

Clustering and Distance Methods

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Outline

- 1 Introduction
- 2 Comparing Data
- 3 Non Hierarchical Methods
- 4 Clustering Based on Statistical Models

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- Oftentimes multivariate data exhibits a natural grouping or clustering behavior that is challenging to analytically describe.
- The goal of clustering analysis is to find the natural groupings that exist within the data without requiring assumptions by the statistician.

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- It requires that the statistician choose a number of groupings before analysis. This can be problematic on large data sets as well as on data whose shape is not well known before.
- Clustering is a more base and general process that allows the data to naturally sift into groupings based on the user defined “distance”. We will focus on this method for this project.

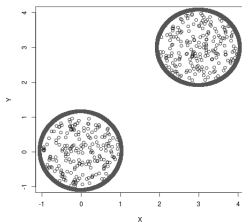
What is Distance?

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- We see that these two clusters of points have natural circular boundaries.

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- The problem is defining a meaningful metric to compare points.

Common Measures of Distance

- Euclidean distance: $d(x, y) = \sqrt{\sum_i (x_i - y_i)^2}$
- Minkowski Metric (Taxicab distance), $d(x, y) = [\sum_{i=1}^p |x_i - y_i|^m]^{1/m}$
- Canberra metric $d(x, y) = \sum_i \frac{|x_i - y_i|}{x_i + y_i}$

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- Other examples include hamming distance and the “max metric”

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So, d is a measurement of “difference” in the two variables

Similarity Coefficient Continued

In this way, we can write the comparisons in the table,

	$k = 1$	$k = 0$	Totals
$i = 1$	a	b	$a + b$
$i = 0$	c	d	$c + d$
Totals	$a + c$	$b + d$	$a + b + c + d$

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From this table, we can define many similarity coefficients depending on our desired measurement. These coefficients are then collected into matrices, upon which we can perform our more familiar statistical operations.

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 - Agglomerative (Start with n clusters and combine)
 - Divisive (Start with 1 cluster and partition)

Agglomerative Method Steps

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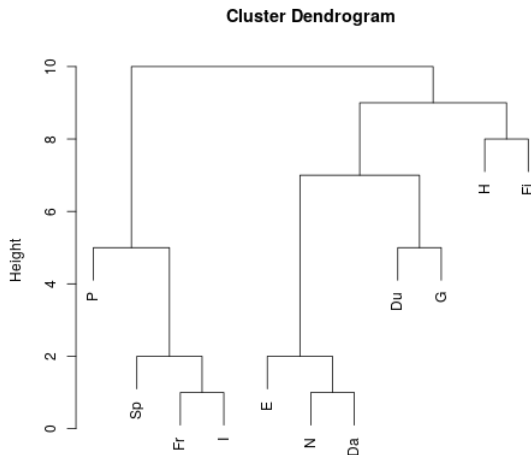
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- 4 Rinse and repeat $n - 1$ times. Keep a record of each merge as it happens for plotting purposes.

Dendrogram Example

Using the agglomerative method on Concordance data from several different languages, we arrive at,



This result shows that French and Italian are closely related, and that Danish and Norwegian are as well. To the French and Italian, we see that Spanish is also closely related, and so on.

Hierarchical Methods

When we compute the steps of these agglomerative methods, we must consider the different ways we can join the subgroups.

- Single Linkage - Find the smallest values in the distance matrix and join those groups together
- Complete Linkage - Much the same as the former, however our updated matrix uses the maximum distance between the subgroups when updating
- Average Linkage - Updating the matrix uses the average of all possible distances from the original grouping.
- Wards Method - Seeks to minimize ESS. Treats each cluster as an n-sphere and computes the distance to the centroid. Each successive merger results in the smallest possible ESS based on these centers and groupings.

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- Due to the nature of the methods, we have no statistically sound way of identifying and cleaning outliers from the data, which can result in bad clustering.
- There is also no way to change a points assignment once it has been merged into a new cluster. As such, points can be grouped incorrectly.
- Using multiple different methods and comparing the results is often the best way to decide if your results are stable and correct.

Nonhierarchical Methods

- We saw before that Hierarchical methods are designed to group variables into groups.
- We now consider methods that group items into an arbitrary number of groups by recursively updating.
- This is commonly done by choosing “seed” points at random within the space of the data.

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- 3 Update the affected centroids
- 4 Rinse and Repeat until each point is in the best possible centroid for itself

K-means Example

Consider the data,

Item	x_1	x_2
A	5	3
B	-1	1
C	1	-2
D	-3	-2

To perform a K-means clustering, let us at random choose an initial partition of (AB) and (CD) . We find the centroid of each of these to be,

Cluster	x_1	x_2
(AB)	$\frac{5-1}{2} = 2$	$\frac{3+1}{2} = 2$
(CD)	$\frac{1-3}{2} = -1$	$\frac{-2-2}{2} = -2$

So, we have our centroids, $(2, 2)$ and $(-1, -2)$. Next, we compute the distances between each point and the centroids,

$$d(A, (AB)) = \sqrt{10}$$

$$d(A, (CD)) = \sqrt{61}$$

$$d(B, (AB)) = \sqrt{10}$$

$$d(B, (CD)) = \sqrt{9}$$

Here, we see that B is actually closer to the centroid of (CD) , and as such, we shall move it, and update the centroids. Continuing,

$$d(C, (A)) = \sqrt{41}$$

$$d(C, (BCD)) = \sqrt{5}$$

$$d(D, (A)) = \sqrt{89}$$

$$d(D, (BCD)) = \sqrt{5}$$

K-means Continued

- We see now that none of the points need to be moved, and as such, we are done with our calculations.
- The resulting groups (A) and (BCD) make sense upon inspection, as BCD all lie at least partly within the negative ranges around the origin, while A sits alone in quadrant I with a ways between it and the other points.
- As before, it is recommended to re-run the algorithm with different initial conditions, to test stability.

- It is important to make sure that our initial conditions do not inadvertently overlap.

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- Outliers will tend to isolate themselves into their own groups.
- Even if you know the number of groups a population has, if the sampling method does not have points in each cluster, a forced number of clusters might have bad results.

Clustering Based on Statistical Models

- The methods that we have considered before have all been fairly intuitive in terms of the derivation of the steps, but there exist more statistically model based methods for analyzing the data.
- For example, consider the model where each cluster k has an expected proportion p_k of the points, and the associated pdf $f_k(x)$. Then, for k clusters,

$$f_{Mix}(x) = \sum p_k f_k(x)$$

- As one might expect, a common mixture model is a multivariate normal distribution.

Further Analysis of the Normal Model

- As one might expect, this normal distribution model will yield results close to the K-means and Ward's method processes.
- Choosing the number of clusters for an arbitrary data set can be challenging, but the formula

$$\#Clusters \propto -2 \ln(L_{max}) - Penalty$$

where L_{max} is defined as the product of our mixture functions.

- Under the AIC the penalty can be written as,

$$-2 \ln(L_{max}) - 2N\left(\frac{k}{2}(p+1)(p+2) - 1\right)$$

Multidimensional Scaling

- As one might expect, a natural consequence of looking at these high dimensional abstract data sets will be to attempt to project them into a lower dimensional form for graphical interpretation.
- This, however will not always be a simple procedure, as information is lost through the projection into lower dimensions, as well as the fact that trying to encode categorical data always results in loss of precision as well.
- As such, we might consider a process similar to PCA to attempt to look at the data while retaining as much information as possible.
- Multidimensional scaling looks at a slightly different problem than PCA, however, as the relative distances between the points should be retained when possible, as this can provide interesting analytical results on high dimensional data.

Multidimensional Scaling Continued

- Looking at N observations, there are $M = N(N - 1)/2$ similarities between points. Assuming the data is fine enough that equal values are not possible, it is trivially possible to order these distances,

$$s_1 < s_2 < \dots < s_M$$

As such we would expect that the associated distances to follow the form,

$$d_1^q > d_2^q > \dots > d_M^q$$

Where q is the desired dimensionality of the projection.

- For cases when a the transformation from s to d is not as simple, a computation of the Stress is used to align the data accordingly.
- An interesting consequence of this analysis is that on data about physical distances, the resultant plot will artificially start to resemble a map without the data ever being placed in that shape.

Correspondence Analysis

- Another way of graphically representing complex higher dimensional data is Correspondence analysis.
- This method is used for frequency data.
- By representing relative frequency in the form of stacked bar charts, both the categories and the levels can be compared simultaneously.

Correspondence Analysis Numerics

- To perform the correspondence analysis, we follow these steps,
 - 1 Construct $P = \frac{1}{n}X$
 - 2 Compute, $r = P'1_j$, $c = P'1_i$
 - 3 Then, $D_r = \text{diag}(r_1, r_2, \dots, r_i)$, $D_c = \text{diag}(c_1, c_2, \dots, c_j)$
 - 4 We also define the matrices $D^{\pm 1/2}$ as you might expect
 - 5 Then, a SVD is performed on,

$$A = D_r^{1/2}(P - rc')D_c^{-1/2}$$

- 6 Depending on the number and weight of the non-zero singular values, the original correspondence may be written as a lower order sum of the eigenvalues and outer product of the eigenvectors.

Summary

- When confronted with a large and high order data set, there are many options for how to analyze it. From a traditional approach of regression and multivariate extensions of classical statistical tests to PCA, there are many things that can be done.
- If the goal of the researcher is to isolate and naturally group the data, Clustering analysis provides many tools for finding the underlying structure in the data.
- While each method is not without its relative cost and risk, both computationally and accuracy wise, the relative speed and ease of interpretation of the results of clustering provide powerful tools for researchers to show their data to a larger group of people in a meaningful way.

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