## Machine Learning, Machine Learning (extended)

7 – Unsupervised Learning: Clustering Kashif Rajpoot

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#### Outline

- Supervised vs unsupervised learning
- Clustering
- Similarity measure
- K-means clustering
- Kernelized k-means clustering
- Hierarchical agglomerative clustering

# Supervised vs unsupervised learning

- So far, we have looked at supervised learning
- Supervised learning
  - The algorithm learns from  $x_1, x_2, ..., x_N$  and  $t_1, t_2, ..., t_N$  so that it can later classify  $x_{new}$
  - The availability of  $t_n$  makes learning a supervised task
- What if we only have  $x_n$ , but not  $t_n$ ?
  - Unsupervised learning
- Clustering: create a grouping of objects
  - Each group containing "similar" objects

## Clustering: examples

- Shopping recommendation system
  - Recommend products likely to be purchased by a customer
- Grouping people from social network
  - Based on user activity
- Brain region network analysis
  - Determine "similar" brain regions that "activate" together or "rest" together

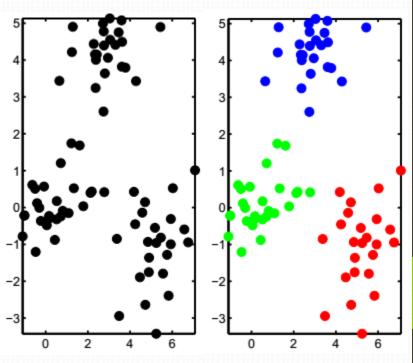
## Clustering

- Clustering: partition data into groups such that each group contains "similar" objects
  - Increase intra-group similarity
  - Similar objects should belong to same cluster
- How do we measure "similarity" between objects?
  - · "closeness"

$$D_{ij} = (x_i - x_j)^T (x_i - x_j)$$

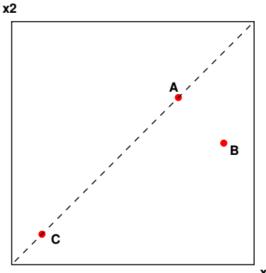
$$= (x_i^{(1)} - x_j^{(1)})^2 + (x_i^{(2)} - x_j^{(2)})^2$$

$$= \sum_{d=1}^{M} (x_i^{(d)} - x_j^{(d)})^2$$



## Clustering: Similarity measure

- Clustering: partition data into groups such that each group contains "similar" objects
  - Increase intra-group similarity
  - Similar objects should belong to same cluster
- How do we measure "similarity" between objects?
  - "closeness"
- Which points are "closest" or "most similar"?
- Choice of similarity measure is dependent on the nature of data and your application



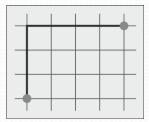
## Clustering: Similarity measure

- How to estimate the distance (i.e. similarity) between two objects/points p and q?
  - Various measures can be used
- K-means algorithm is flexible to use any distance measure

Euclidean distance  $(x_2, y_2)$  $(x_1, y_1)$ 

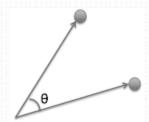
$$D_{e}(\mathbf{p}, \mathbf{q}) = \sqrt{\sum_{d=1}^{M} (p^{(d)} - q^{(d)})^{2}}$$

Manhattan distance



$$D_m(\mathbf{p}, \mathbf{q}) = \sum_{d=1}^{M} |p^{(d)} - q^{(d)}|$$

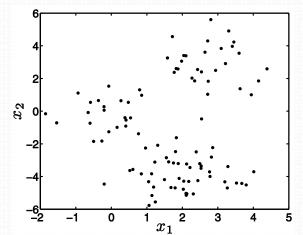
Cosine angle



$$D_{c}(\mathbf{p}, \mathbf{q}) = \cos^{-1} \left( \frac{\sum_{d=1}^{M} p^{(d)} q^{(d)}}{\sqrt{\sum_{d=1}^{M} p^{(d)^{2}}} \sqrt{\sum_{d=1}^{M} q^{(d)^{2}}}} \right) / \pi$$
 7

- Consider that there are total K clusters
- In 2D, each cluster k is represented by a cluster centre (i.e. mean)

$$\boldsymbol{\mu}_k = \left[\mu_k^{(1)}, \mu_k^{(2)}\right]^T$$



- Each object  $x_n$  is assigned to its closest cluster k on the basis of its distance to cluster mean  $\mu_k$
- Distance to cluster k can be estimated in various ways
  - For example: squared Euclidean distance

$$D_{nk} = (\boldsymbol{x}_n - \boldsymbol{\mu}_k)^T (\boldsymbol{x}_n - \boldsymbol{\mu}_k)$$

- No analytical solution for finding the closest cluster mean  $\mu_k$  for all objects
  - Iterative solution

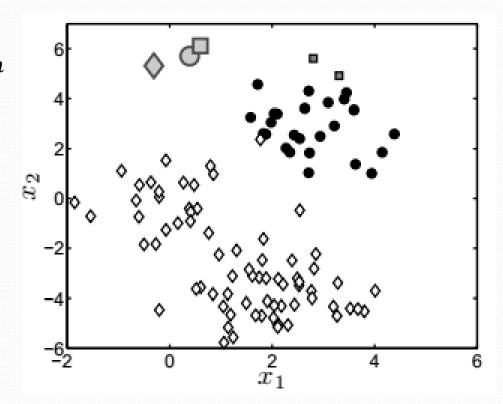
- Iterative algorithm: <u>Method 1</u>
  - 1. Randomly initialize cluster means  $\mu_1, \mu_2, ..., \mu_K$
  - 2. Compute distances  $D_{n1}, D_{n2}, ..., D_{nK}$  for each object  $x_n$  to all K cluster means
  - 3. Assign the object  $x_n$  to cluster k with lowest distance  $D_{nk}$ 
    - i.e. assign  $x_n$  to its closest cluster k with mean  $\mu_k$
  - 4. Update each cluster mean  $\mu_k$ , to represent the mean of newly updated cluster

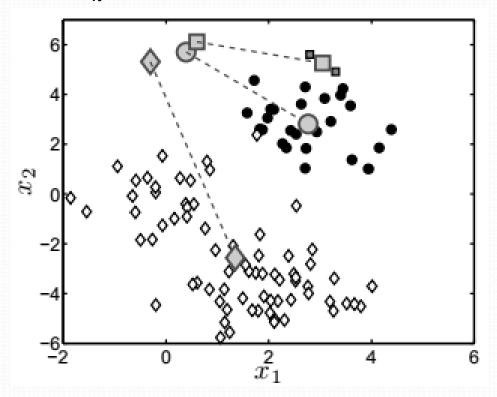
$$\mu_k = \frac{1}{N_k} \sum_{n=1}^{N_k} x_n$$

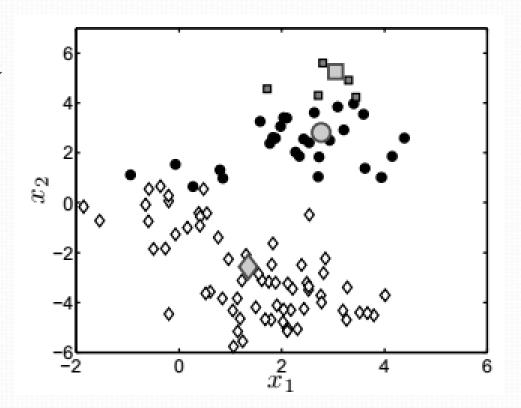
where  $N_k$  is the number of objects in  $k^{th}$  cluster

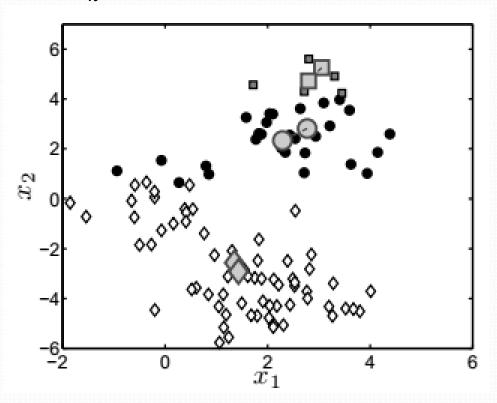
5. Stop if assignments don't change, or reached max number of iterations, else jump to step 2

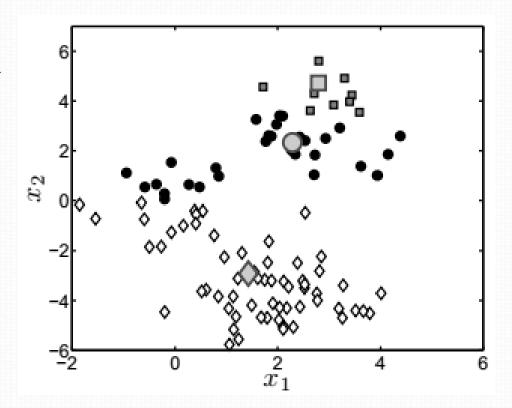
- Randomly initialize cluster means  $\mu_1, \mu_2, ..., \mu_k$
- Assign each  $x_n$  to its closest cluster with mean  $\mu_k$

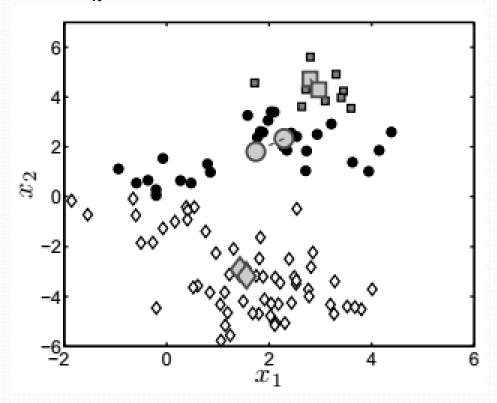


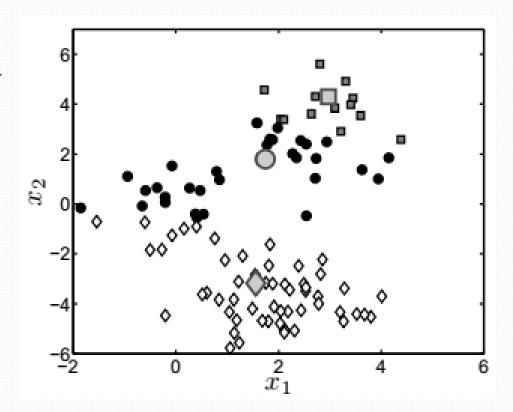


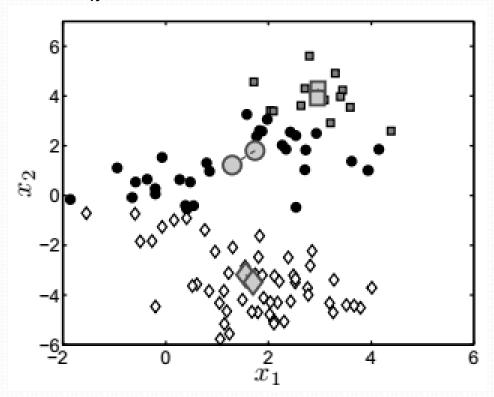


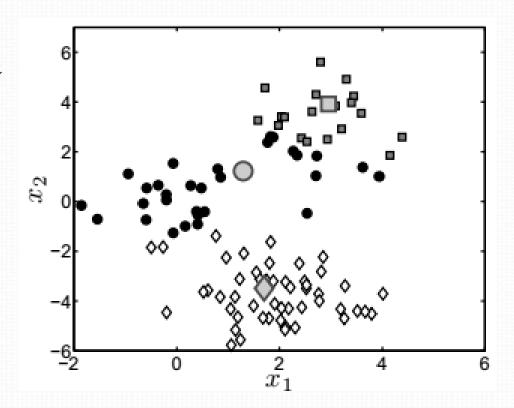


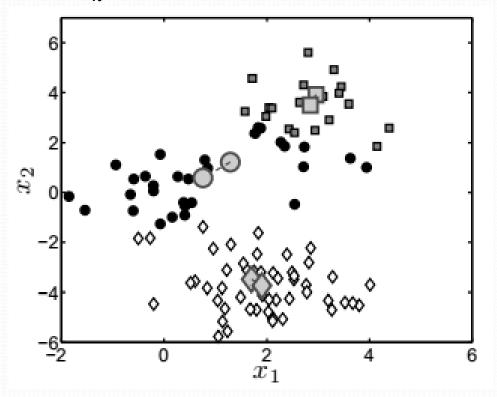


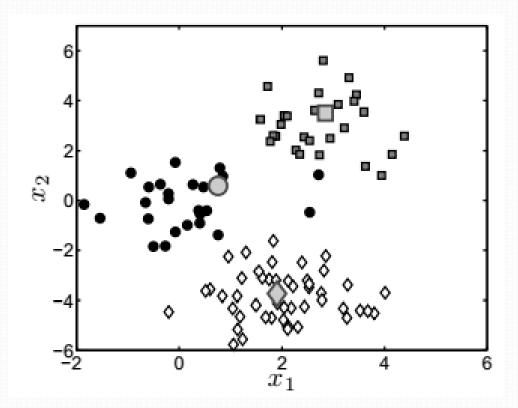


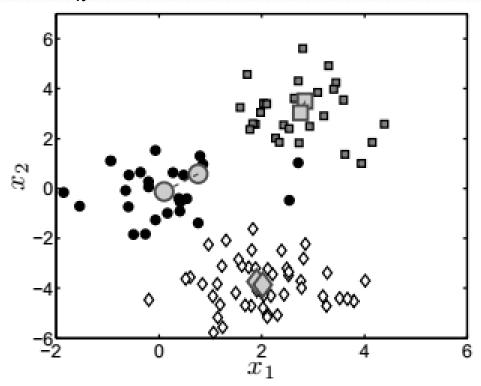


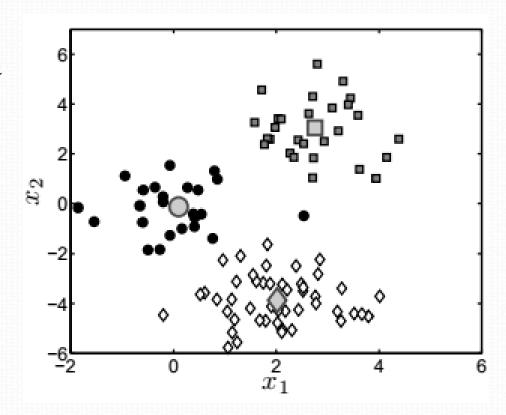


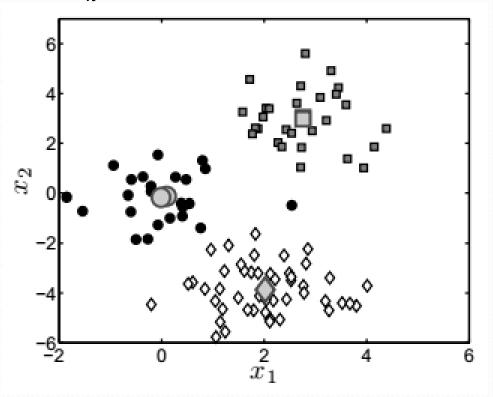




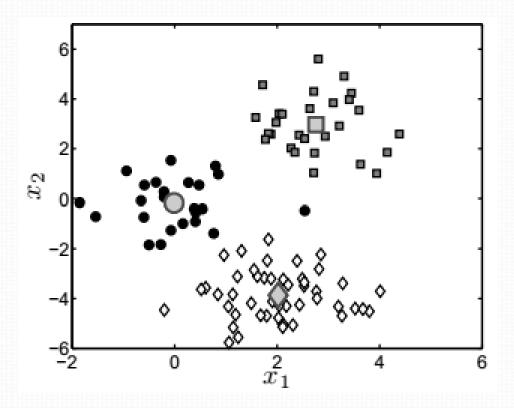








Solution at convergence



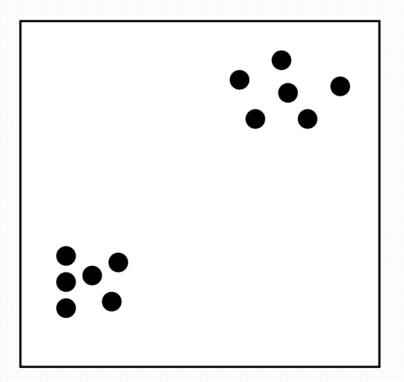
- Iterative algorithm: Method 2
  - 1. Randomly assign each object  $x_n$  to one of K clusters
  - 2. Compute cluster means  $\mu_1, \mu_2, ..., \mu_K$  for each cluster k, to represent the mean of newly updated cluster

$$\mu_k = \frac{1}{N_k} \sum_{n=1}^{N_k} x_n$$

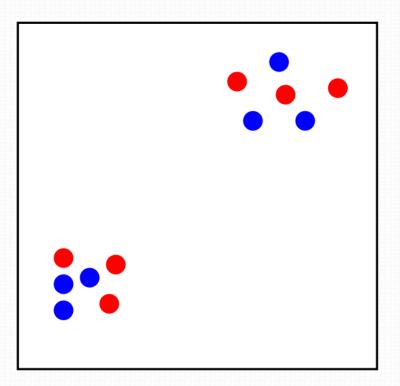
where  $N_k$  is the number of objects in  $k^{th}$  cluster

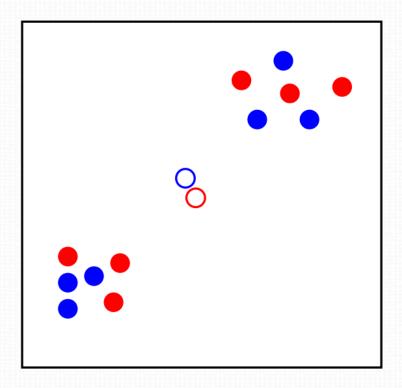
- 3. Compute distances  $D_{n1}, D_{n2}, ..., D_{nK}$  for each object  $x_n$  to all K cluster means
- 4. Assign the object  $x_n$  to cluster k with lowest distance  $D_{nk}$ 
  - i.e. assign  $x_n$  to its closest cluster k with mean  $\mu_k$
- Stop iterations if assignments don't change, or maximum number of iterations reached, else jump to step 2

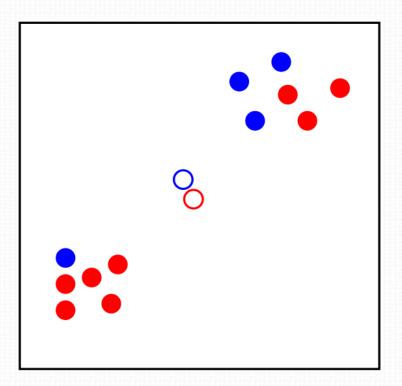
Cluster data points with Method 2

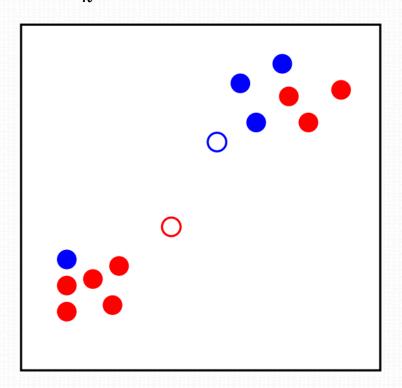


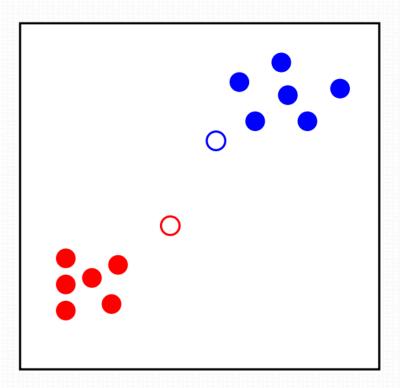
• Randomly assign each object  $x_n$  to one of K clusters

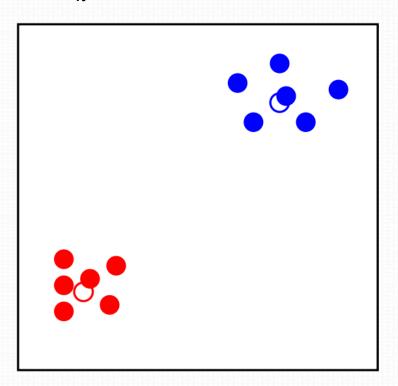












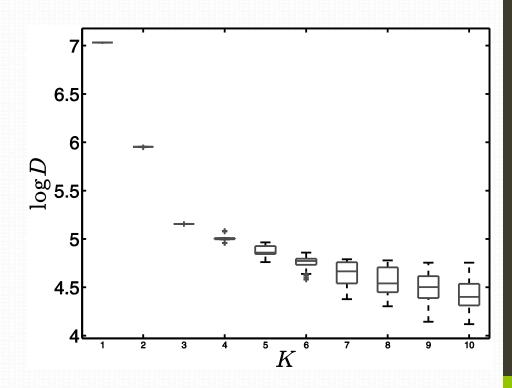
 The k-means clustering algorithm converges to a local minimum of the following intra-group distance estimate:

$$\mathcal{D} = \sum_{k=1}^{K} \sum_{n=1}^{N_k} (x_n - \mu_k)^T (x_n - \mu_k)$$

- i.e. total distance between all the objects and their respective cluster means
- Global optimum vs local optimum
  - Depends on random initialization
- Repeat runs of k-means clustering, pick one with lowest  $\mathcal{D}$

#### How to choose K?

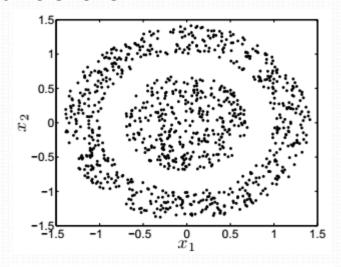
- Model selection
- 50 repeat runs
- D decreases with increasing K
  - Mhh5
- N = K

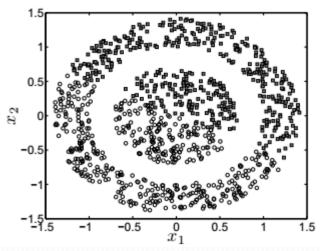


 K often depends on application/problem/data

#### K-means limitations

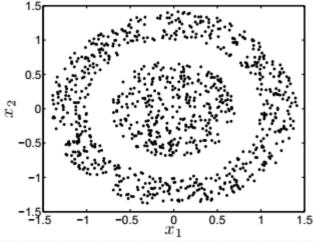
- K-means assumes clear cluster structure
  - What if it's lacking?
- Cluster means?
- Do objects conform to the similarity concept that K-means clustering heavily relies on?
- Outer cluster can't be represented by a single cluster mean





#### Data transformation

- How about transforming data in to a new space where it can be separable?
  - $x \to \phi(x)$
- For this example, consider  $\phi(x) = x_n^{(1)^2} + x_n^{(2)^2}$ ?
- In practice, there is a very neat trick which doesn't even require the explicit data transformation
  - Kernel trick
- Kernel function (a kind of similarity measure)
  - A function that is equivalent to the dot product of vectors in the transformed space
  - $k(\mathbf{x}_m, \mathbf{x}_n) = \phi(\mathbf{x}_m)^T \phi(\mathbf{x}_n)$



#### Kernel function

- There is a number of off-the-shelf kernels that have been shown to work well
- Linear kernel

• 
$$k(\mathbf{x}_m, \mathbf{x}_n) = \mathbf{x}_m^T \mathbf{x}_n$$

- Gaussian kernel
  - $k(\mathbf{x}_m, \mathbf{x}_n) = exp\{-\gamma(\mathbf{x}_m \mathbf{x}_n)^T(\mathbf{x}_m \mathbf{x}_n)\}$
  - $k(x_m, x_n) = exp\{-\gamma ||x_m x_n||^2\}$
- Polynomial kernel
  - $k(\mathbf{x}_m, \mathbf{x}_n) = (\mathbf{x}_m^T \mathbf{x}_n + c)^{\beta}$

- Can we kernelize k-means?
- Kernel function
  - A function that corresponds to inner product of vectors in some other transformed space
- As long as an algorithm has data appearing only in inner products in model learning, kernels can be used

$$k(\mathbf{x}_m, \mathbf{x}_n) = \phi(\mathbf{x}_m)^T \phi(\mathbf{x}_n)$$

Distance estimate (squared Euclidean)

$$D_{nk} = (\boldsymbol{x}_n - \boldsymbol{\mu}_k)^T (\boldsymbol{x}_n - \boldsymbol{\mu}_k)$$

Cluster mean calculation

$$\mu_k = \frac{1}{N_k} \sum_{n=1}^{N_k} x_n$$

Distance can be estimated as:

$$D_{nk} = \left(x_n - \frac{1}{N_k} \sum_{m=1}^{N_k} x_m\right)^T \left(x_n - \frac{1}{N_k} \sum_{m=1}^{N_k} x_m\right)$$

$$D_{nk} = x_n^T x_n - \frac{2}{N_k} \sum_{m=1}^{N_k} x_m^T x_n + \left(\frac{1}{N_k}\right)^2 \sum_{m=1}^{N_k} \sum_{l=1}^{N_k} x_m^T x_l$$

Distance can be estimated as:

$$D_{nk} = \mathbf{x}_n^T \mathbf{x}_n - \frac{2}{N_k} \sum_{m=1}^{N_k} \mathbf{x}_m^T \mathbf{x}_n + \left(\frac{1}{N_k}\right)^2 \sum_{m=1}^{N_k} \sum_{l=1}^{N_k} \mathbf{x}_m^T \mathbf{x}_l$$

With kernel trick, dot products could be replaced:

 $D_{nk}$ 

$$= k(\mathbf{x}_n, \mathbf{x}_n) - \frac{2}{N_k} \sum_{m=1}^{N_k} k(\mathbf{x}_m, \mathbf{x}_n) + \left(\frac{1}{N_k}\right)^2 \sum_{m=1}^{N_k} \sum_{l=1}^{N_k} k(\mathbf{x}_m, \mathbf{x}_l)$$

• Let's recall  $\mu_k$  will be

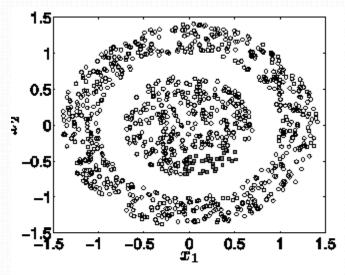
$$\mu_k = \frac{1}{N_k} \sum_{n=1}^{N_k} \phi(x_n)$$

but what's  $\phi(x_n)$ ?

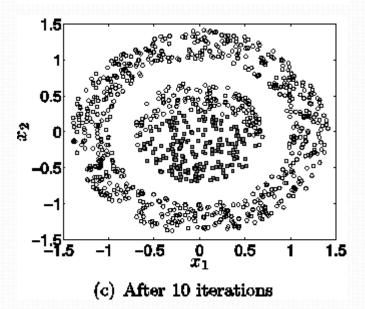
- Algorithm:
  - 1. Randomly assign each object  $x_n$  to one of K clusters
  - 2. Compute distances  $D_{n1}, D_{n2}, ..., D_{nK}$  for each object  $x_n$ , using the kernel trick

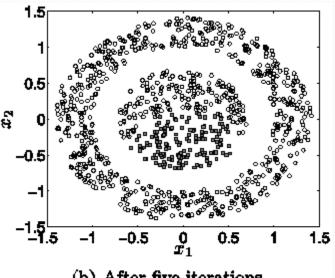
$$D_{nk} = k(\mathbf{x}_n, \mathbf{x}_n) - \frac{2}{N_k} \sum_{m=1}^{N_k} k(\mathbf{x}_m, \mathbf{x}_n) + \left(\frac{1}{N_k}\right)^2 \sum_{m=1}^{N_k} \sum_{l=1}^{N_k} k(\mathbf{x}_m, \mathbf{x}_l)$$

- 3. Assign the object  $x_n$  to cluster k with lowest distance  $D_{nk}$ 
  - ullet i.e. assign  $x_n$  to its closest cluster
- 4. Stop iterations if assignments don't change, or maximum number of iterations reached, else jump to step 2

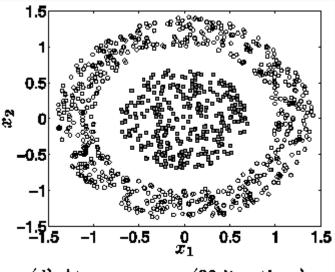


(a) Kernel K-means after one iteration





(b) After five iterations



(d) At convergence (30 iterations)

- Brings flexibility to simple k-means algorithm
- Need to set additional kernel parameters
- Sensitive to initialization
  - Often, it's good to run standard k-means algorithm before kernelized k-means for decent initialization

#### K-means: limitations

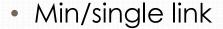
- Sensitive to initialization
- Computationally expensive
- Often needs repeated runs
- Makes hard assignments (i.e. each object can belong only to one cluster)
  - Soft assignment (each object has a probability of belonging to each cluster)

- Bottom-up (agglomerative) approach
  - Each object starts as a singleton cluster
  - Two most similar clusters are merged iteratively
  - Stop when all objects are in same cluster
- Top-down (divisive) approach
  - All objects belong to same cluster
  - "outsider" objects from least cohesive cluster are removed iteratively
  - Stop when each object is a singleton cluster

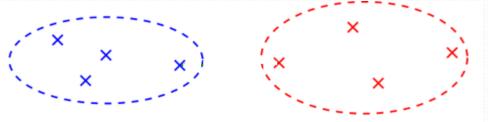
# Clustering: similarity measure

- We know how to estimate similarity (or distance between pair of objects
- How do we measure similarity between a pair of clusters (R and S)?



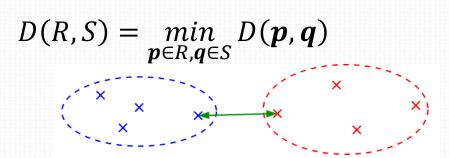


- Max/complete link
- Average link

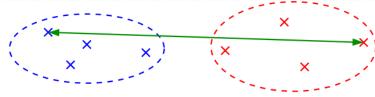


# Clustering: similarity measure

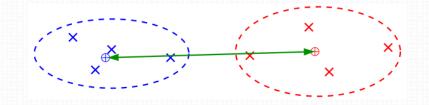
- Min/single link:
  - Clusters can get very large
- Max/complete link:
  - Results in small, round clusters
- Average link:
  - Compromise between min and max links



$$D(R,S) = \max_{\boldsymbol{p} \in R, \boldsymbol{q} \in S} D(\boldsymbol{p}, \boldsymbol{q})$$

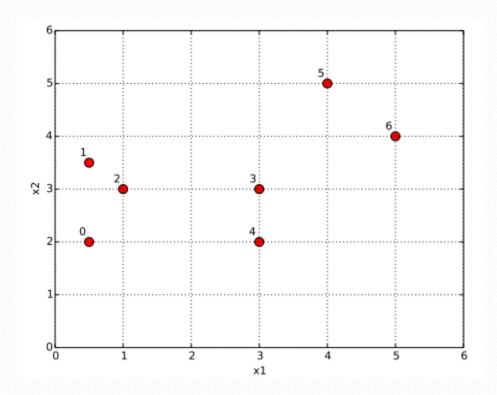


$$D(R,S) = \frac{1}{|R||S|} \sum_{\boldsymbol{p} \in R, \boldsymbol{q} \in S} D(\boldsymbol{p}, \boldsymbol{q})$$

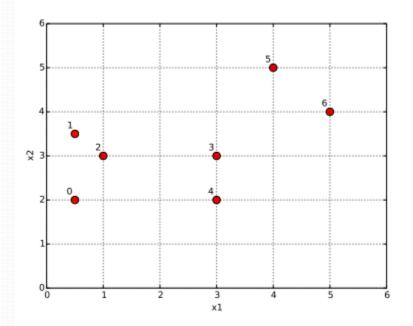


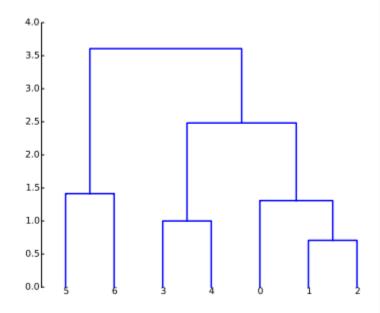
- Hierarchical agglomerative clustering (HAC) is a bottom-up clustering strategy
  - Each object is a singleton cluster
  - Successively merge closest pair of clusters together, till all clusters merge
- 1. Consider each object as a singleton cluster
- 2. Compute distance between all pairs of clusters
- Remove the closest pair of clusters from the data, and replace them as a single cluster
- 4. If not all objects belong to same single cluster, jump to step 2

Hierarchically cluster



- The results of hierarchical clustering can be visualized as a <u>dendrogram</u> where the height of each "junction" in the dendrogram represents the distance between the pair of clusters
- Hierarchical clustering generates all number of clusters
  - Cut horizontally across the dendrogram





# Flat clustering vs hierarchical clustering

- Flat clustering (K-means) results in a single partitioning of objects
- Hierarchical clustering can produce different partitioning results depending on the level-ofresolution we are looking at
- Flat clustering needs to know K in advance
- Hierarchical clustering doesn't need to know K

# Flat clustering vs hierarchical clustering

- Flat clustering is usually more efficient (run-time) if we know K
  - NxK distance computations at each iteration
- Hierarchical clustering can be slow (has to make several merge/split decisions)
  - Choosing 2 closest objects from N objects requires  $\frac{N!}{2!(N-2)!} => N*(N-1)/2 \text{ calculations just for first pairing}$
- No clear consensus on which of the two produces better clustering
  - Results depend upon nature of data

### Summary

- Two common clustering methods
  - By associating an object with its nearest cluster mean
  - By successively merging smaller clusters to hierarchically form larger clusters
- Free choices
  - How many clusters?
  - What distance metric?
  - What linkage method?

- Optimal combination is often found by trial-and-error
- Kernelized k-means

- Use K-means algorithm and Euclidean distance metric to cluster the following 2-dimensional objects in to 3 clusters
  - O1 (2,10), O2 (2,5), O3 (8,4), O4 (5,8), O5 (7,5), O6 (6,4), O7 (1,2), O8 (4,9)
- Suppose that initial cluster centres are O1, O4, and O7. Run the K-means algorithm steps for 1 iteration and show:
  - The clusters (i.e. objects belonging to each cluster)
  - The centres of new clusters
  - Draw a 10x10 grid with all the 8 objects and show the clusters and centres after the first iteration

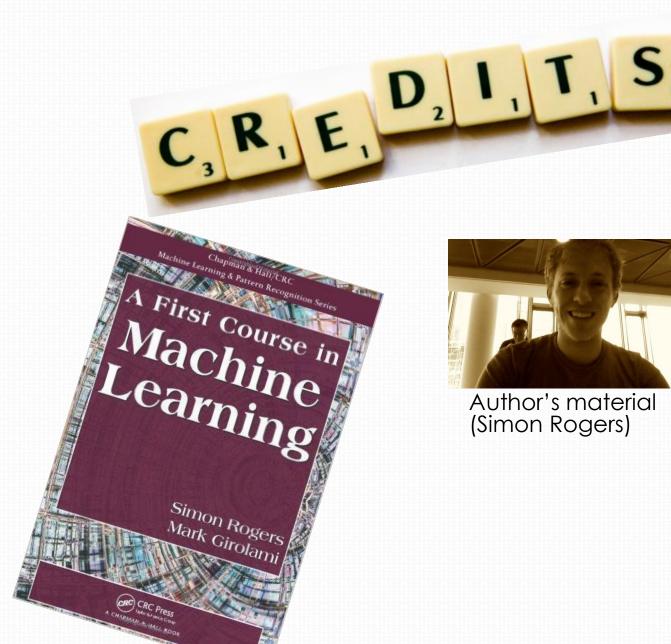
- Use min/single link to perform agglomerative clustering by showing the dendrogram
  - The data is described by the distance matrix

	Α	В	С	D
Α	0	1	4	5
В		0	2	6
С			0	3
D				0

- Use max/complete link to perform agglomerative clustering by showing the dendrogram
  - The data is described by the distance matrix
- Note: the height of each "junction" in the dendrogram represents the distance between the pair of clusters

- Use single and complete link agglomerative clustering to cluster the following 8 objects by showing the dendrograms
  - O1 (2,10), O2 (2,5), O3 (8,4), O4 (5,8), O5 (7,5), O6 (6,4), O7 (1,2), O8 (4,9)

- Try MATLAB code kmeansexample.m (from FCML book website)
- Try MATLAB code kmeansK.m (from FCML book website)
- Try MATLAB code kernelkmeans.m (from FCML book website)
- Try MATLAB code kmeans\_cluster.m (from Canvas)
- Try MATLAB code run\_kmeans.m (from Canvas)



Iain Styles



# Thankyou