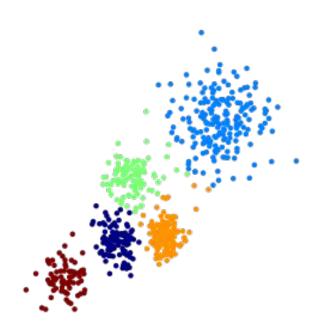


Clustering

K-means& hierarchical clustering

DSI SEA5, jf.omhover



Clustering

K-means& hierarchical clustering

DSI SEA5, if.omhover

OBJECTIVES



- Relate clustering to unsupervised learning
- Illustrate the utility of clustering in real-world problems
- Describe and implement the k-means algorithm
- Describe and implement the HAC algorithm
- Compare purpose and utility of k-means and HAC
- Discuss the role of metrics for applying clustering to different problems
- Analyze how the (high) dimensionality of data impacts metrics based clustering techniques



Supervised / Unsupervised Learning

Supervised Learning

 (x_1, y_1)

 (x_n, y_n)

x y



REALITY

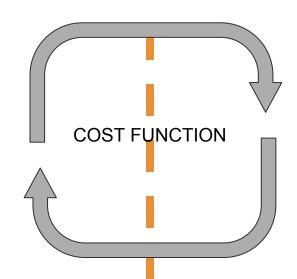
| | type | income | education | prestige |
|------------|------|--------|-----------|----------|
| accountant | prof | 62 | 86 | 82 |
| pilot | prof | 72 | 76 | 83 |
| architect | prof | 75 | 92 | 90 |
| author | prof | 55 | 90 | 76 |
| chemist | prof | 64 | 86 | 90 |
| minister | prof | 21 | 84 | 87 |
| professor | prof | 64 | 93 | 93 |
| dentist | prof | 80 | 100 | 90 |
| reporter | wc | 67 | 87 | 52 |
| engineer | prof | 72 | 86 | 88 |
| undertaker | prof | 42 | 74 | 57 |
| lawyer | prof | 76 | 98 | 89 |

data

OBJECTIVE:

descriptive predictive normative

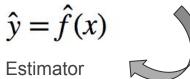
. . .



MODEL

$$y = f(x) + \epsilon$$

take a function as an assumption



of the function



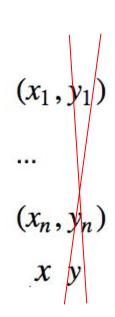
Unsupervised Learning



REALITY

| | type | income | education | prestige |
|------------|------|--------|-----------|----------|
| accountant | prof | 62 | 86 | 82 |
| pilot | prof | 72 | 76 | 83 |
| architect | prof | 75 | 92 | 90 |
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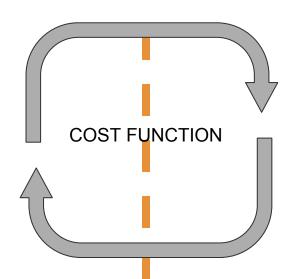
data



OBJECTIVE:

descriptive predictive normative

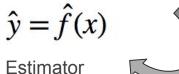
. . .



MODEL

$$y = f(x) + \epsilon$$

take a function as an assumption



of the function



Machine Learning: different types of learning



[Samuel, 1959]: Machine learning is the "field of study that gives computers the ability to learn without being explicitly programmed"

Supervised learning:

- The model is derived from observations of input/output pairs
- You have data samples with labelled output (quantitative / qualitative)

Unsupervised learning:

- The model is derived from the confrontation of a meta-model with observations
- You have data samples without no output class, and you want to explain or describe them (but you have an idea of what you're looking for)

Reinforcement learning:

- The model is derived from interactions with an external agent or environment



Unsupervised-type questions









- I have a database of clients with their purchase history and I want to draw profiles
- I have the proceedings of the last 2016 data science conference and I want to see the hot topics
- I have obtained usage traces of users on my GUI (clicks, forward/backward, inputs, time spent on each page etc) and I want to understand what different behavior and trajectories they may have
- I have this dataset of gene expressions and I want to extract groups of genes that have mutual influences
- I have this dataset of tweets on the presidential debate and
 I want my candidate to know which people were tweeting about what



Clustering

Brainstorm: what's a cluster?



dataset

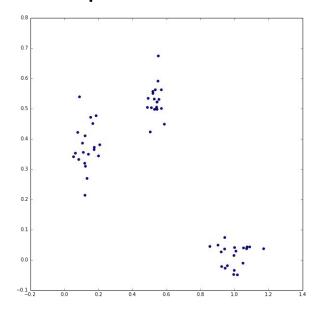
[[0.06497338 0.35259884] [0.20073913 0.34345291] [0.0540259 0.34076791] [0.1415917 0.34904249] [0.1066884 0.38564734] [0.07874009 0.42121996] [0.16728357 0.45044774] [0.18659554 0.47644782] [0.08462494 0.3317815] [0.17597371 0.37192779] [0.1547712 0.47111988] [0.089005 0.53872432] [0.11967159 0.3192513] [0.12118539 0.21355644] [0.17501382 0.36435908] [0.12403482 0.30928354] [0.12190772 0.40995677]

How many clusters do you see ?

Why does it jump out?

What makes it a "cluster" ?

scatter plot

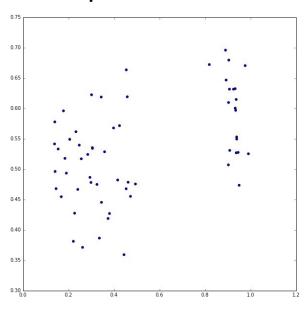


. . .

Brainstorm: what's a cluster?



scatter plot 2



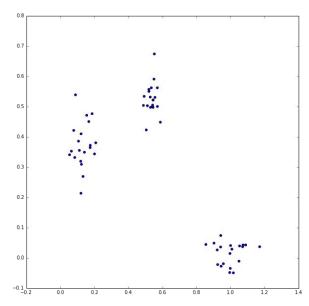
How many clusters do you see ?

Why does it jump out?

What makes it a "cluster"?

What makes it NOT a "cluster"?

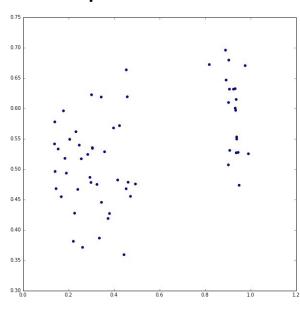
scatter plot



Clusters: a cognitive definition



scatter plot 2

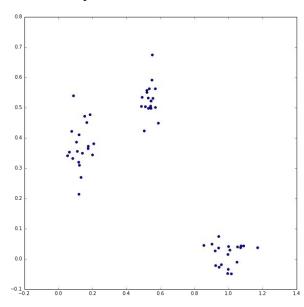


A partition of the dataset (not necessarily crisp)

A strong <u>internal</u> similarity (small intra/within cluster distance)

A strong <u>external</u> dissimilarity (large extra cluster distance)

scatter plot

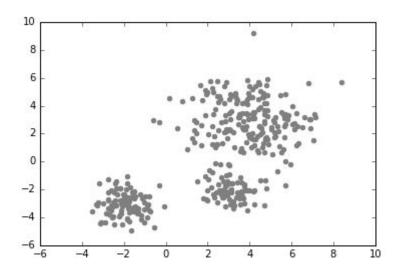




the k-Means algorithm

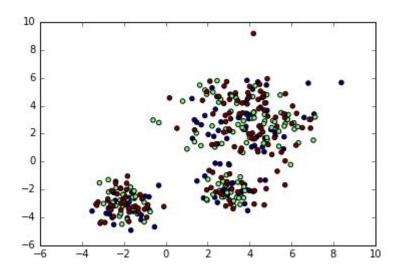


- 1. Randomly assign a number, from 1 to K, to each of the observations.
- Iterate until the cluster assignments stop changing:
 - a. For each of the K clusters, compute the cluster *centroid*: the vector of the *p* features means for the observations in the k-th cluster
 - Assign each observation to the cluster whose centroid is closest (defined using Euclidian distance)



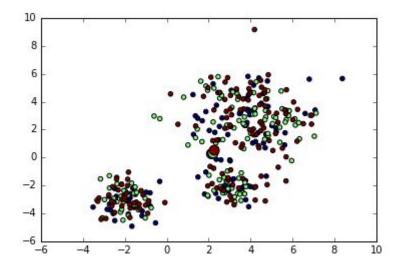


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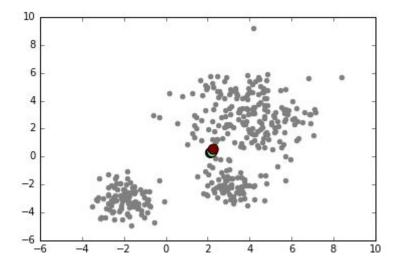


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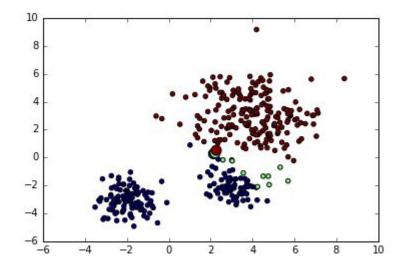


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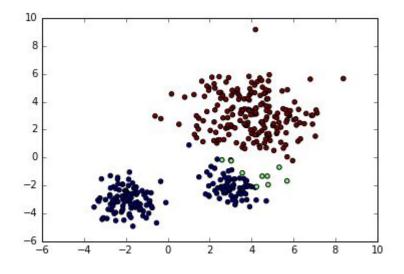


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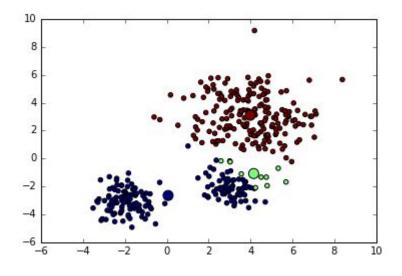


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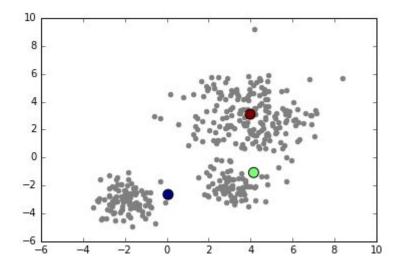


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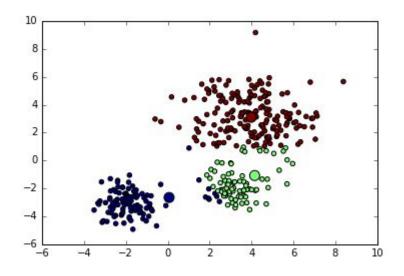


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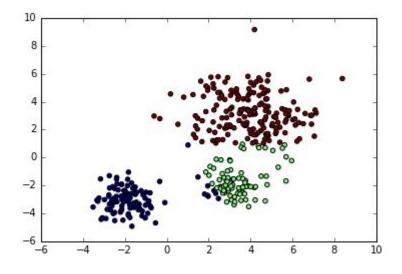


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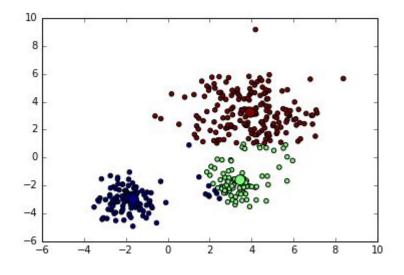


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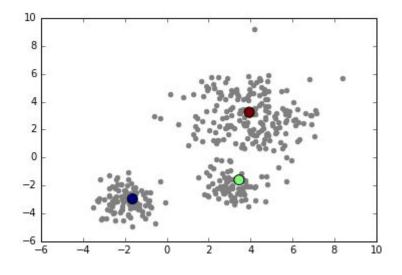


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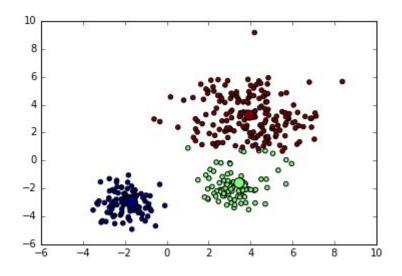


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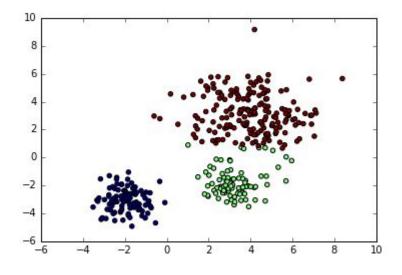


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Convergence of the algorithm



- 1. Randomly assign a number, from 1 to K, to each of the observations.
- 2. **Iterate** until the cluster assignments stop changing:
 - a. For each of the K clusters, compute the cluster *centroid*: the vector of the *p* features means for the observations in the k-th cluster
 - Assign each observation to the cluster whose centroid is closest (defined using Euclidian distance)

Objective: minimize "within cluster similarity"

$$\underset{C_1,...,C_K}{\text{minimize}} \left\{ \sum_{k=1}^K \frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 \right\}$$

Because of the effect of scale on euclidian distance

Pre-scale is almost mandatory

K-Means concerns



- 1. Randomly assign a number, from 1 to K, to each of the observations.
- 2. **Iterate** until the cluster assignments stop changing:
 - a. For each of the K clusters, compute the cluster *centroid*: the vector of the *p* features means for the observations in the k-th cluster
 - Assign each observation to the cluster whose centroid is closest (defined using Euclidian distance)



PB2: Robustness to initialization

PB3: Initialization strategy?



PB4: What's the best k?



PB1: When to stop?

PB1: When to stop?



Convergence is assured, but is not necessarily fast...

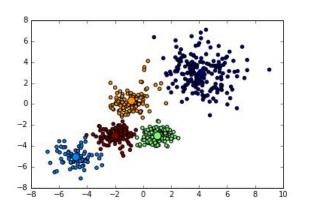
Solution 1: when the centroids don't change at all (you may wait a long time)

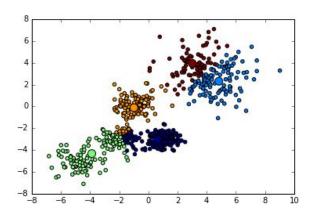
Solution 2: when the centroids don't change that much (tol)

Solution 3: when we get tired of waiting (max iter)

PB2: Robustness to initialization



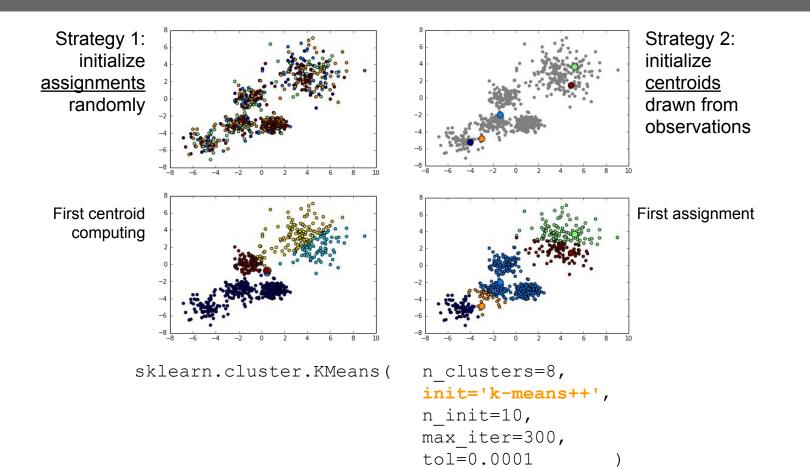




Depending on your initialization, you may (will) have different results. Solution: try several times and see if the result is stable

PB3: Initialization strategy



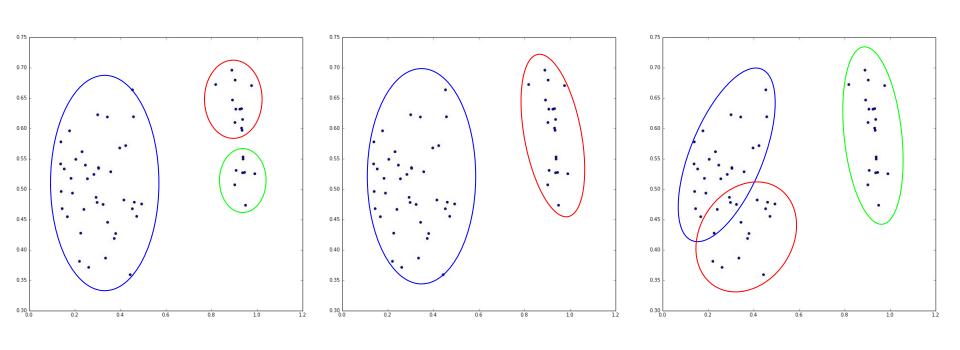




PB4: how to choose k? (Evaluating clustering)

What makes a good clustering?





Elbow method

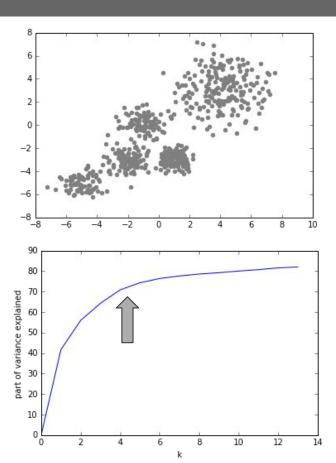


Compute total WCSS (k=1)

Compute WCSS for each iteration of k

Equiv. to a "total variance explained" plot

Observe where does increasing k stop that increase in WCSS?



Silhouette plot



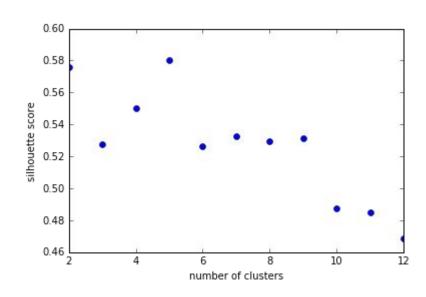
from sklearn.metrics import silhouette_score, silhouette_samples

For each point x_i :

- a(i) average dissimilarity of x_i with points in the same cluster
- b(i) average dissimilarity of x_i with points in the nearest cluster
 - "nearest" means cluster with the smallest b(i)

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

What's the range of silhouette scores?



GAP Statistic



For each K, compare W_K (within-cluster sum of squares) with that of randomly generated "reference distributions"

Generate B distributions

$$Gap(K) = \frac{1}{B} \sum_{b=1}^{B} \log W_{Kb} - \log W_{K}$$

Choose smallest K such that $Gap(K) \ge Gap(K+1) - s_{N+1}$ where s_K is the standard error of Gap(K)

paper by Hastie et al



Individual Assignment

(there's a hidden gem)