INTRO TO DATA SCIENCE K-MEANS CLUSTERING

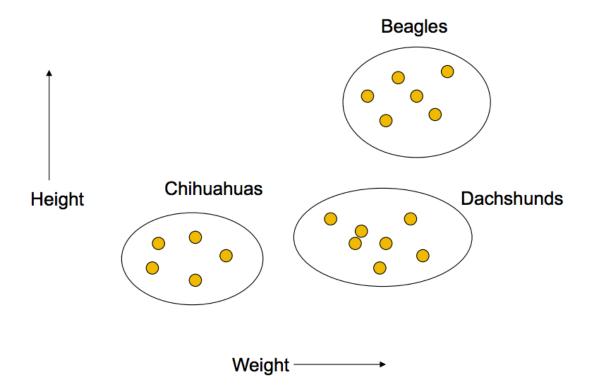
I. CLUSTER ANALYSIS

CLUSTERING

Given a set of data points, group them into clusters so that

- 1) Points within each cluster are similar to each other
- 2) Points from different clusters are dissimilar

EXAMPLE: DOGGIE DATA



UNDERSTANDING

- Genes with common functions
- Web search groupings (IR)
- Climate Pressure Patterns
- Astronomy Galaxies
- Medicine Disease subsets
- Business Customer Segments

UTILITY

- Data Sampling
- Compression
- NN Reduction

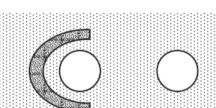
DIFFERENT TYPES OF CLUSTERS



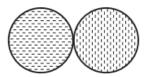


(a) Well-separated clusters. Each point is closer to all of the points in its cluster than to any point in another cluster.

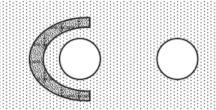




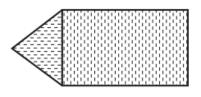
(c) Contiguity-based clusters. Each point is closer to at least one point in its cluster than to any point in another cluster.

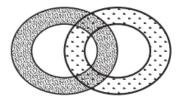


(b) Center-based clusters. Each point is closer to the center of its cluster than to the center of any other cluster.



(d) Density-based clusters. Clusters are regions of high density separated by regions of low density.





(e) Conceptual clusters. Points in a cluster share some general property that derives from the entire set of points. (Points in the intersection of the circles belong to both.)

K-MEANS

K-MEANS CLUSTERING

Q: What is k-means clustering?

A: A greedy learner that partitions a data set into k clusters.

greedy — captures local structure (depends on initial conditions)

partition — performs complete clustering (each point belongs to exactly one cluster)

BASIC K-MEANS ALGORITHM

- 1: Select K points as initial centroids.
- 2: repeat
- 3: Form K clusters by assigning each point to its closest centroid.
- Recompute the centroid of each cluster.
- 5: **until** Centroids do not change.

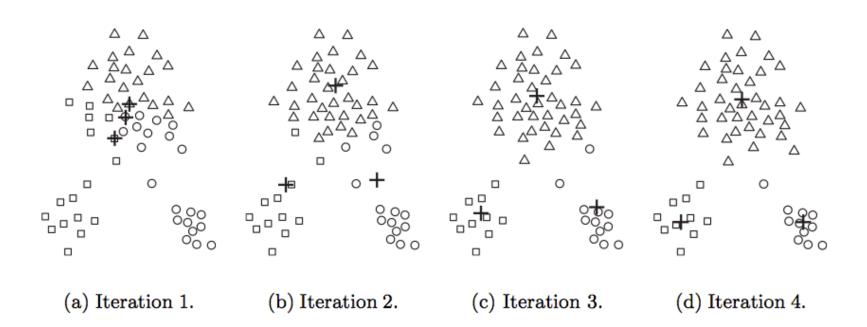


Figure 8.3. Using the K-means algorithm to find three clusters in sample data.

STRENGTHS & WEAKNESSES

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Difficulties can sometimes be overcome by increasing the value of k and combining subclusters in a post-processing step.

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 - perform alternative clustering task, use resulting centroids as initial k-means centroids
 - start with global centroid, choose point at max distance, repeat (but might select outlier)

STEP 2 - SIMILARITY MEASURES

Q: How do you determine which centroid is the nearest?

The "nearness" criterion is determined by the similarity/distance measure we discussed earlier.

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This measure makes quantitative inference possible.

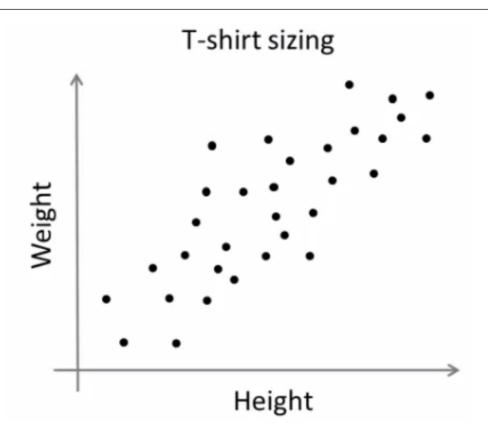
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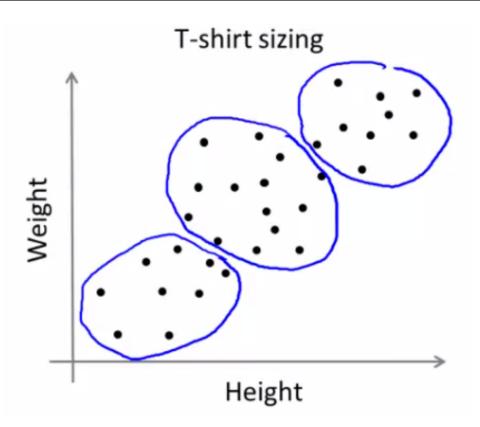
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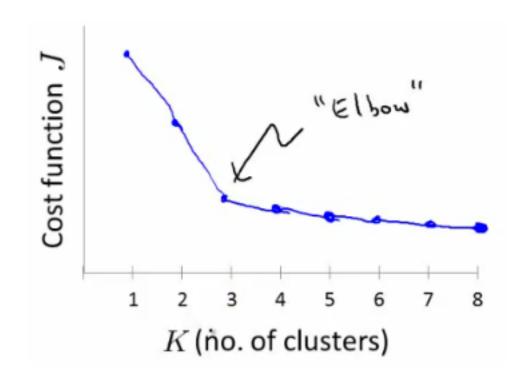
NOTE

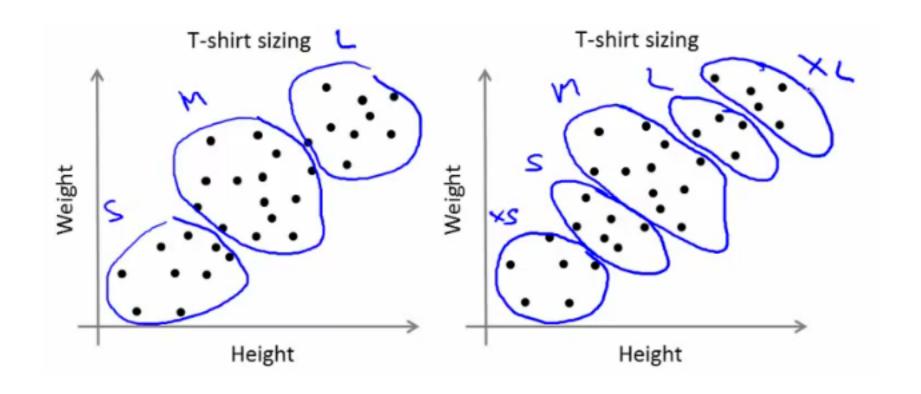
Technically, by defining a similarity measure we are mapping our observations into a *metric space*.

K-MEANS FOR NON SEPARATED CLUSTERS

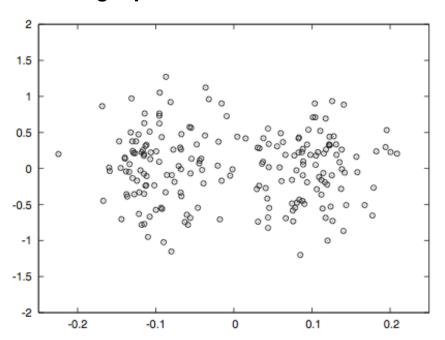


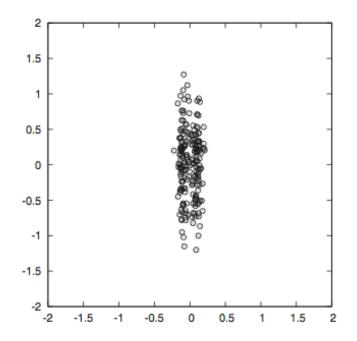






These graphs show two different representations of the same data:





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NOTE

Another useful property is smoothness.

(symmetry)

(triangle inequality)

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We can express different semantics about our data through the choice of metric.

Ex: One popular metric for text mining problems (or any problem with sparse binary data) is the **Jaccard coefficient**,

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Applying this metric to a problem expresses the sparse nature of the data, and makes a variety of text mining techniques accessible.

STEP 2 — SIMILARITY MEASURES

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For this reason, it's really the choice of metric that determines the definition of a cluster.

STEP 3 — OBJECTIVE FUNCTION

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A: By optimizing an **objective function** that tells us how "good" the clustering is.

The iterative part of the algorithm (recomputing centroids and reassigning points to clusters) explicitly tries to minimize this objective function.

Ex: Using the Euclidean distance measure, one typical objective function is the **sum of squared errors** from each point x to its centroid c_i :

$$SSE = \sum_{i=1}^{K} \sum_{x \in C_i} d(x, c_i)^2$$

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Given two clusterings, we will prefer the one with the lower SSE since this means the centroids have converged to better locations (a better local optimum).

STEP 4 — CONVERGENCE

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Recall that, in general, different runs of the algorithm will converge to different local optima (centroid configurations).

III. CLUSTER VALIDATION

In general, k-means will converge to a solution and return a partition of k clusters, even if no natural clusters exist in the data.

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We will look at two validation metrics useful for partitional clustering, **cohesion** and **separation**.

Cohesion measures clustering effectiveness within a cluster.

$$\hat{C}(C_i) = \sum_{x \in C_i} d(x, c_i)$$

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Separation measures clustering effectiveness between clusters.

$$\hat{S}(C_i, C_j) = d(c_i, c_j)$$

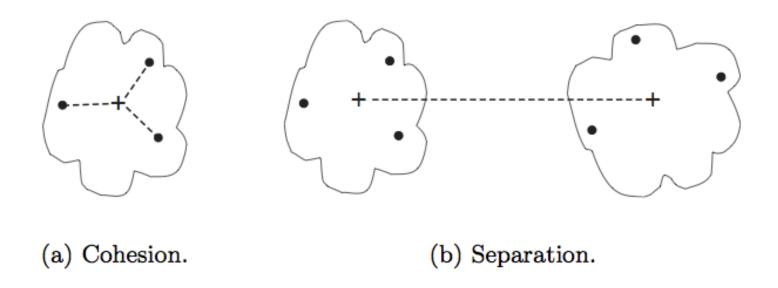


Figure 8.28. Prototype-based view of cluster cohesion and separation.

We can turn these values into overall measures of clustering validity by taking a weighted sum over clusters:

$$\hat{V}_{total} = \sum_{1}^{K} w_i \hat{V}(C_i)$$

Here V can be cohesion, separation, or some function of both.

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The weights can all be set to 1 (best for k-means), or proportional to the cluster *masses* (the number of points they contain).

Cluster validation measures can be used to identify clusters that should be split or merged, or to identify individual points with disproportionate effect on the overall clustering.

One useful measure than combines the ideas of cohesion and separation is the **silhouette coefficient**. For point x_i , this is given by:

$$SC_i = \frac{b_i - a_i}{max(a_i, b_i)}$$

such that:

 a_i = average in-cluster distance to x_i b_{ij} = average between-cluster distance to x_i b_i = $min_j(b_{ij})$ The silhouette coefficient can take values between -1 and 1.

In general, we want separation to be high and cohesion to be low. This corresponds to a value of SC close to +1.

A negative silhouette coefficient means the cluster radius is larger than the space between clusters, and thus clusters overlap.



Figure 8.29. Silhouette coefficients for points in ten clusters.

The silhouette coefficient for the cluster C_i is given by the average silhouette coefficient across all points in C_i :

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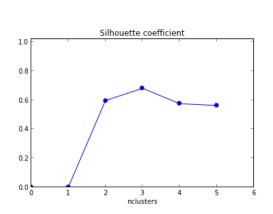
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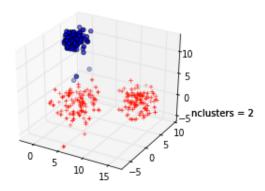
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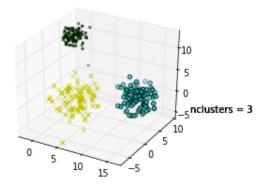
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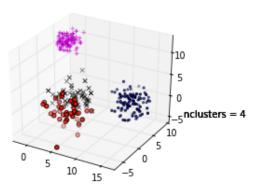
This gives a summary measure of the overall clustering quality.

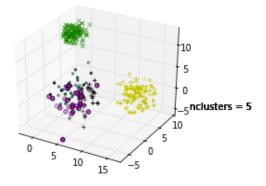
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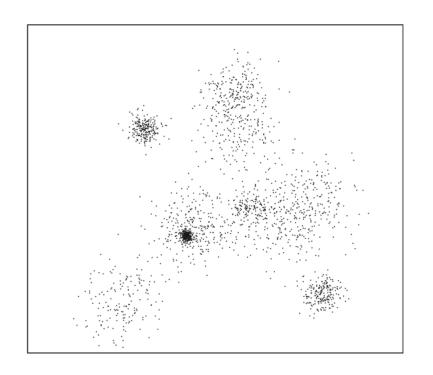


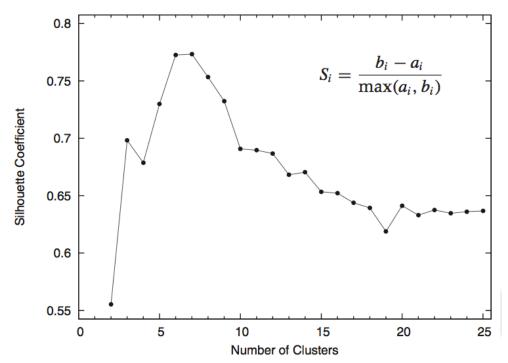






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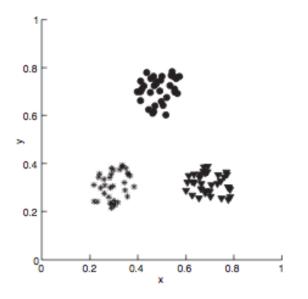




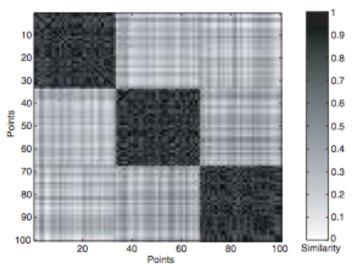
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This can be done either graphically or using correlations.



(a) Well-separated clusters.



(b) Similarity matrix sorted by K-means cluster labels.

One useful application of cluster validation is to determine the best number of clusters for your dataset.

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Q: How would you do this?

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Q: How would you do this?

A: By computing the overall SSE or SC for different values of k.

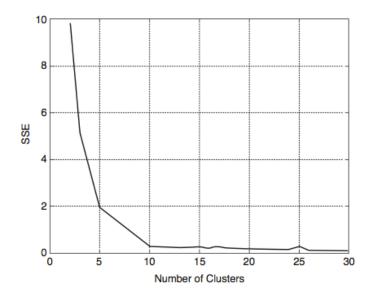


Figure 8.32. SSE versus number of clusters for the data of Figure 8.29.

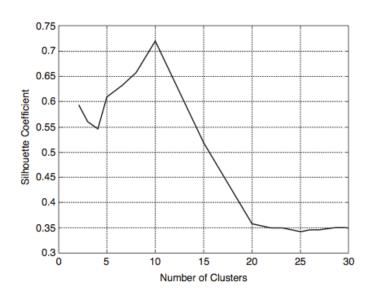


Figure 8.33. Average silhouette coefficient versus number of clusters for the data of Figure 8.29.

Q: How can you determine your level of confidence in these validation metrics?

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A: Statistically; eg, by computing frequency distributions for these metrics (over several runs of the algorithm) and determining statistical significance.

Ultimately, cluster validation and clustering in general are suggestive techniques that rely on human interpretation to be meaningful.

II. K-MEANS CLUSTERING