

An Introduction to Some Popular Clustering Methods

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12 Aug 2015

Outline of the Talk

- 1 Introduction
- 2 Partition Based Clustering
- 3 Hierarchical Clustering

Clustering

- Type of unsupervised learning
- Seeks to group data into subsets
- Typically points within the same cluster are more closely related to each other than other points

- Partition
 - Given a non-empty set A , a partition of A is a collection of disjoint subsets of A whose union is A
- Clustering
 - Division of data into groups of similar objects [Berkhin(2006)]
 - Definition surprisingly useful, as it encompasses both hard partitions and soft partitions
- Unsupervised
 - No labelled data are available [Xu and Wunsch II(2009)]
- Dissimilarities
 - Typically, dissimilarities are metrics
 - Common example is the Euclidean distance

- Classification / taxonomy
[Everitt et al.(2001)Everitt, Landau, and Leese]
 - Psychology: personality types
 - Astronomy: star types
- Bioinformatics
 - Finding co-expressed genes from microarray data
- Business analytics
 - Grouping customers based on consumption patterns
 - Provide customized marketing strategies to each group

Biclustering of Microarray Data

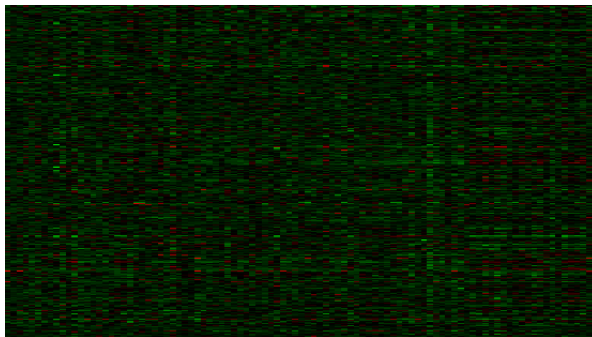


Figure: Biclustering of breast cancer microarray data [Hoshida et al.(2007)Hoshida, Brunet, Tamayo, Golub, and Mesirov] using the algorithm of Cheng and Church [Cheng and Church(2000)]. Rows of data matrix are 1213 genes, whilst the columns are the 97 samples. Colours range from bright green (negative, under-expressed) to bright red (positive, over-expressed).

Classification of Some Clustering Algorithms

[Berkhin(2006)]

- Partitioning Methods
 - k -means
 - k -medoids
- Fuzzy Partitioning Methods
- Hierarchical Methods
 - Agglomerative
 - Divisive
- Density based algorithms

The first versions of the k -means algorithm are attributed to Lloyd [Lloyd(1957)] and Forgy [Forgy(1965)]. The algorithm converges to a local optimum because both types of steps optimize the within-cluster sum of squares (WCSS) objective.

- 1 Each data point is assigned to closest centroid, with ties broken arbitrarily
- 2 The centroid positions are recomputed, based on the new memberships

Problems with k -means [Berkhin(2006)]

- Results dependent upon initialization of centroids
- Computed local optimum may be far from the global optimum
- Not obvious what value of k to use
- Process is sensitive to outliers
- Algorithm lacks scalability
- Only numerical data can be clustered
- Resulting clusters can be unbalanced

Clustering via Expectation-Maximization (EM)

- k -means is a limiting case of fitting data by a mixture of k Gaussians with identical, isotropic covariance matrices
- Soft assignment of data points to mixture components are hardened to label each data point using the most likely component.
- If data does not consist of well separated spherical clouds, k -means can have problems.
- EM clustering allows for “ellipsoidal” clouds of data
- Latent variable is class label
- Expectation step (E step): Calculate expected value of the log likelihood function
- Maximization step (M step): Find parameter that maximizes this quantity

Kernel k -means

- Kernel based methods enable us to deal with clusters which are not linearly separated
- The intuition behind the kernel method is to transform the original problem into a linearly separable problem
- Input data points are mapped nonlinearly into feature space via kernel function
- Kernel $K : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}$ measures similarity between any pair of inputs $\mathbf{x}, \mathbf{c} \in \mathcal{X}$
- For the often used RBF kernel, $K(\mathbf{x}, \mathbf{c}) = \exp(-\frac{\|\mathbf{x}-\mathbf{c}\|^2}{2\sigma^2})$

Kernel k -means Algorithm

- 1 Initialize cluster centers
- 2 Assign each data point to nearest cluster
- 3 Recalculate cluster centers
- 4 Repeat 2 and 3
- 5 Data point whose image is closest to the center is selected as representative of cluster

Finding k

- There are many methods for finding the number of clusters in an automated manner
- One can for instance try
 - Bootstrapping approach (`fpc::clusterboot`)
 - Bayesian approach

Hierarchical Clustering

- Agglomerative

- Each observation starts in its own cluster
- Pairs of clusters are merged as one moves up the hierarchy

- Divisive

- All observations start in one cluster
- Splits are performed recursively as one moves down the hierarchy

Linkage criteria

Linkage	Formula
complete	$\max\{d(a, b) : a \in A, b \in B\}$
single	$\min\{d(a, b) : a \in A, b \in B\}$
average	$\frac{1}{ A B } \sum_{a \in A} \sum_{b \in B} d(a, b)$
centroid	$\ c_i - c_j\ $, where c_i and c_j are centroids of clusters i and j

Table: Some linkage criteria between two sets of observations A and B .

Problems with Hierarchical Clustering

[Xu and Wunsch II(2009)]

- Lack of robustness
- Sensitivity to noise and outliers
- Once object is assigned to cluster, membership is not reconsidered
- Computational complexity, but there are efficient variants
[Murtagh(1983)]



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Acknowledgements

Many thanks to Vik Gopal and Alex You for helpful comments