

Data Clustering: A Very Brief Overview

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Outline

- Introduction
- Five Ws of Clustering
Who, What, When, Where, Why?
- One H of Clustering
How?
- Algorithms
- Conclusion

Introduction

- Unsupervised Learning: a very important problem in machine learning
 - Big amount of data
 - Unlabeled data
 - Time and effort to label
 - Not enough information to label
- Data Mining: an interdisciplinary field in computer science
 - A very large set of data in a database
 - Intersection of
 - Machine learning
 - Database systems

Introduction

- Some examples
 - Classification of plants given their features
 - Finding patterns in a DNA sequence
 - Recognizing objects, actions from images
 - Image segmentation
 - Document classification
 - Customer shopping patterns
 - Analyzing web searching patterns

5Ws of Clustering

- *Who, What, When, Where, Why?*
- As a researcher, you are given a (large) set of points without labels
- Grouping unlabeled data
 - Points within each cluster should be similar (close) to each other
 - Points from different clusters should be dissimilar (far)

5Ws of Clustering

- Given points are usually in a high-dimensional space
- Similarity is defined using a distance measure
 - Euclidean Distance,
 - Mahalanobis Distance,
 - Minkowski Distance,
 - ...

1H of Clustering

- *How do we cluster?*
- In general two types of algorithms:
 - Partition Algorithms
 - Obtain a single level of *partition*
 - Hierarchical Algorithms
 - Obtain a *hierarchy* of clusters

Partition Algorithms

- K-Means

- Set the number of clusters (k)

- Initialize k centroids

- Group points close to centroid

$$\sum_{i=0}^N \min_{\mu_j \in C} (\|x_i - m_j\|^2)$$

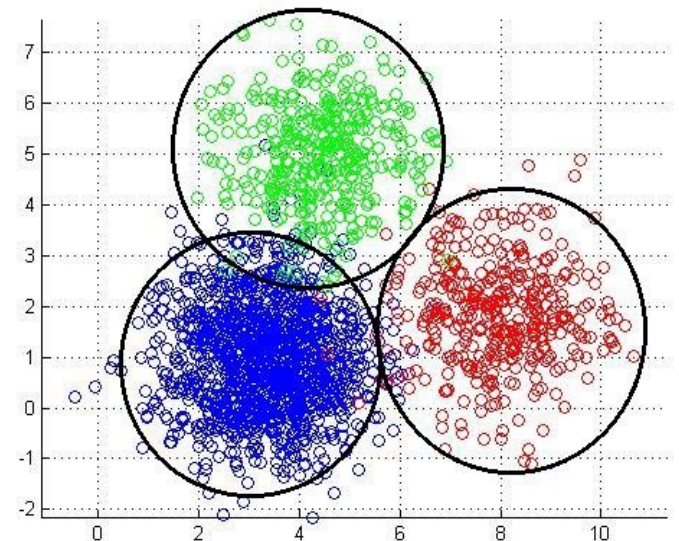
- Re-calculate centroids

- Always converges (may be to local minimum)

- Kmeans++

- Not highly scalable, Computation

- Minibatch K-means



Partition Algorithms

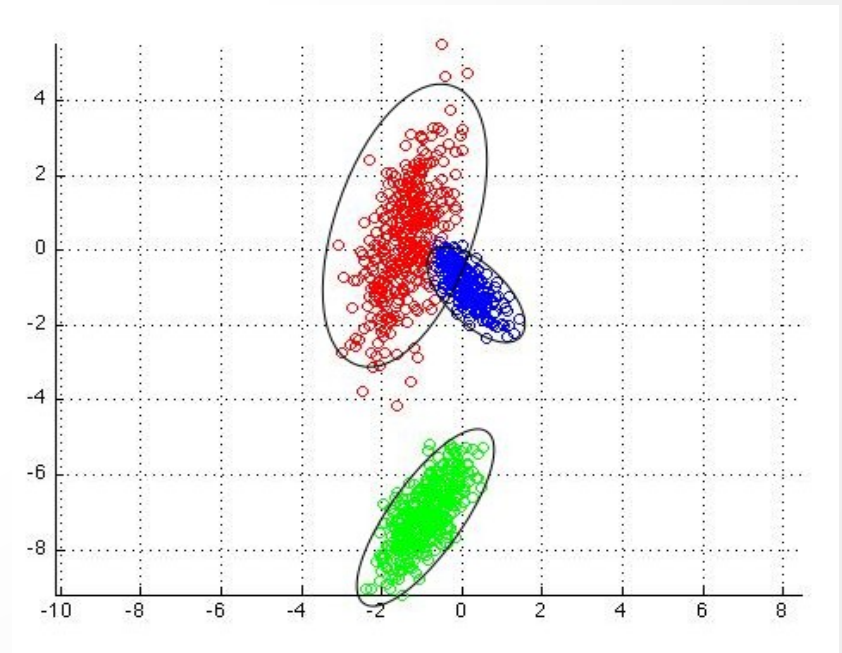
- Mean Shift
 - Set the bandwidth (max. distance)

$$\|x_i - m_j\|^2 \leq BW^2$$

- Mixture of Gaussian
 - Mahalanobis distance

$$\sum_{i=0}^N \min_{\mu_j \in C} ((x_i - \mu_j)^T \Sigma_j^{-1} (x_i - \mu_j))$$

- Not highly scalable



Partition Algorithms

- Spectral Clustering

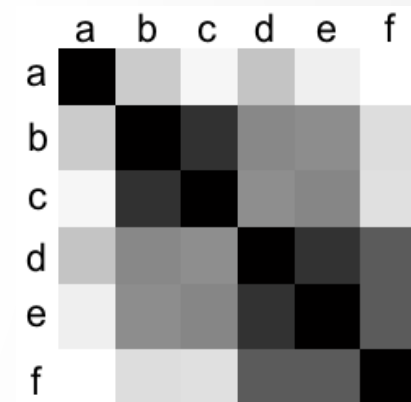
- Set the number of clusters (k)
- Similarity Matrix (pair-wise distance)

$$L = D - S \quad D_{ii} = \sum_j S_{ij}$$

- Laplacian Matrix

- Eigenvalues $0 = \lambda_1 \leq \dots \leq \lambda_n$

- Take first k eigenvectors and cluster using K-means



- Eigenvector computation could be a problem for large datasets

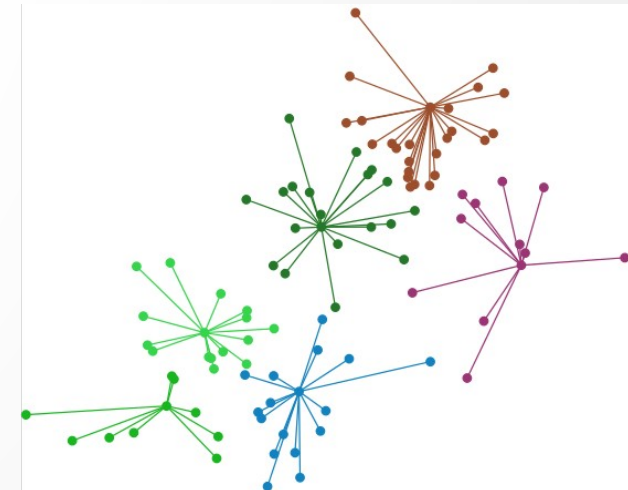
Partition Algorithms

- Affinity Propagation

- No need to specify number of clusters
- Similarity Matrix
- Responsibility Matrix
 - $r(i,k)$ -> Quantify how well x_k will be to serve as “exemplar” for x_i
- Availability Matrix
 - $a(i,k)$ -> Quantify how appropriate it will be for x_i to pick x_k as its “exemplar”
- “Message-passing” between data points
 - Initialize matrices R and A to zero
 - Iteratively update

$$r(i,k) \leftarrow s(i,k) - \max_{k' \neq k} \{a(i,k') + s(i,k')\}$$

$$a(i,k) \leftarrow \min \{0, r(k,k) + \sum_{i' \neq i,k} \max \{0, r(i',k)\}\}$$



Partition Algorithms

- Affinity Propagation
 - Computation complexity
 - Time
 - Memory
 - Not suitable for large datasets

How do we cluster?

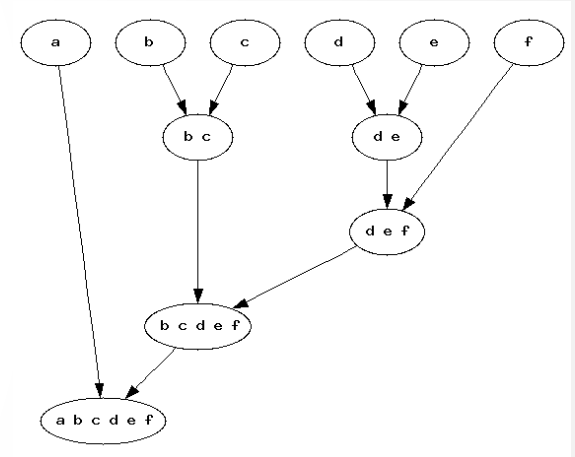
- In general two types of algorithms:
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Hierarchical Algorithms

- Bottom up – agglomerative
 - Iteratively merging small clusters into larger ones
- Top down – divide
 - Iteratively splitting larger clusters
- Can scale to large number of samples

Bottom up Algorithms

- Incrementally build larger clusters out of smaller clusters
 - Initially, each instance in its own cluster
 - Repeat:
 - Pick the two closest clusters
 - Merge them into a new cluster
 - Stop when there's only one cluster left
 - Obtain *dendrogram*



- Need to define “closeness” (metric and linkage criteria)

Bottom up Algorithms

- Linkage criteria
 - Ward: minimizing the sum of squared differences within all clusters (~K-means)
 - Single linkage: minimizes the distance between samples in a cluster (~K-NN)
 - Complete linkage: minimizes the maximum distance between samples in a cluster
 - Average linkage: minimizes the average of distances between samples in a cluster
- Distance Metric

Top down Algorithms

- Put all samples in one cluster and iteratively split the clusters
 - Distance metric to measure dissimilarity

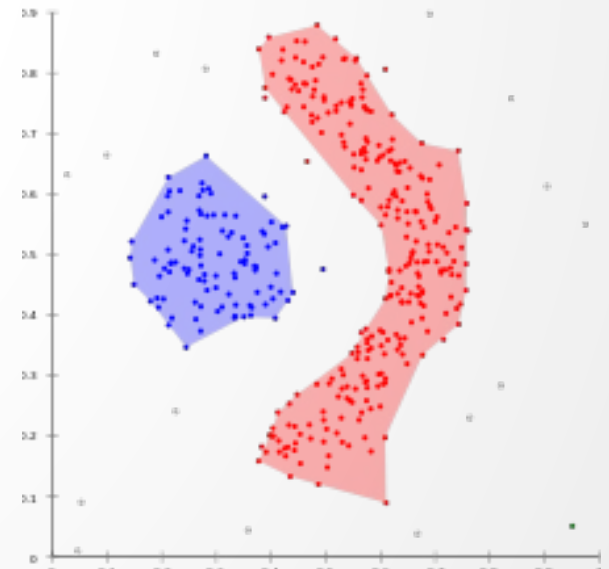
Other Algorithms

- DBSCAN*
 - Core samples: samples that are very close to each other
 - Non-core samples: samples that are close to core samples (except core samples themselves)
 - Set *epsilon* (ϵ) (distance) and *min. number of samples* to form a dense region
 - Take an arbitrary point
 - Check its ϵ -neighborhood
 - If it contains more samples than *min. number of samples*, create a cluster
 - If not mark as noise (outlier)

*Density-based spatial clustering of applications with noise

Other Algorithms

- DBSCAN
 - Can find arbitrarily shaped clusters
 - Can detect outliers
 - Can scale to very large datasets



Conclusion

- Clustering is a huge domain
- Need to select the approach suitable for the problem
 - Parameters to set (e.g., number of clusters)
 - Data geometry
 - Convergence: local / global optimum
 - Number of samples
 - Computation time

Conclusion

- Clustering performance evaluation

- Adjusted Rand Index

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

- Mutual Information

- Homogeneity, completeness

- Silhouette Coefficient

- Davies-Bouldin Index

$$DB = \frac{1}{n} \sum_{i=1}^n \max_{i \neq j} \left(\frac{\sigma_i + \sigma_j}{d(c_i, c_j)} \right)$$

- ...

THANK YOU

- References

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<http://scikit-learn.org/stable/modules/clustering.html>
- Anil K. Jain, M. N. Murty, and P. J. Flynn. “Data clustering: a review”, ACM Computing Surveys, 31(3):264–323, 1999
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- Brendan J. Frey and Delbert Dueck, “Clustering by Passing Messages Between Data Points”, Science Feb. 2007