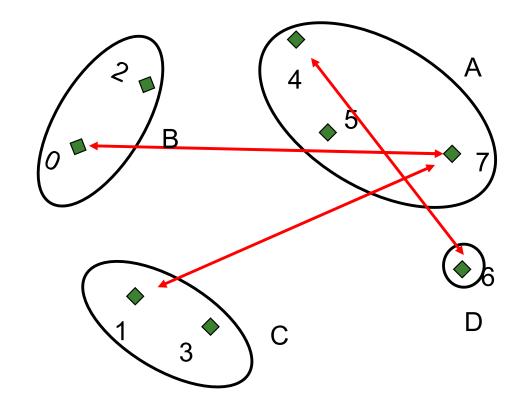
d (ci, cj)	Given by
A, B	d (2,4)
A, C	d (5,3)
A, D	d (7,6)



#### Distance of two clusters: Complete Link

- Distance of two farthest members
- Prone to noise and outliers

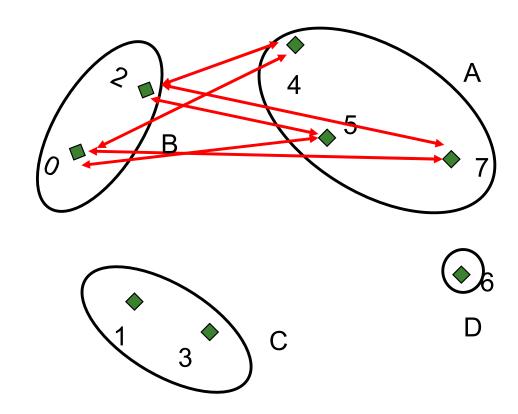
d (ci, cj)	Given by
A, B	d (7,0)
A, C	d (7,1)
A, D	d (4,6)



#### Distance of two clusters: Average Link

- Average distance of all pairs between two clusters
- More robust against noise and outliers

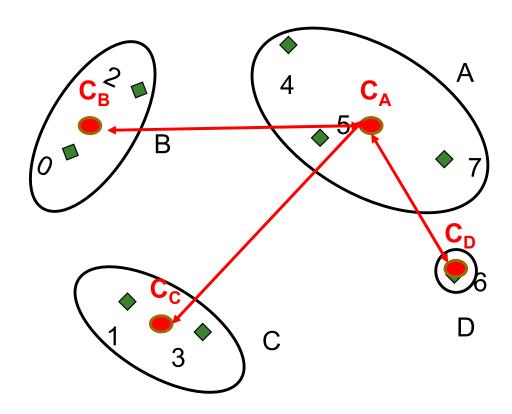
d (ci, cj)	Given by
A, B	Average of all 6 links
A, C	Average of all 6 links
A, D	Average of d(4,6), d(5,6), and d(7,6)



#### Distance of two clusters: Centroid Link

- Distance between the centroids of two clusters
- More robust against noise and outliers

d (ci, cj)	Given by
A, B	$d(C_A, C_B)$
A, C	$d(C_A, C_C)$
A, D	$d(C_A, C_D)$

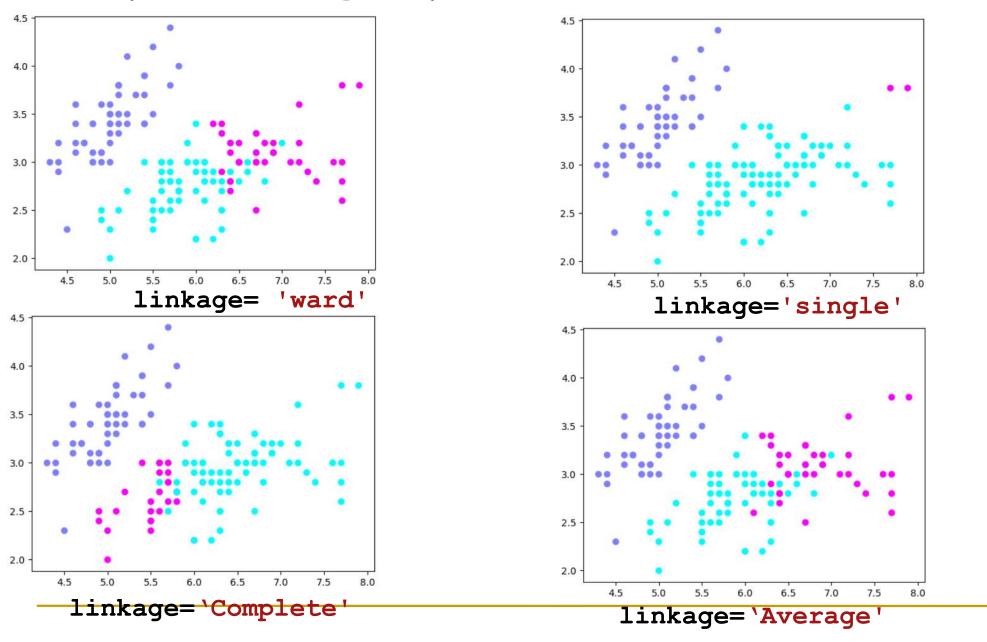


# Distance of two clusters: Ward Minimum Variance Link

- Minimizes the total within-cluster variance
- Merges two clusters that results in a minimum increase of within cluster variance
- Tends to merge smaller clusters together

d(A, B) = 
$$\frac{||CA - CB||^2}{\frac{1}{n_A} + \frac{1}{n_B}}$$

#### Compare linkage options



Iris dataset from sklearn, using Euclidean distance

#### Agglomerative Clustering: Summary

- Simple and easy to understand
- No need to specify the number of clusters in advance.
- Complexity is usually higher than K-Means Number of optimal clusters is subjective.
   No one knows the correct clusters!

# Today

- Hierarchical Clustering
- 2. K-means Clustering
- 3. Hierarchical Clustering

