## Data Clustering

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### Outline

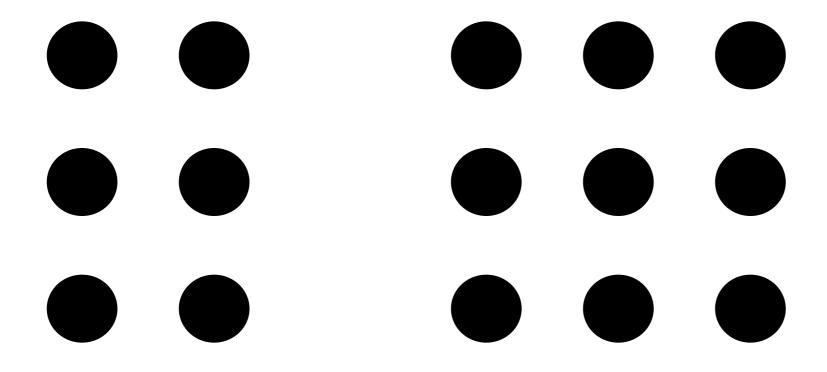
- Why cluster data?
- Clustering as unsupervised learning
- Clustering algorithms
  - k-means, k-medoids
  - agglomerative clustering
  - Brown's clustering
  - Spectral clustering
- Cluster evaluation measures
  - Purity
  - Normalised Mutual Information
  - Rand Index
  - B-CUBED
  - Precision, Recall, F-score
- Supervised clustering

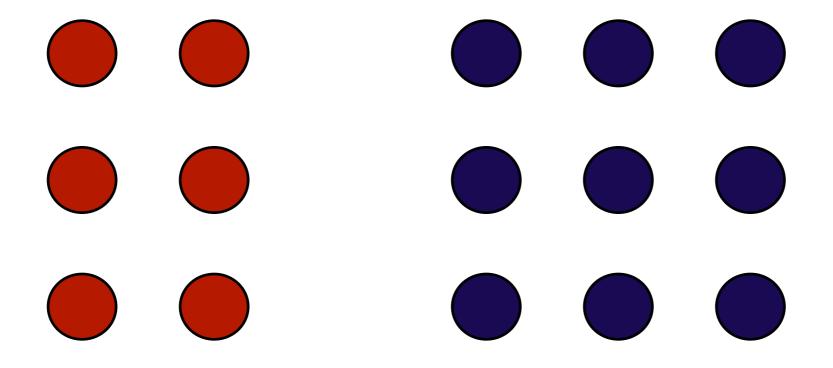
## Why cluster data?

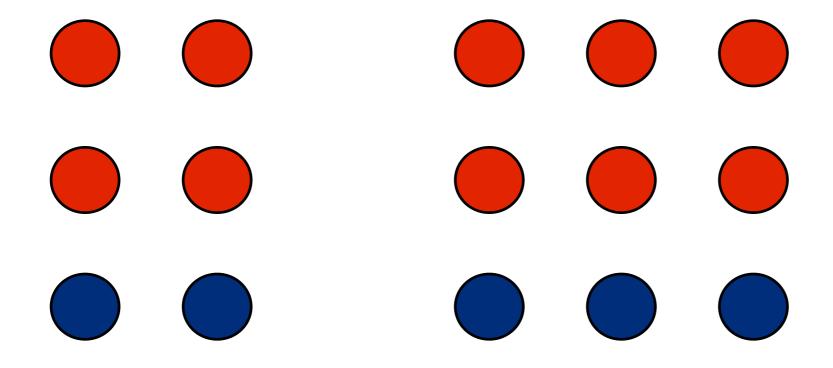
- Data Mining has two main objectives
  - Prediction: classification, regression etc.
  - Description: pattern mining, rule extraction, visualisation *clustering*
- Clustering is:
  - Unsupervised learning
    - no label data is required (consider classification algorithms we discussed so far in the lecture which are supervised algorithms)
  - Generalisation / Abstraction of concepts
  - Topic detection
  - Visualisation
  - Outlier detection

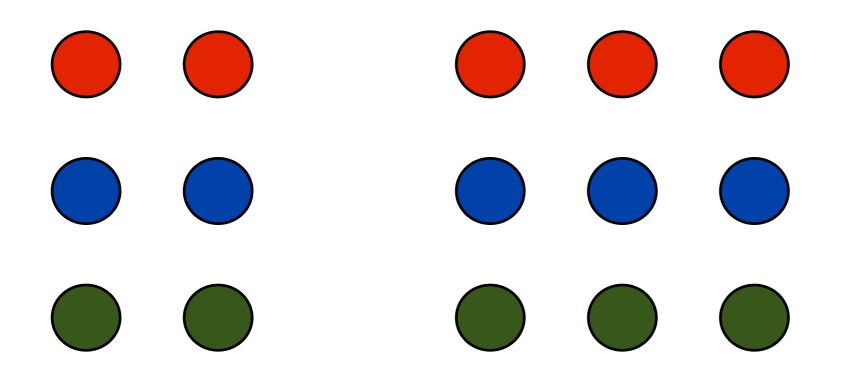
## Unsupervised Learning

- Supervised learning
  - labels for training instances are provided
- Unsupervised learning
  - No labels for training instances are provide
- Semi-supervised learning
  - Both labeled and unlabeled training instances are provided
- What can we learn about training data if we do not have any labels?
  - The similarity and distribution of the features can still be learnt and this can be used to create rich feature spaces for supervised learning (if required)









How many clusters?

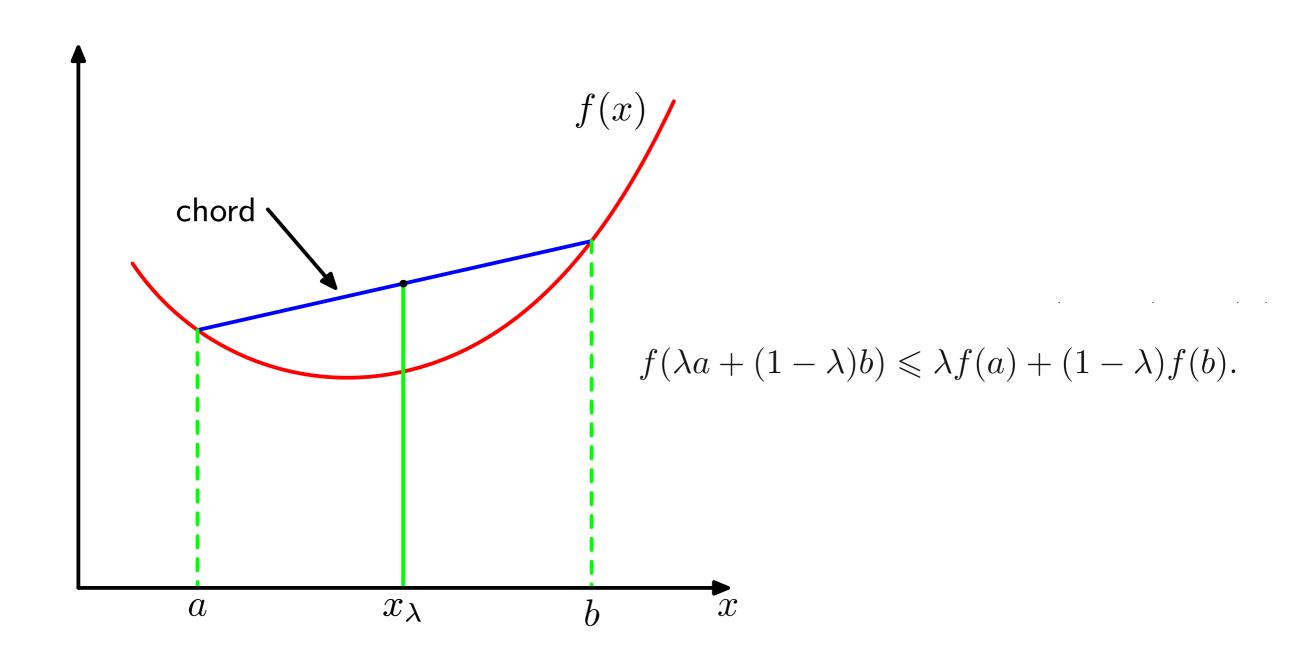
### General Remarks

- A single dataset can be clustered into several ways
- There is no single right or wrong clustering
  - Simply different views on the same data
- If so how can we measure the quality of a clustering algorithm?
  - Two ways
    - Compare the clusters produced by a clustering algorithm against some reference (gold standard) set of clusters (direct evaluation)
    - Use the clusters as features for some other (eg. supervised learning) task and measure the difference in the performance of the second task (indirect evaluation)

## Clustering as Optimisation

- Given a dataset  $\{\mathbf{x}_1, ..., \mathbf{x}_N\}$  of N instances represented as d dimensional real vectors  $(\mathbf{x}_i \in R^d)$ , partition these N instances into k clusters  $S_1,...,S_k$  such that some objective function  $f(S_1,...,S_k)$  is minimised.
- Observations
  - k and f are given
  - f can be the similarity between the clusters (good to create dissimilar clusters as much as possible), information gain, correlation and various other such goodness measures (heuristics)
  - Often clustering is an NP hard and a non-convex problem
    - http://rangevoting.org/VattaniKmeansNPC.pdf
    - approximations, relaxations are required in practice

### Convex Functions



## Clustering Algorithms

- Partitioning
  - Construct k partitions and iteratively update the partitions
    - k-Means, k-Medoids
- Hierarchical
  - Create a hierarchy of clusters (dendrogram)
    - Agglomerative clustering (bottom-up)
    - Conglomerative clustering (top-down)
- Graph-based clustering
  - Graph-cut algorithms (Spectral Clustering)
- Model-based clustering
  - Mixture of Gaussians
- Other types: Non-parametric Bayesian (Latent Dirichlet Allocation), Expectation Maximisation (EM) algorithm, and many more ...

### k-Means Derivation

$$\arg\min_{S_1,...,S_k} \sum_{i=1}^k \sum_{\boldsymbol{x}_i \in S_i} ||\boldsymbol{x}_j - \boldsymbol{\mu}_i||^2$$

We want to minimize the distance between data instances ( $x_i$ ) and some cluster centres ( $\mu_i$ )

$$f(S_1, \dots, S_k) = \sum_{i=1}^k \sum_{x_i \in S_i} ||x_j - \mu_i||^2$$

This objective function is called the within cluster sum of squares (WCSS) objective

$$\frac{\partial f(S_1, \dots, S_k)}{\partial \mu_i} = 0$$

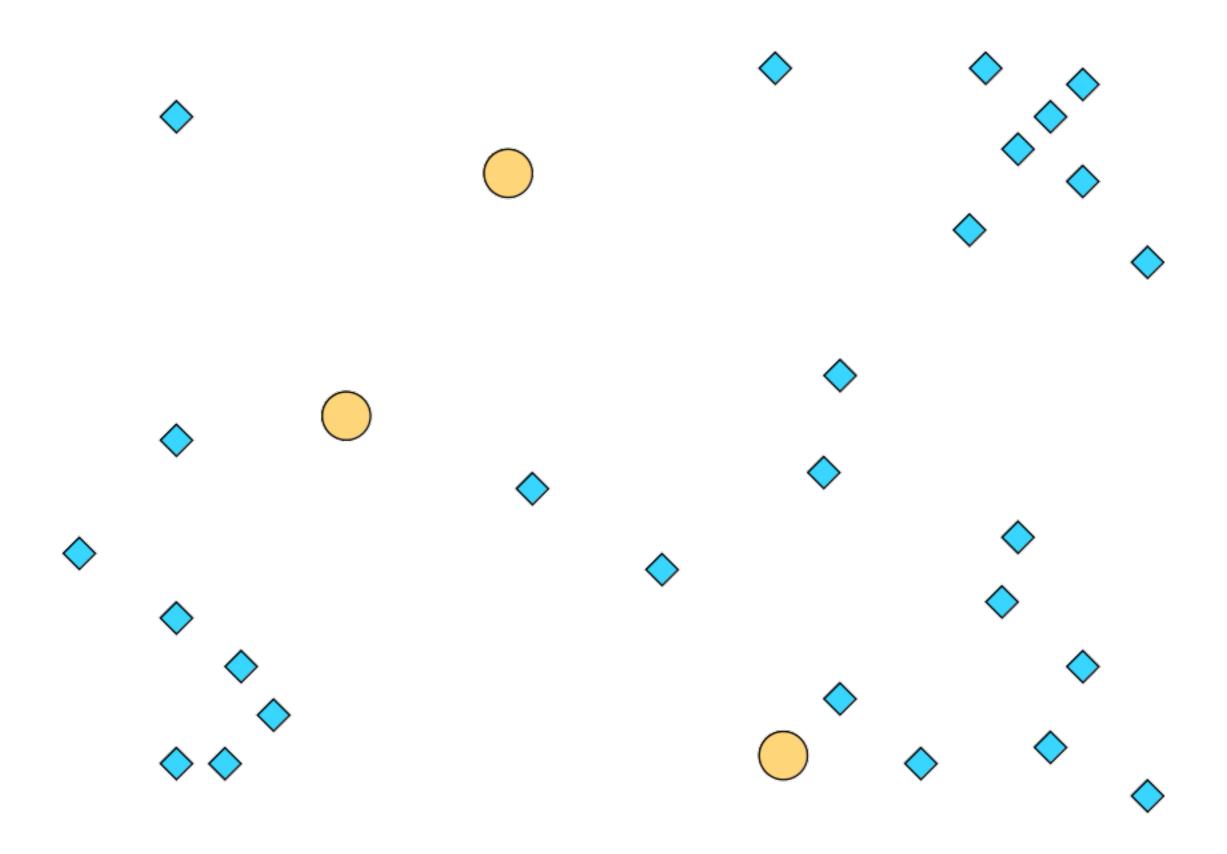
$$\frac{\partial f(S_1, \dots, S_k)}{\partial \mu_i} = \sum_{\boldsymbol{x}_j \in S_i} 2(\boldsymbol{x}_j - \boldsymbol{\mu}_i)$$

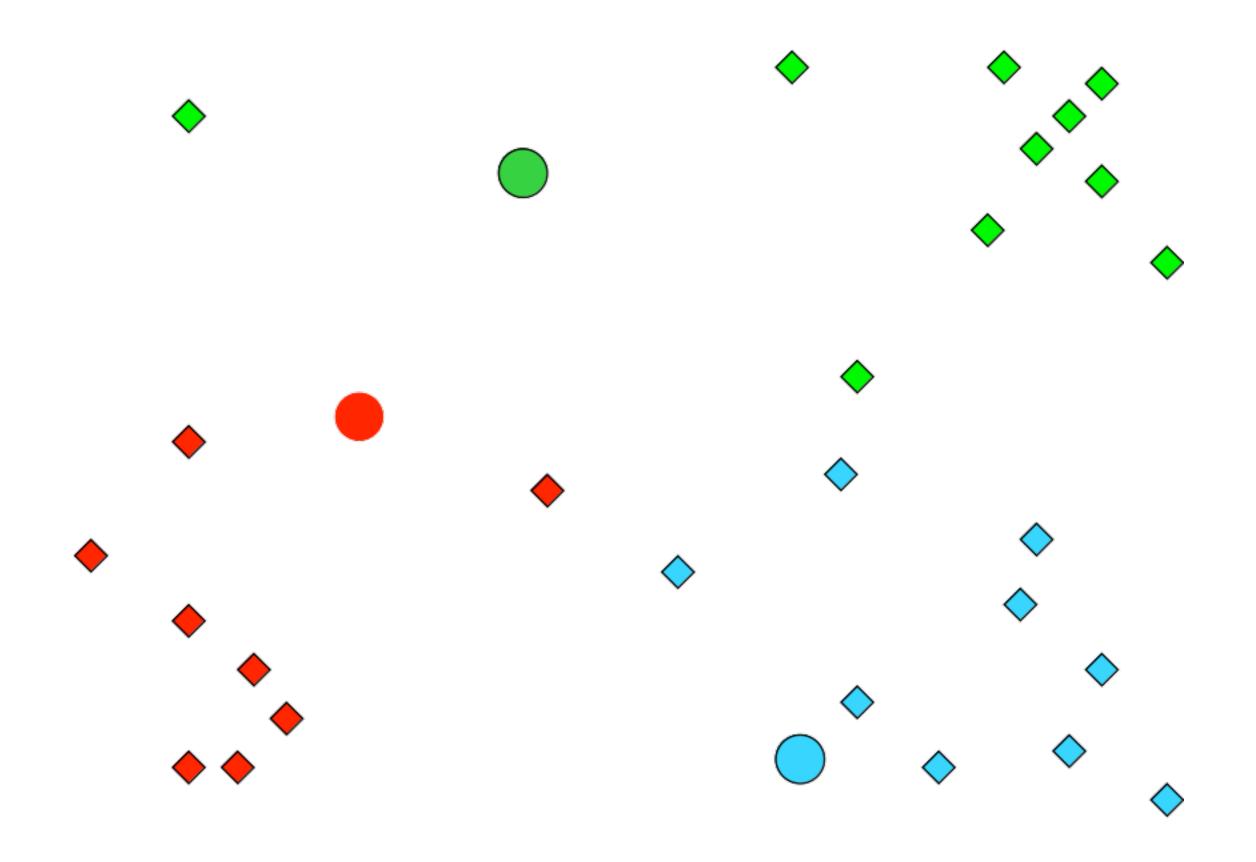
$$\mu_i = \frac{1}{|S_i|} \sum_{\boldsymbol{x}_j \in S_i} \boldsymbol{x}_j$$

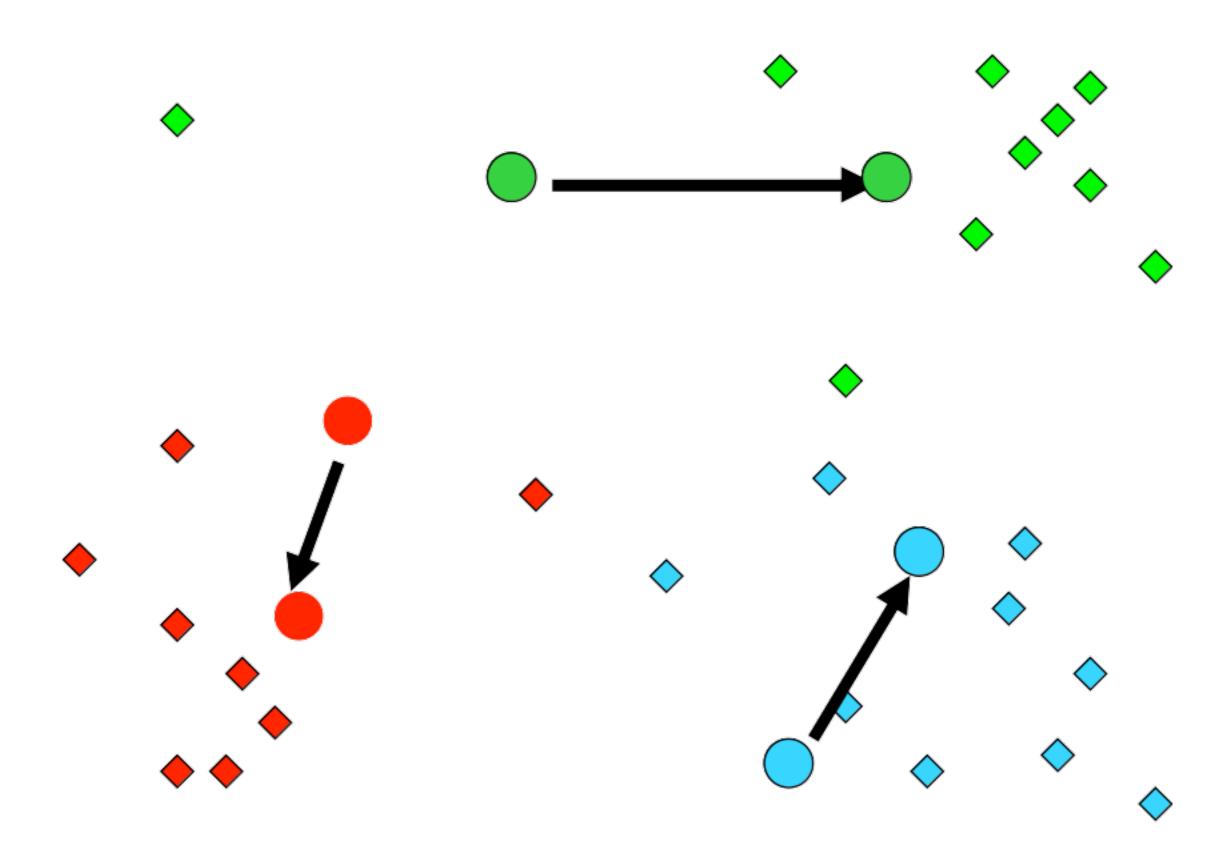
Just compute the centroid (mean) of each cluster and that will give you the cluster centers

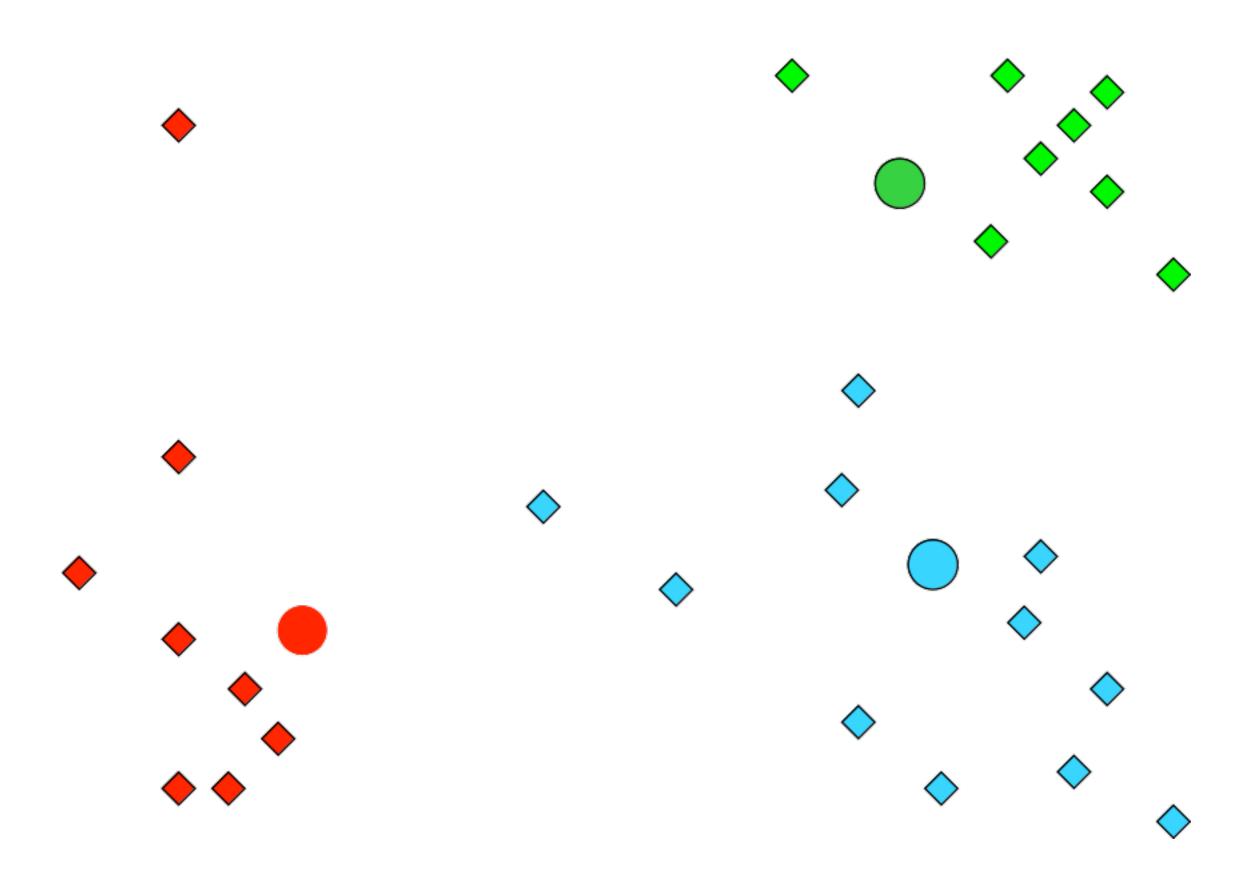
## k-Means Clustering

- INPUT
  - The number of clusters k
  - Dataset {x<sub>1</sub>, ..., x<sub>N</sub>} of N instances represented as d dimensional real vectors (x<sub>i</sub> ∈ R<sup>d</sup>)
- 1. Set k instances from the dataset randomly. (initial cluster means/centers)
- 2. Assign all other instances to the closest cluster centre.
- 3. Compute the mean of each cluster
- 4. until convergence repeat between steps 2 and 3 convergence = no instances have moved among clusters (often after a fixed number of iterations specified by the user)









### Issues with k-Means

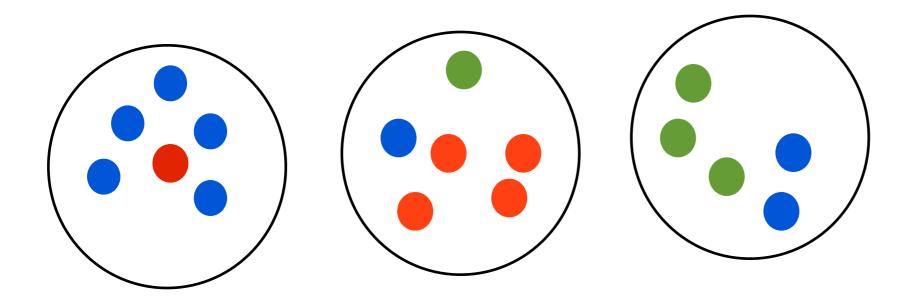
- Results can vary depending on the initial random choices
- Can get trapped in a local minimum that isn't the global optimal solution
  - Repeat the clustering procedure multiple times with different initialisations and select the best final clustering
    - best? according to what? many heuristics exist.
      - smallest number of iterations before convergence
      - largest total distance between the final cluster means
- Outliers have a larger effect on the mean value, hence cluster centre and the cluster
- cluster centres (means) are not actual instances in the cluster
  - We could pick actual instances as initial cluster centroids.

### Evaluating Clustering — Purity

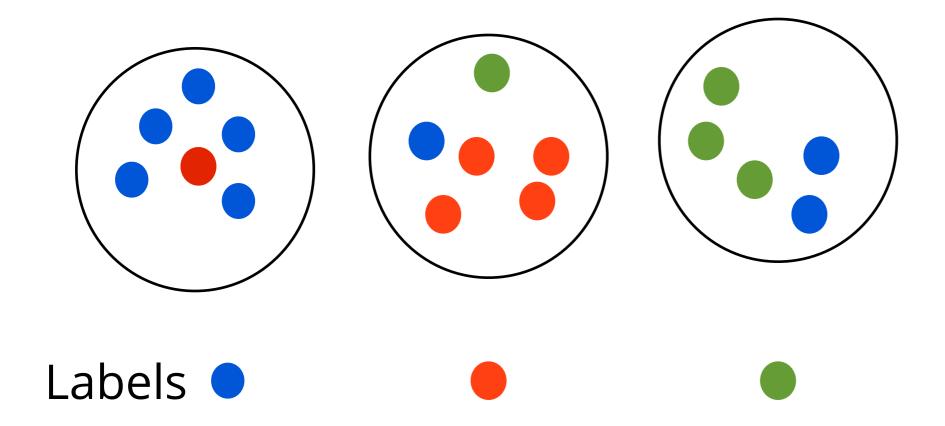
- Let us assume that we have a set  $\Omega = \{\omega_1, ..., \omega_K\}$  clusters for a set of classes  $C = \{c_1, ..., c_J\}$
- Assign each cluster the label that appears most in that cluster (aka. majority labelling)
- Purity measures the ratio of the items that are in the cluster with the same class as its own.

purity(
$$\Omega$$
, C) =  $\frac{1}{N} \sum_{k} \max_{j} |\omega_{k} \cap c_{j}|$ 

Here, N is the total number of items.



Quiz: Compute purity for this clustering.



purity = 
$$(5 + 4 + 3) / 17 = 12/17 = 0.71$$

Purity achieves its maximum value of 1 for singletons (each item is in a cluster containing only that single item)!

Obviously this is not good "clustering" and purity does not recognise this.

## Evaluating Clustering — NMI

- Let us assume that we have a set  $\Omega = \{\omega_1, ..., \omega_K\}$  clusters for a set of classes  $C = \{c_1, ..., c_J\}$
- Assign each cluster the label that appears most in that cluster (aka. majority labelling)
- Normalised Mutual Information (NMI) computes the ratio of information that we can know about the classes C given the clusters  $\Omega$  to the averaged information that is contained in C and  $\Omega$ .

$$NMI(\Omega, C) = \frac{I(\Omega, C)}{[H(\Omega) + H(C)]/2}$$

$$I(\Omega, \mathcal{C}) = \sum_{k} \sum_{j} p(\omega_k \cap c_j) \log \left( \frac{p(\omega_k \cap c_j)}{p(\omega_k)p(c_j)} \right)$$
$$= \sum_{k} \sum_{j} \frac{|\omega_k \cap c_j|}{N} \log \left( \frac{N|\omega_k \cap c_j|}{|\omega_k||c_j|} \right)$$

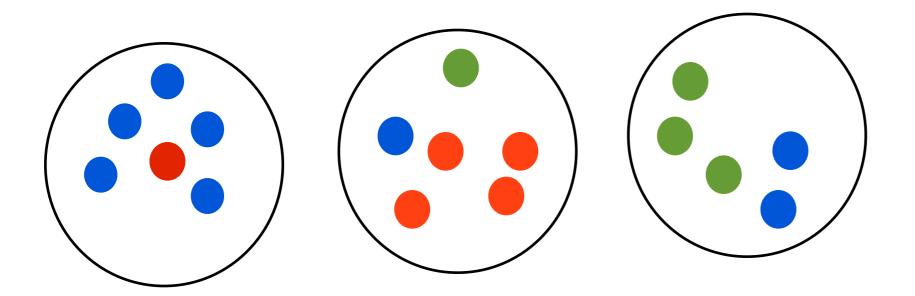
$$H(\Omega) = -\sum_{k} p(\omega_k) \log p(\omega_k)$$
$$= -\sum_{k} \frac{|\omega_k|}{N} \log \frac{|\omega_k|}{N}$$

Mutual Information (MI)

Entropy

#### Why we do we normalise by the average?

- $I(X,Y) \le [H(X) + H(Y)]/2$
- Proof (sketch):
  - I(X,Y) = H[X] H[X|Y] = H[Y] H[Y|X]
  - Add those two and use the fact that (conditional) entropy is nonnegative
    - $H[X|Y] + H[Y|X] \ge 0$



Quiz: Compute NMI for this clustering.

Let 
$$C_1 = Blue$$
,  $C_2 = Pad$  and  $C_3 = Green$ .

$$P(C_1) = \frac{8}{17}, \quad P(C_2) = \frac{C}{17}, \quad P(C_3) = \frac{4}{17}.$$

$$\therefore H(C) = -\frac{3}{2}P(C_1) \log P(C_2)$$

$$= -\left[\frac{8}{17}\log \frac{E}{17} + \frac{5}{17}\log \frac{C}{17} + \frac{4}{17}\log \frac{4}{17}\right] = 1.055$$

Chewise,
$$P(\omega_1) = \frac{6}{17}, \quad P(\omega_2) = \frac{6}{17}, \quad P(\omega_3) = \frac{C}{17}$$

$$H[\Omega] = -\left[\frac{6}{17}\log \frac{6}{17} + \frac{C}{17}\log \frac{6}{17} + \frac{5}{17}\log \frac{7}{17}\right] = 1.095$$

$$P(\omega_1 \cap C_4) = \frac{5}{17} \quad P(\omega_1 \cap C_2) = \frac{1}{17} \quad P(\omega_1 \cap C_3) = \frac{0}{17}$$

$$P(\omega_2 \cap C_1) = \frac{1}{17} \quad P(\omega_2 \cap C_2) = \frac{4}{17} \quad P(\omega_2 \cap C_3) = \frac{1}{17}$$

$$P(\omega_3 \cap C_1) = \frac{2}{17} \quad P(\omega_3 \cap C_2) = \frac{0}{17} \quad P(\omega_3 \cap C_3) = \frac{3}{17}$$

$$T(\omega_1 \cap C_1) = \frac{2}{17} \quad P(\omega_2 \cap C_2) \log \frac{1}{17} \quad P(\omega_3 \cap C_3) = \frac{3}{17}$$

$$T(\omega_1 \cap C_1) = \frac{2}{17} \quad P(\omega_2 \cap C_2) \log \frac{1}{17} \quad P(\omega_3 \cap C_3) = \frac{3}{17}$$

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```
import math

C = [5,1,1,4,1,2,3]

p = []
N = sum(C)
for x in C:
    p.append(float(x) / float(N))

E = 0
for x in p:
    E += - x * math.log(x)

print E
```

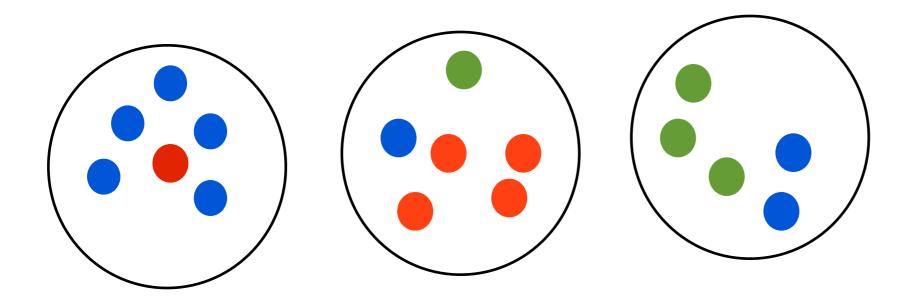
#### Evaluating Clustering — Rand Index (RI)

- Assign each cluster the label that appears most in that cluster (aka. majority labelling)
- Build a contingency table considering pairs of items in each cluster
  - Positive = same cluster
  - Negative = different clusters
  - True = same class
  - False = different classes
- TP = No. of item pairs that are in the same cluster and belong to the same class
- FP = No. of item pairs that are in the same cluster but belong to different classes
- TN = No. of item pairs that are in different clusters and belong to different classes
- FN = No. of item pairs that are in different clusters but belong to the same class

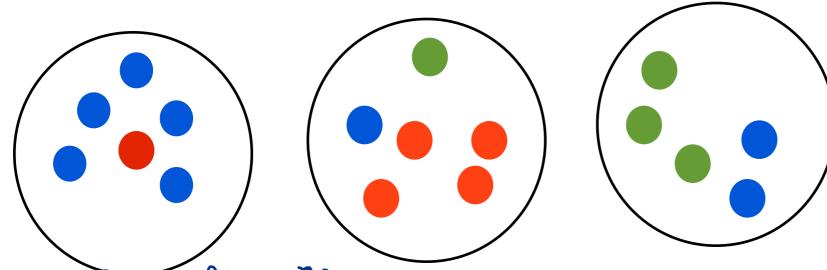
contingency table	same cluster	different clusters		
same class	TP	FN		
different classes	FP	TN		

$$RI = \frac{TP + TN}{TP + FP + TN + FN}$$

(accuracy of the clustering)



Quiz: Compute RI for this clustering.



TP+FP = 
$$6C_2 + 6C_2 + 3C_2 = 15 + 15 + 10 = 40$$
.

$$\left[ n_{Y} = \frac{n!}{r! (n-r)!} \right]$$

$$6(2 = \frac{6!}{2! \cdot 4!} = \frac{6 \times 5}{2} = 15$$
  $5(2 = \frac{5!}{2! \cdot 3!} = \frac{5 \times 4}{2} = 10.$ 

$$TP = {}^{5}C_{2} + {}^{4}C_{2} + {}^{3}C_{2} + {}^{2}C_{2} = 10 + 6 + 3 + 1 = 20$$

$${}^{4}C_{2} = \frac{4!}{2! \cdot 2!} = \frac{4 \times 3}{2} = 6$$

$$FN = (5x1) + (1x4) + (5x2) + (1x2) + (1x3)$$
  
=  $5+4+10+2+3=24$ 

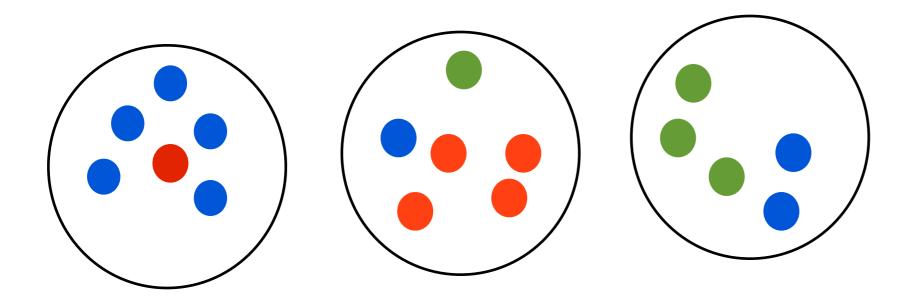
$$TN + FN = (5+1)(1+1+4) + (5+1)(3+2) + (1+1+4)(3+2)$$

$$= 6\times6 + 6\times5 + 6\times5 = 36 + 30 + 30 = 96.$$

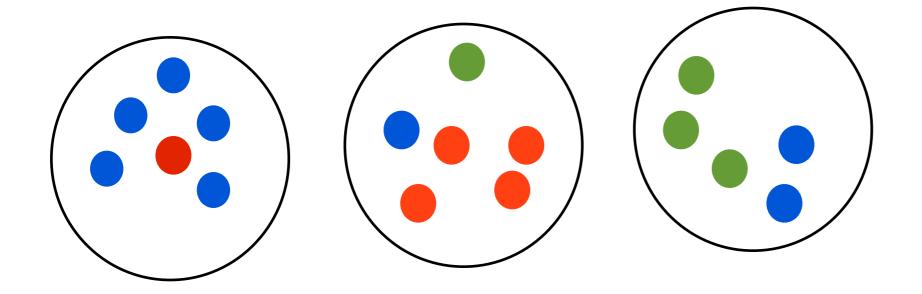
	same cluster	different clusters
same class	20	24
different classes	20	72

## Evaluating Clustering — P/R/F

- We can use Precision (P), Recall (R), and F-measure
   (F) at to evaluate the accuracy of a clustering.
- For this purpose we must first create the contingency table as we did for RI and then compute P, R, F as follows



Quiz: Compute P/R/F for this clustering.



	same cluster	different clusters
same class	TP=20	FN=24
different classes	FP=20	TN=72

$$P = TP / (TP + FP) = 20 / (20+20) = 0.5$$
  
 $R = TP / (TP + FN) = 20 / (20 + 24) = 0.45$   
 $F = 2PR / (P + R) = 0.47$ 

### **B-CUBED** Measure

- Proposed in (Bagga B. Baldwin = B<sup>3</sup>)
  - A. Bagga and B. Baldwin. Entity-based cross document coreference resolution using the vector space model, In Proc. of 36th COLING-ACL, pages 79--85, 1998.
- We would like to evaluate clustering without labelling any clusters.

$$\operatorname{precision}(x) = \frac{\text{No. of items in C(x) with A(x)}}{\text{No. of items in C(x)}}$$
$$\operatorname{recall}(x) = \frac{\text{No. of items in C(x) with A(x)}}{\text{Total no. of items with A(x)}}$$

C(x): The ID of the cluster that x belongs to

A(x): label of x

### **B-CUBED Measure**

 Compute the average over all the items (instances) that appear in all clusters (N)

$$Precision = \frac{1}{N} \sum_{p \in DataSet} Precision(p)$$

$$Recall = \frac{1}{N} \sum_{p \in DataSet} Recall(p)$$

$$F-Score = \frac{1}{N} \sum_{p \in DataSet} F(p)$$

## Hierarchical Clustering

- Sometimes we might want to organise the data into a hierarchy of subsuming concepts for visualisation (abstraction) purposes
- Two methods exists
  - Conglomerative clustering
    - Start from one big cluster with all data instances and repeatedly partition it
    - Top-down approach
  - Agglomerative clustering
    - Start singletons (clusters with exactly one instance) and iteratively merge the most similar two clusters
      - Bottom-up approach
      - computationally more efficient (O(logn) merges required)

## Merging two clusters

- Single linkage
  - Distance between two clusters A and B is the smallest distance between any instance  $a \in A$  and  $b \in B$

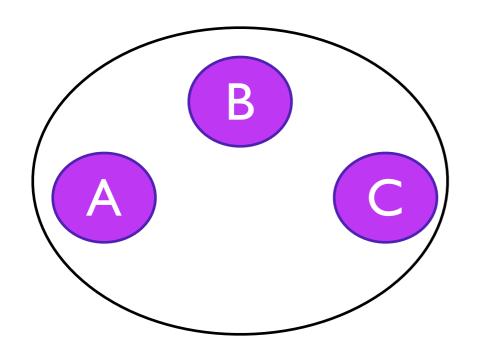
$$D(\mathcal{A}, \mathcal{B}) = \min_{a \in \mathcal{A}, b \in \mathcal{B}} dist(a, b)$$

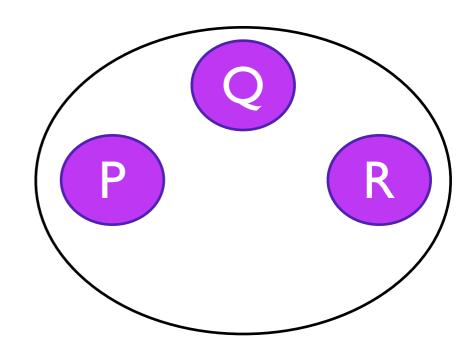
- Complete linkage
  - Distance between two clusters A and B is the largest distance between any instance  $a \in A$  and  $b \in B$

$$D(\mathcal{A},\mathcal{B}) = \max_{\substack{a \in \mathcal{A}, b \in \mathcal{B}}} dist(a,b)$$
   
 • Average linkage (Group-Average)

- - Average of all the pairs selected from each cluster

$$D(\mathcal{A}, \mathcal{B}) = \frac{1}{|\mathcal{A}||\mathcal{B}|} \sum_{a \in \mathcal{A}, b \in \mathcal{B}} dist(a, b)$$





Quiz: Let us assume that in the 2D space there are two clusters {A,B,C} and {P,Q,R}. Which of the distances correspond to the single link and complete link distances between the shown clusters?

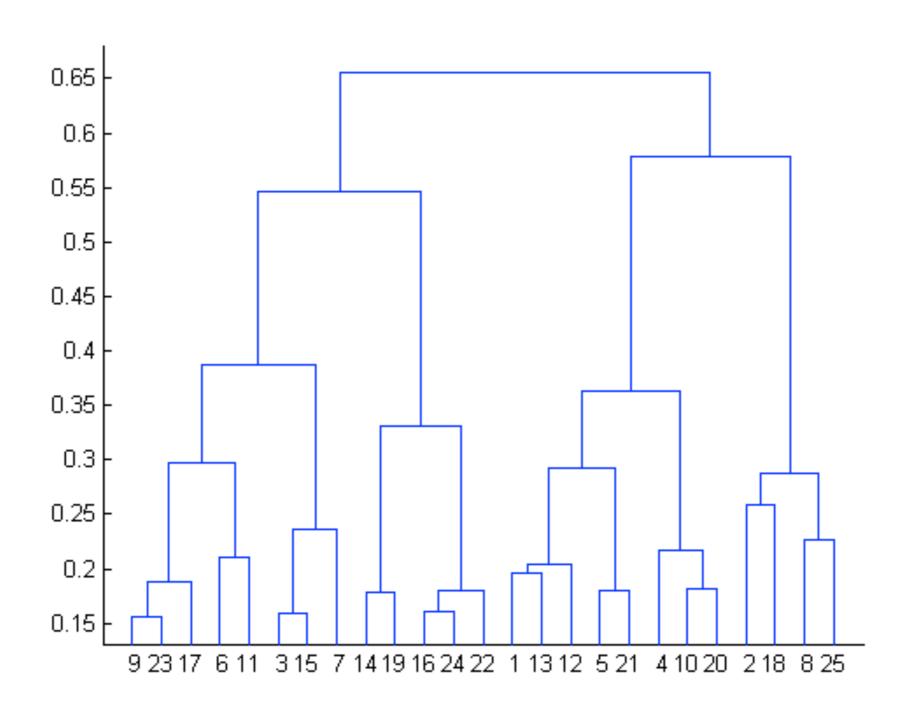
#### Group-Average Agglomerative Clustering

#### • INPUT:

- A set of N data instances  $\{x_1, ..., x_N\}$ , Number of clusters k
- Initialise
  - Create singletons  $S_i = \{x_i\}$  for i = 1, ..., N
- Repeat until only we are left with one cluster
  - Merge the two clusters S<sub>i</sub> and S<sub>j</sub> with the minimum distance (cf. maximum similarity)

$$D(\mathcal{S}_i, \mathcal{S}_j) = \frac{1}{|\mathcal{S}_i||\mathcal{S}_j|} \sum_{a \in \mathcal{S}_i, b \in \mathcal{S}_j} dist(a, b)$$

# Dendrogram



### Clusters as Features

- We can use clustering to find similar features in instances without requiring any supervision (no label data is required for clustering)
  - Distributional similarity of features over instances
- Once we have clustered the features, we can use the cluster IDs as features
- Benefits
  - Reduces the dimensionality of the feature space
    - dogs and cats are mapped to pets
  - Reduces feature sparseness
    - If at least one of the features in a cluster appears in an instance, then we assume that the entire cluster appeared as a feature for that instance

#### Clustering the word-document Matrix

	d <sub>1</sub>	$d_2$	d <sub>3</sub>	$d_4$	$d_5$	d <sub>6</sub>
dog	2	3	0	0		5
cat	I	I	0	0	3	2
apple	0	0	I	2	3	0
banana	0	0	2	5	0	0

## Clustering the 2D matrix

- Each row vector can be seen as the feature vector for each word (row elements)
  - We can measure similarity between row vectors and cluster the words
- Each column vector can be seen as the feature vector for each document (column elements)
  - We can measure similarity between column vectors and cluster the documents

## Co-clustering

- Cluster both rows and columns simultaneously!
  - You will get both row and column clusters
- Information Theoretic Co-Clustering (ITCC)
  - Inderjit Dhillion and Subramanyam Mallela and Dharmendra Modha, pp. 89--98, International Conference on Knowledge Discovery and Data Mining (KDD), 2003.

### Clustering as Graph Partitioning

- Given a set of instances  $\{x_1, ..., x_N\}$ , we can represent these instances using a weighted undirected graph G, where the weight of the edge that connects two vertices in the graph corresponds to the similarity between the corresponding vertices.
- Then, the clustering problem becomes a graph partitioning problem where we must delete k edges from this graph to create k number of clusters
  - Spectral clustering algorithms (discussed later at the lecture for Graph Mining)