Lecture 18: Unsupervised Learning

COMP90049

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Qiuhong Ke, CIS

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Roadmap

So far:

- · Supervised machine learning algorithms
- Train and evaluation the performance of the classifiers

Today:

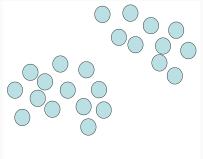
- Introduction of clustering (unsupervised learning)
- · K-means clustering
- · Hierarchical clustering
- · Evaluate clustering performance



Introduction

What is clustering

- Unsupervised learning: The class of an example is not known (or at least not used)
- Goal: find groups of similar data points
- "Similar" measure: e.g., small Euclidean distance.

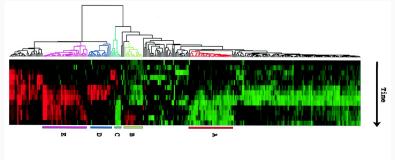




Why clustering I

Applications:

· Gene expression data analysis



Eisen, M.B. et al. Cluster analysis and display of genome-wide expression patterns. Proceedings the National Academy of Sciences, 95(25), 1998, pp.14863-14868.



Why clustering II

Applications:

· Image segmentation



(a) original image



(b) segmentation output (2 clusters)



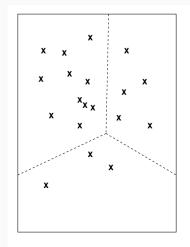
(c) segmentation output (3 clusters)

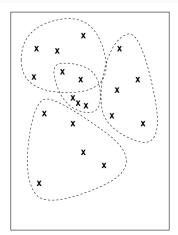
· Identify groups of customer



Exclusive vs. overlapping clustering

· Can an item be in more than one cluster?







Deterministic vs. probabilistic clustering

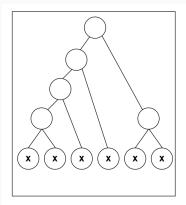
• Can an item be partially or weakly in a cluster?

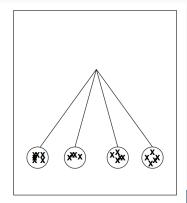
Instance	Cluster		Cluster				
IIIStarice		-	Instance	1	2	3	4
1	2		1	0.01	0.87	0.12	0.00
2	3		2	0.05	0.25	0.67	0.03
3	2		3	0.00	0.98	0.02	0.00
4	1		4	0.45	0.39	0.08	0.08
5	2		5	0.01	0.99	0.00	0.00
6	2		6	0.07	0.75	0.08	0.10
7	4		7	0.23	0.10	0.20	0.47
:	:						• • • • • • • • • • • • • • • • • • • •
	•		:	:			



Hierarchical vs. partitioning clustering

 Do the clusters have subset relationships between them? e.g. nested in a tree?

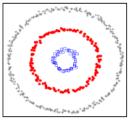


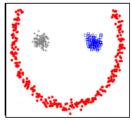


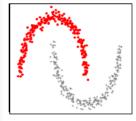


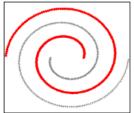
Heterogenous vs. homogenous clustering

· Clusters of widely different sizes, shapes, and densities











Clustering, basic contrasts

- · Exclusive vs. overlapping clustering
 - Can an item be in more than one cluster?
- Deterministic vs. probabilistic clustering (Hard vs. soft clustering)
 - Can an item be partially or weakly in a cluster?
- · Hierarchical vs. partitioning clustering
 - Do the clusters have subset relationships between them? e.g. nested in a tree?
- · Heterogenous vs. homogenous
 - · Clusters of widely different sizes, shapes, and densities
- · Partial vs. complete clustering
 - · In some cases, we only want to cluster some of the data
- · Incremental vs. batch clustering
 - · Is the whole set of items clustered in one go?



k-means

k-means Clustering

Given k (the number of clusters), the *k*-means algorithm is implemented in four steps:

- 1. Initialize: Select random k points to act as seed cluster centroids
- 2. Iterate:
 - Step 1 Cluster assignment: Assign each instance to the cluster with the nearest centroid
 - Step 2 Recompute the cluster centroids: Update centroid of each cluster to the average of its assigned instances
- 3. Until the centroids don't change
 - · Exclusive, deterministic, partitioning, batch clustering method

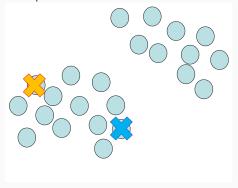


Measuring Similarity / Proximity / Closeness

- · Data points in Euclidean space
 - · Euclidean distance
 - · Manhattan (L1) distance
- · Discrete values
 - · Hamming distance: discrepancy between the bit strings
- · Other measures
 - · Cosine similarity
 - · Jaccard measure

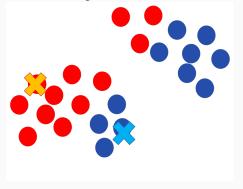


Pick 2 (k = 2) random points as cluster centroids.



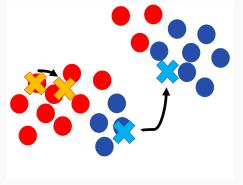


(Iteration 1) Step 1: Cluster assignment



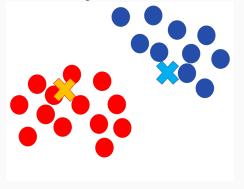


(Iteration 1) Step 2: Recompute the cluster centroids



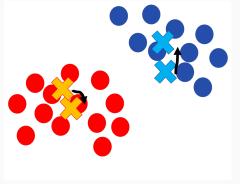


(Iteration 2) Step 1: Cluster assignment



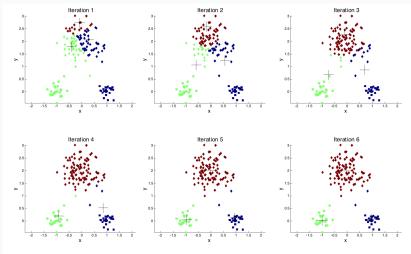


(Iteration 2) Step 2: Recompute the cluster centroids



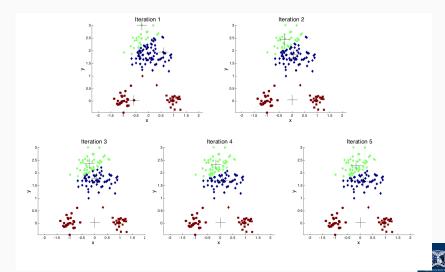


Example, Iterations

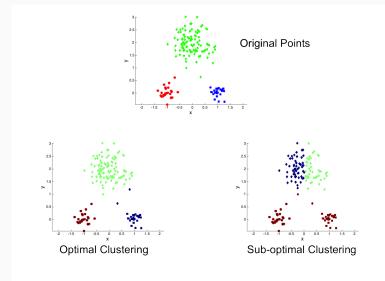




Example, Impact of initial seeds



Example, Different outcomes





Pros of *k*-means

- · relatively efficient:
 - O(ndki), where n is no. instances, d is no. attributes, k is no. clusters, and i is no. iterations; normally k, i ≪ n
 - Unfortunately we cannot a priori know the value of i!
- · can be extended to hierarchical clustering



Cons of k-means

- results sensitive to random centroid selection:
 - · try multiple iterations with different seeds
 - try better initialization method (k-means++)
- "mean" ill-defined for nominal or categorical attributes
- · may not work well when the data contains outliers
- not able to handle non-convex clusters, or clusters of differing densities or sizes





• need to specify k in advance.

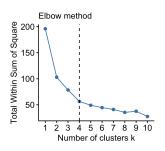


How to choose the number of clusters?

 calculate within-cluster Sum of Squared Error SSE_w for different number of clusters K

$$SSE_w = \sum_{i=1}^K \sum_{x \in C_i} (x - m_i)^2$$

- x is a data point in cluster C_i and m_i is the centroid of cluster C_i .
- As K increases, we will have a smaller number of instances in each cluster → SSE_w decreases
- Elbow method: K increases to K+1, the drop of SSE_w starts to diminish





Hierarchical Clustering

Hierarchical Clustering

Bottom-up (= agglomerative) clustering

- · Start with single-instance clusters
- At each step, join the two "closest" clusters (in terms of margin between clusters, distance between mean, ...)

Top-down (= divisive) clustering

- · Start with one universal cluster
- · Find two partitioning clusters
- · Proceed recursively on each subset

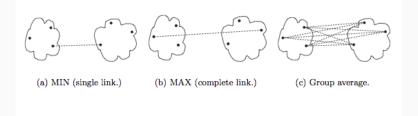


Bottom-up (Agglomerative) Clustering

- 1. Each point starts as a cluster. Compute the proximity matrix.
- 2. repeat
- 3. Merge the closest two clusters
- 4. Update the proximity matrix to reflect the proximity between the new cluster and the original clusters
- 5. until Only one cluster remains



Graph-based measure of Proximity



Updating the proximity matrix:

- Single Link: *Minimum* distance between any two points in the two clusters. (most similar members)
- Complete Link: Maximum distance between any two points in the two clusters. (most dissimilar members)
- Group Average: Average distance between all points (pairwise).



Agglomerative Clustering Example

				4	
1	1.00	0.90	0.10	0.65 0.60 0.40 1.00 0.80	0.20
2	0.90	1.00	0.70	0.60	0.50
3	0.10	0.70	1.00	0.40	0.30
4	0.65	0.60	0.40	1.00	0.80
5	0.20	0.50	0.30	0.80	1.00

What are the two closest points?



Agglomerative Clustering Example

		2			
1	1.00	0.90	0.10	0.65	0.20
2	0.90	1.00	0.70	0.60	0.50
3	0.10	0.70	1.00	0.40	0.30
4	0.65	0.60	0.40	1.00	0.80
5	0.20	0.90 1.00 0.70 0.60 0.50	0.30	0.80	1.00

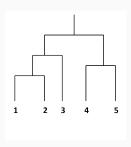
Merge points 1 & 2 into a new cluster: 6

Update (single link):

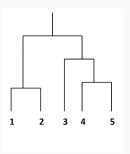
Update (complete link):



	1	2	3	4	5
1	1.00	0.90	0.10	0.65	0.20
2	0.90	1.00	0.10 0.70	0.60	0.50
3	0.10	0.70	1.00	0.40	0.30
4	0.65	0.60	0.40	1.00	0.80
5	0.20	0.50	1.00 0.40 0.30	0.80	1.00



Single link

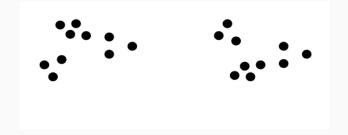


Complete link



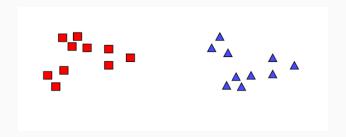
Evaluation

What is a good clustering?





Two clusters?





Four clusters?





Six clusters?





Types of Evaluation

Unsupervised:

- · cluster cohesion: compactness, tightness
- · cluster separation: isolation, distinctiveness.

Supervised: measure how well cluster labels match externally supplied class labels.

- entropy
- purity



Unsupervised Evaluation I

A "good" cluster should have one or both of:

- High Cluster Cohesion: instances in a given cluster should be closely related to each other
- High Cluster Separation instances in different clusters should be distinct from each other



Unsupervised Evaluation II

Within-cluster Sum of Squared Error SSE_w : the smaller, the better

$$SSE_w = \sum_{i=1}^K \sum_{x \in C_i} (x - m_i)^2$$

- x: a data point in cluster C_i
- *m_i*: the centroid for cluster *C_i*.

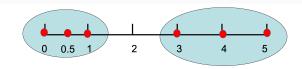
Between-cluster Sum of Squared Error SSE_b : the larger, the better

$$SSE_b = \sum_{i=1}^K n_i (m_i - m)^2$$

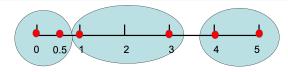
- m: mean of all data points in the dataset
- m_i : centroid for cluster C_i .
- n_i : number of instances in cluster C_i .



Example



$$SSE_w = 2.5, SSE_b = 18.375$$



$$SSE_w = ?, SSE_b = ?$$



Supervised Evaluation

Entropy: the smaller, the better

$$entropy = \sum_{i=1}^{k} \frac{|n_i|}{N} H_i$$

Purity: the larger, the better

$$purity = \sum_{i=1}^{k} \frac{|n_i|}{N} \max_{j} P_i(j)$$

- n_i : number of instances in cluster C_i .
- N: total number of instances in all clusters.
- $P_i(j)$: probability of the class j in the cluster i.
- H_i: entropy of class distribution in cluster i.

$$H_i = -\sum_j P_i(j) log_2 P_i(j)$$



Supervised Evaluation Example I

· Calculate the entropy and purity of the following cluster output

Cluster	Play = yes	Play = no
1	4	0
2	4	4

$$entropy_1 = -1 \times log(1) - 0 \times log(0) = 0$$

$$entropy_2 = -0.5 \times log(0.5) - 0.5 \times log(0.5) = 1$$

$$purity_1 = max(1,0) = 1$$

$$purity_2 = max(0.5,0.5) = 0.5$$

$$entropy = \frac{4}{12} \times 0 + \frac{8}{12} \times 1 = 0.67$$

$$purity = \frac{4}{12} \times 1 + \frac{8}{12} \times 0.5 = 0.67$$



Supervised Evaluation Example II

· Calculate the entropy and purity of the following cluster output

Cluster	Play = yes	Play = no
1	2	0
2	6	4



Summary

- · What basic contrasts are there in different clustering methods?
- How does k-means operate, and what are its strengths and weaknesses?
- What is hierarchical clustering, and how does it differ from partitioning clustering?
- · How to evaluation the clustering performance?



References

- Tan, Steinbach, Kumar (2006) Introduction to Data Mining. Chapter 8, Cluster Analysis
 - http://www-users.cs.umn.edu/~kumar/dmbook/ch8.pdf
- Jain, Dubes (1988) Algorithms for Clustering Data. http://homepages.inf.ed.ac.uk/rbf/BOOKS/JAIN/Clustering_Jain_Dubes.pdf

