

INTRO to DATA SCIENCE

CLUSTER ANALYSIS

- I. CLUSTER ANALYSIS**
- II. THE K-MEANS ALGORITHM**
- III. CHOOSING K**
- IV. EXAMPLE**

I. CLUSTER ANALYSIS

	<i>continuous</i>	<i>categorical</i>
<i>supervised</i>	???	???
<i>unsupervised</i>	???	???

	<i>continuous</i>	<i>categorical</i>
<i>supervised</i>	<i>regression</i>	<i>classification</i>
<i>unsupervised</i>	<i>dimension reduction</i>	<i>clustering</i>

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In general, greater similarity between points leads to better clustering.

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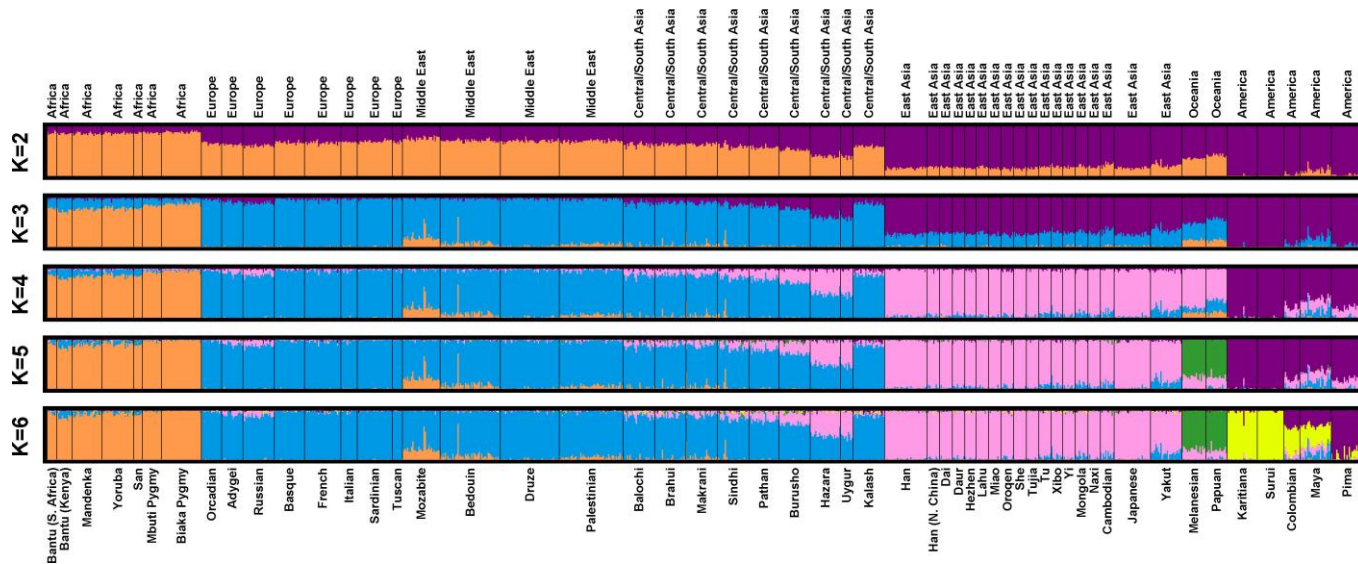
A: To enhance our understanding of a dataset by dividing the data into groups.

Clustering provides a layer of abstraction from individual data points.

The goal is to extract and enhance the natural structure of the data

Clustering can be useful in a wide variety of domains, including genetics, consumer internet and business.

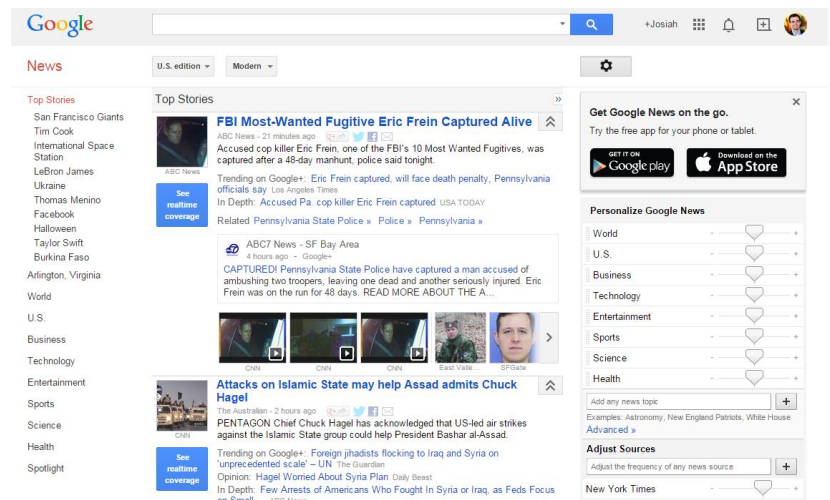
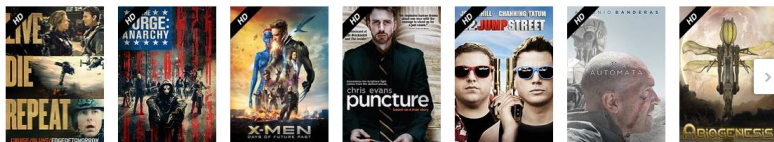
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There are many kinds of clustering procedures. For our class, we will be focusing on K-means clustering, which is one of the most popular clustering algorithms.

K-means is an iterative method that partitions a data set into k clusters.

II. K-MEANS CLUSTERING

Q: How does the algorithm work?

- 1) *choose k initial centroids (note that k is an input)*
- 2) *for each point:*
 - find distance to each centroid*
 - assign point to nearest centroid*
- 3) *recalculate centroid positions*
- 4) *repeat steps 2–3 until stopping criteria met*

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A: There are several options:

- randomly (but may yield divergent behavior)*
- 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)*

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In the case of k-means clustering, the similarity metric is the Euclidian distance:

$$d(x_1, x_2) = \sqrt{\sum_{i=1}^N (x_{1i} - x_{2i})^2}$$

Q: How do we recompute the positions of the centers at each iteration of the algorithm?

A: By calculating the centroid (i.e., the geometric center)

We iterate until some stopping criteria are met; in general, suitable convergence is achieved in a small number of steps.

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Stopping criteria can be based on the centroids (eg, if positions change by no more than ε) or on the points (eg, if no more than $x\%$ change clusters between iterations).

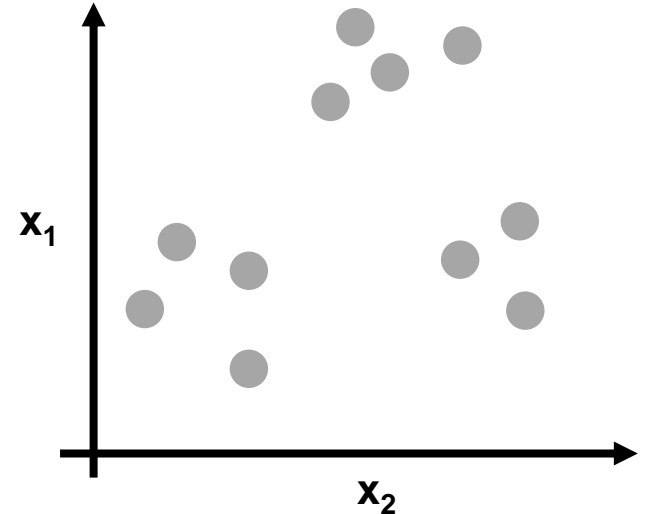
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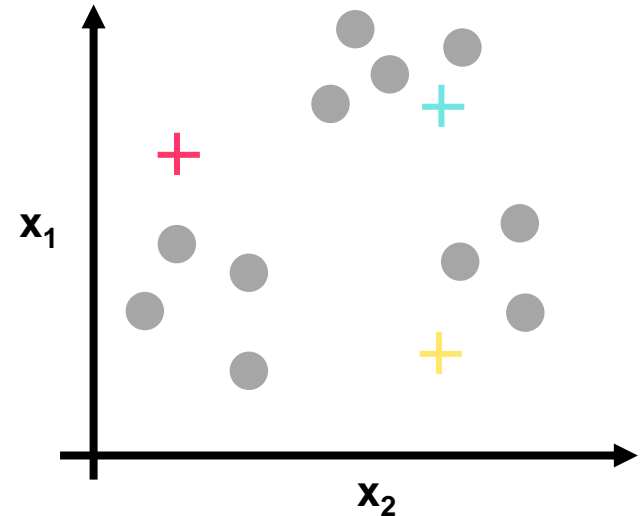
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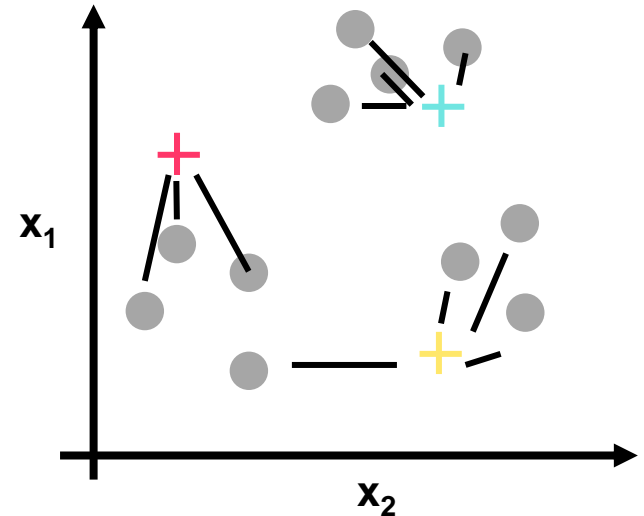
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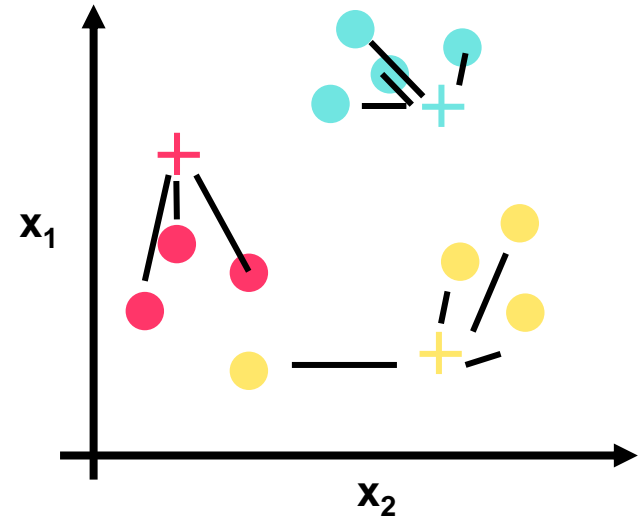
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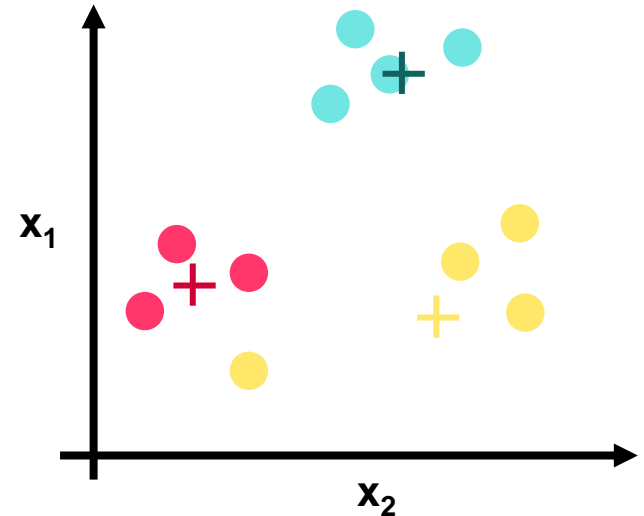
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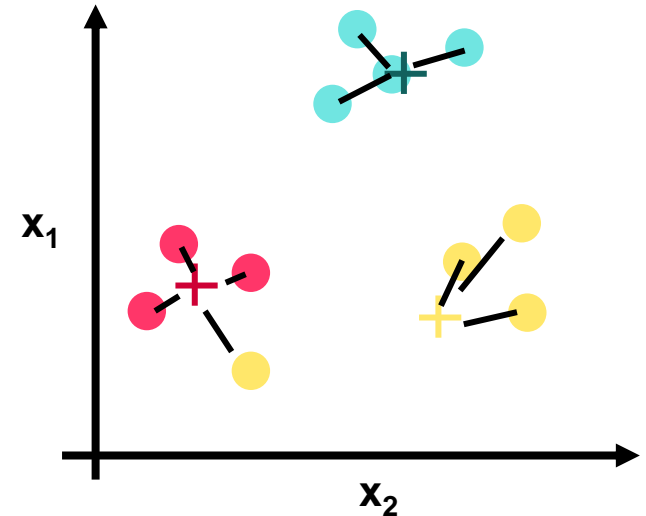
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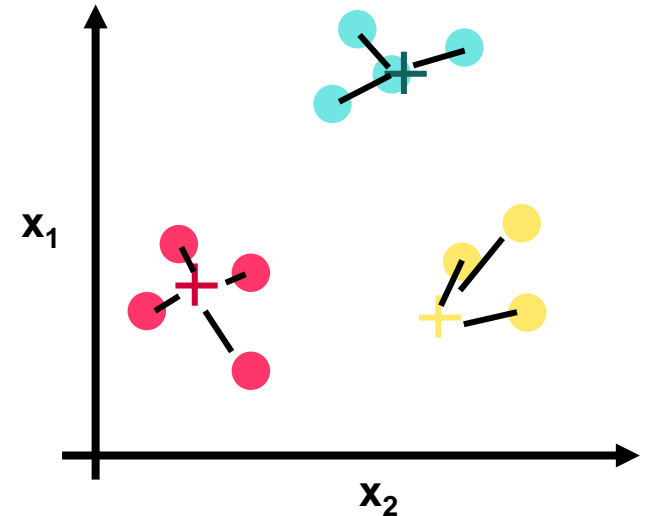
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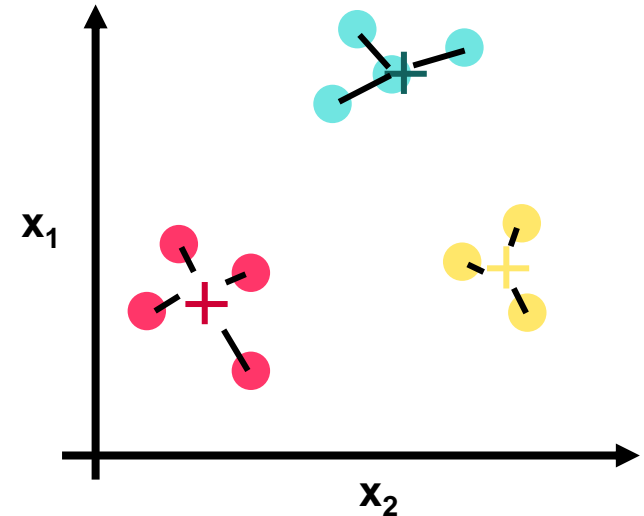
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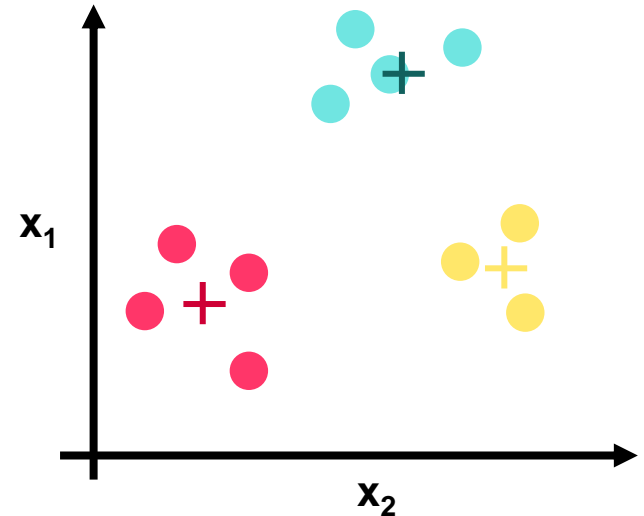
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III. CLUSTER VALIDATION

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.*

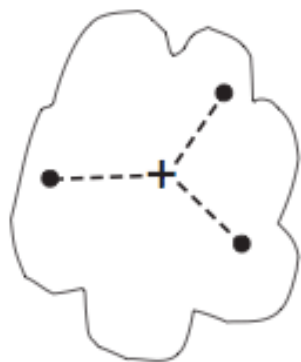
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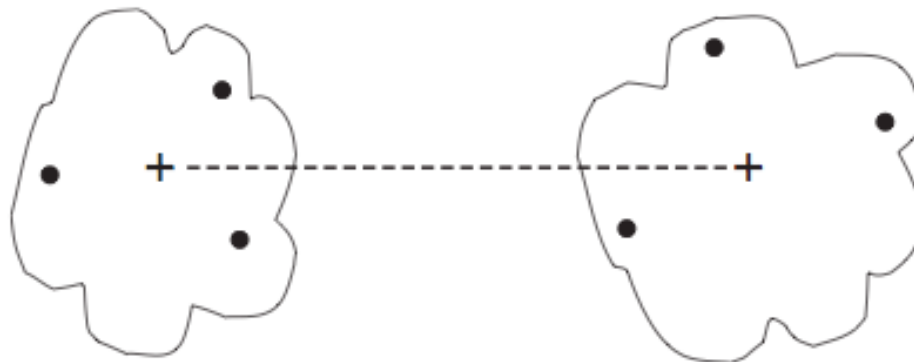
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Separation *measures clustering effectiveness between clusters.*

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




(a) Cohesion.



(b) Separation.


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

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

$$SC = \frac{b - a}{\max(a, b)}$$


a = The mean distance between a sample and all other points in the same class.

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

$$SC = \frac{b - a}{\max(a, b)}$$


b = The mean distance between a sample and all other points in the next nearest cluster.

Generally, this formula simplifies to the following:

$$SC = 1 - \frac{a}{b}$$

This is because generally $b > a$:

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.

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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NOTE

This gives a summary measure of the overall clustering quality.

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Q: How would you do this?

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

A: Another commonly used metric is the within sum of squared errors

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

Strengths:

K-means is a popular algorithm because of its computational efficiency and simple and intuitive nature.

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Weaknesses:

However, K-means is highly scale dependent, and may produce misleading results for data with widely varying shapes and densities.

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EX: K-MEANS CLUSTERING