Clustering. Unsupervised Learning

Maria-Florina Balcan 11/05/2018

Clustering, Informal Goals

Goal: Automatically partition unlabeled data into groups of similar datapoints.

Question: When and why would we want to do this?

Useful for:

- Automatically organizing data.
- · Understanding hidden structure in data.
- Preprocessing for further analysis.
 - Representing high-dimensional data in a low-dimensional space (e.g., for visualization purposes).

Applications (Clustering comes up everywhere...)

Cluster news articles or web pages or search results by topic.







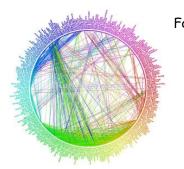


Cluster protein sequences by function or genes according to expression

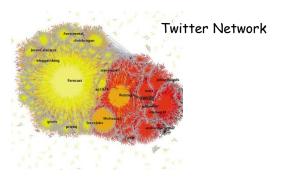
profile.



Cluster users of social networks by interest (community detection).



Facebook network



Applications (Clustering comes up everywhere...)

Cluster customers according to purchase history.





Cluster galaxies or nearby stars (e.g. Sloan Digital Sky Survey)



And many many more applications....

Clustering

Today:

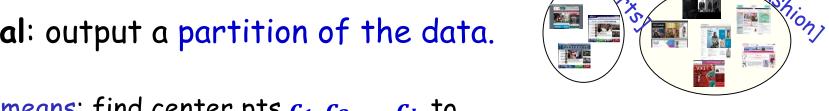
- Objective based clustering
 - K-means clustering
- Hierarchical clustering

Objective Based Clustering

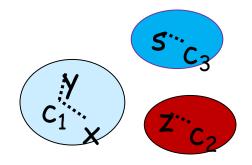
Input: A set 5 of n points, also a distance/dissimilarity measure specifying the distance d(x,y) between pairs (x,y).

E.g., # keywords in common, edit distance, wavelets coef., etc.

Goal: output a partition of the data.



- k-means: find center pts $c_1, c_2, ..., c_k$ to minimize $\sum_{i=1}^{n} \min_{i \in \{1,\dots,k\}} d^2(\mathbf{x}^i, \mathbf{c}_i)$
- k-median: find center pts $c_1, c_2, ..., c_k$ to minimize $\sum_{i=1}^{n} \min_{i \in \{1,...,k\}} d(x^i, c_i)$



K-center: find partition to minimize the maxim radius

Euclidean k-means Clustering

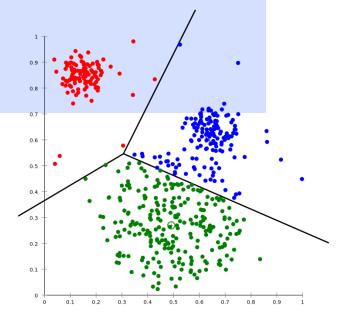
Input: A set of n datapoints $x^1, x^2, ..., x^n$ in \mathbb{R}^d

target #clusters k

Output: k representatives $c_1, c_2, ..., c_k \in \mathbb{R}^d$

Objective: choose $c_1, c_2, ..., c_k \in \mathbb{R}^d$ to minimize

$$\sum_{i=1}^{n} \min_{j \in \{1,\dots,k\}} \left| \left| \mathbf{x}^{i} - \mathbf{c}_{j} \right| \right|^{2}$$



Euclidean k-means Clustering

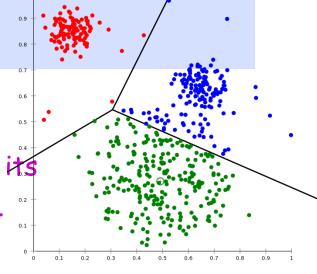
Input: A set of n datapoints $x^1, x^2, ..., x^n$ in R^d target #clusters k

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 $\sum_{i=1}^{n} \min_{j \in \{1,\dots,k\}} \left| \left| \mathbf{x}^i - \mathbf{c}_j \right| \right|^2$

Natural assignment: each point assigned to its closest center, leads to a Voronoi partition.



Euclidean k-means Clustering

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Computational complexity:

NP hard: even for k = 2 [Dagupta'08] or

d = 2 [Mahajan-Nimbhorkar-Varadarajan09]

There are a couple of easy cases...



An Easy Case for k-means: k=1

Input: A set of n datapoints $x^1, x^2, ..., x^n$ in R^d

Output: $c \in \mathbb{R}^d$ to minimize $\sum_{i=1}^n ||\mathbf{x}^i - \mathbf{c}||^2$

Solution: The optimal choice is $\mu = \frac{1}{n} \sum_{i=1}^{n} x^{i}$

Idea: bias/variance like decomposition

$$\frac{1}{n}\sum_{i=1}^{n}\left|\left|\mathbf{x}^{i}-\mathbf{c}\right|\right|^{2}=\left|\left|\boldsymbol{\mu}-\mathbf{c}\right|\right|^{2}+\frac{1}{n}\sum_{i=1}^{n}\left|\left|\mathbf{x}^{i}-\boldsymbol{\mu}\right|\right|^{2}$$

Avg k-means cost wrt c

Avg k-means cost wrt μ

So, the optimal choice for c is μ .

Another Easy Case for k-means: d=1

Input: A set of n datapoints $x^1, x^2, ..., x^n$ in R^d

Output: $c \in \mathbb{R}^d$ to minimize $\sum_{i=1}^n ||\mathbf{x}^i - \mathbf{c}||^2$

Extra-credit homework question

Hint: dynamic programming in time $O(n^2k)$.

Common Heuristic in Practice: The Lloyd's method

[Least squares quantization in PCM, Lloyd, IEEE Transactions on Information Theory, 1982]

Input: A set of n datapoints $x^1, x^2, ..., x^n$ in R^d

Initialize centers $c_1, c_2, ..., c_k \in \mathbb{R}^d$ and clusters $C_1, C_2, ..., C_k$ in any way.

Repeat until there is no further change in the cost.

- For each $j: C_j \leftarrow \{x \in S \text{ whose closest center is } c_j\}$
- For each $j: c_j \leftarrow mean of C_j$

Common Heuristic in Practice: The Lloyd's method

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Holding $c_1, c_2, ..., c_k$ fixed, pick optimal $C_1, C_2, ..., C_k$

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Common Heuristic: The Lloyd's method

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- For each $j: c_i \leftarrow mean of C_i$

Note: it always converges.

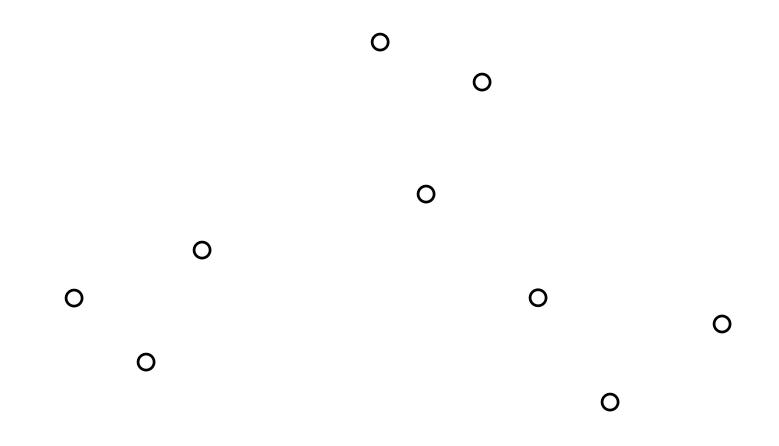
- the cost always drops and
- there is only a finite #s of Voronoi partitions
 (so a finite # of values the cost could take)

Initialization for the Lloyd's method

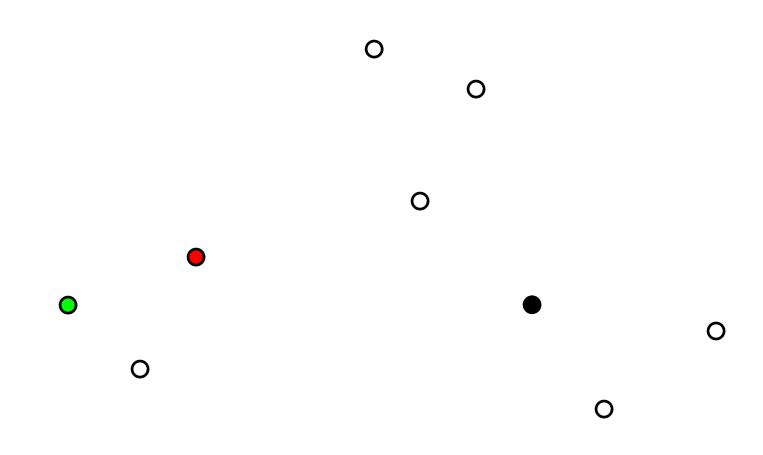
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\label{eq:continuous_set_of_n_def} \begin{subarray}{ll} \textbf{Input: A set of $n$ datapoints $x^1, x^2, ..., x^n$ in $R^d$ \\ \hline & \textbf{Initialize centers $c_1, c_2, ..., c_k \in R^d$ and \\ & clusters $C_1, C_2, ..., C_k$ in any way. \\ \hline & \textbf{Repeat until there is no further change in the cost.} \\ \end{subarray}
```

- For each $j: C_j \leftarrow \{x \in S \text{ whose closest center is } c_j\}$
- For each $j: c_j \leftarrow \text{mean of } C_j$
- Initialization is crucial (how fast it converges, quality of solution output)
- Discuss techniques commonly used in practice
 - Random centers from the datapoints (repeat a few times)
 - Furthest traversal
 - K-means ++ (works well and has provable guarantees)

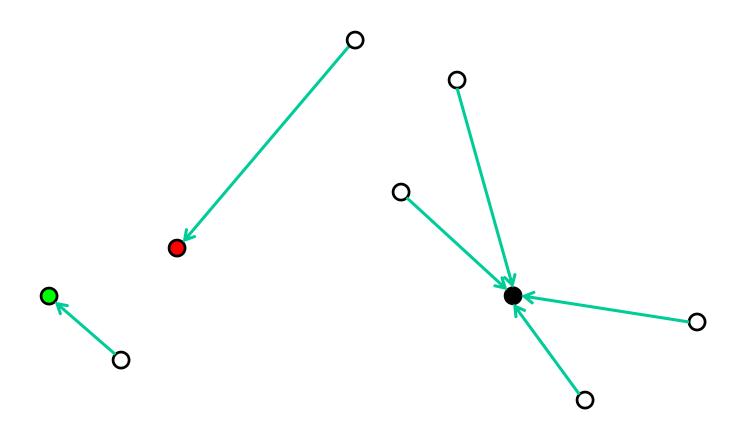
Example: Given a set of datapoints



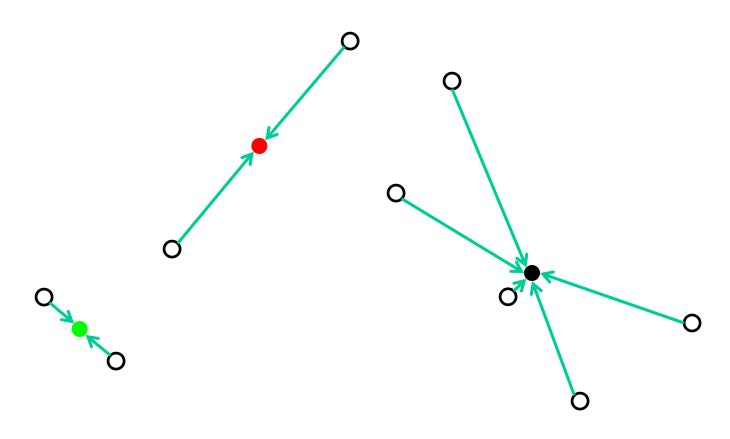
Select initial centers at random



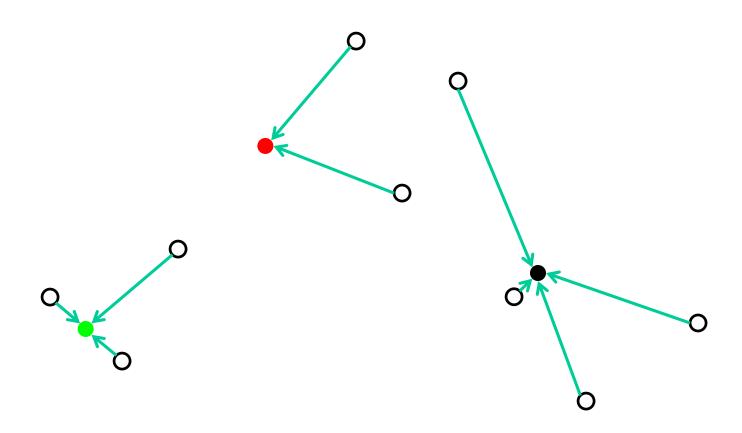
Assign each point to its nearest center



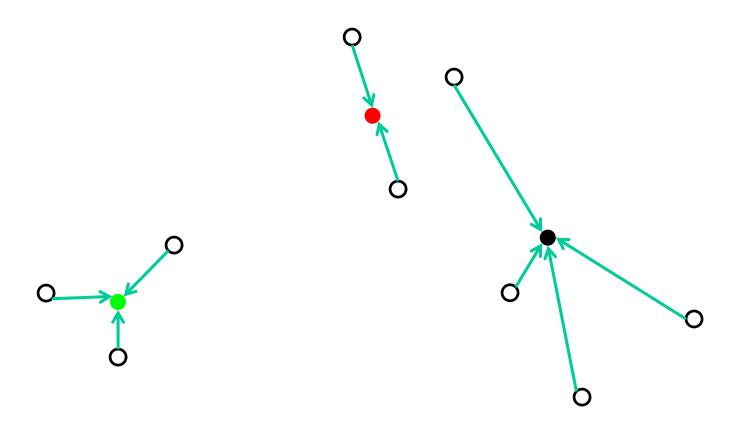
Recompute optimal centers given a fixed clustering



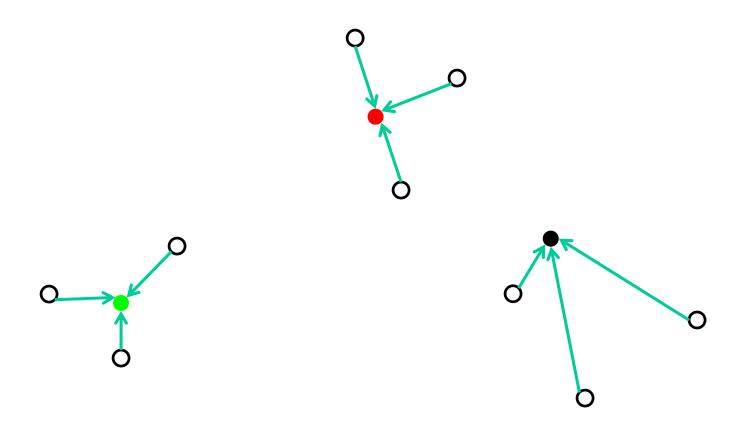
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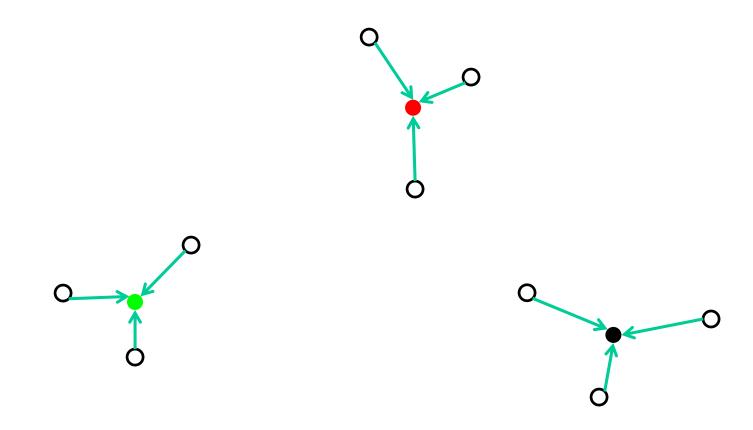
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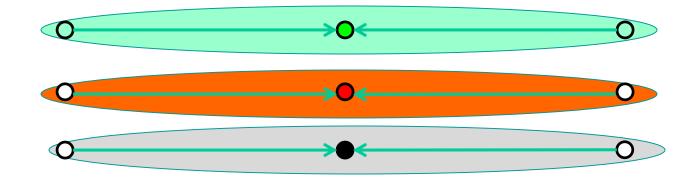
Assign each point to its nearest center



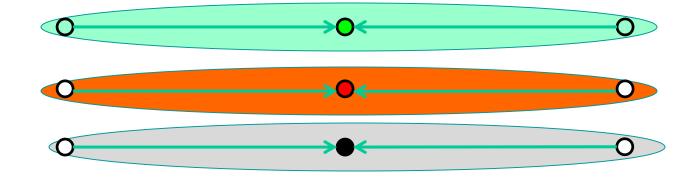
Recompute optimal centers given a fixed clustering



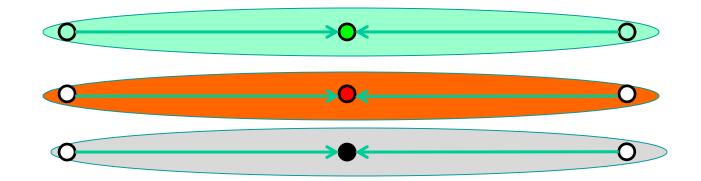
Get a good quality solution in this example.



It always converges, but it may converge at a local optimum that is different from the global optimum, and in fact could be arbitrarily worse in terms of its score.

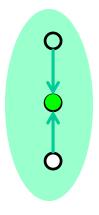


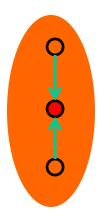
Local optimum: every point is assigned to its nearest center and every center is the mean value of its points.

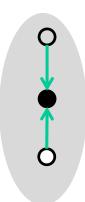


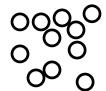
.It is arbitrarily worse than optimum solution....

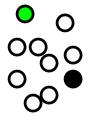


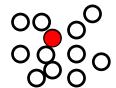




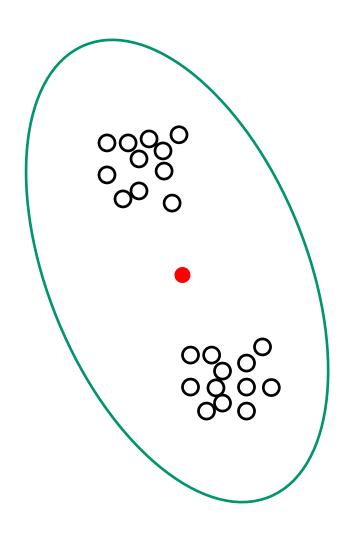


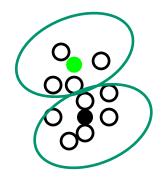






This bad performance, can happen even with well separated Gaussian clusters.





This bad performance, can happen even with well separated Gaussian clusters.

Some Gaussian are combined....



- If we do random initialization, as k increases, it becomes more likely we won't have perfectly picked one center per Gaussian in our initialization (so Lloyd's method will output a bad solution).
 - For k equal-sized Gaussians, Pr[each initial center is in a different Gaussian] $\approx \frac{k!}{k^k} \approx \frac{1}{e^k}$
 - Becomes unlikely as k gets large.

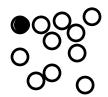
Another Initialization Idea: Furthest Point Heuristic

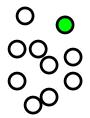
Choose c_1 arbitrarily (or at random).

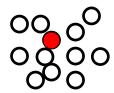
- For j = 2, ..., k
 - Pick c_j among datapoints $x^1, x^2, ..., x^d$ that is farthest from previously chosen $c_1, c_2, ..., c_{j-1}$

Fixes the Gaussian problem. But it can be thrown off by outliers....

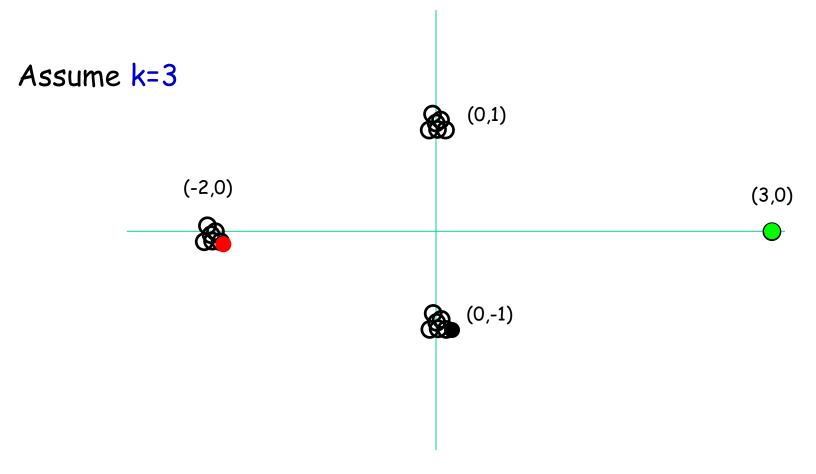
Furthest point heuristic does well on previous example



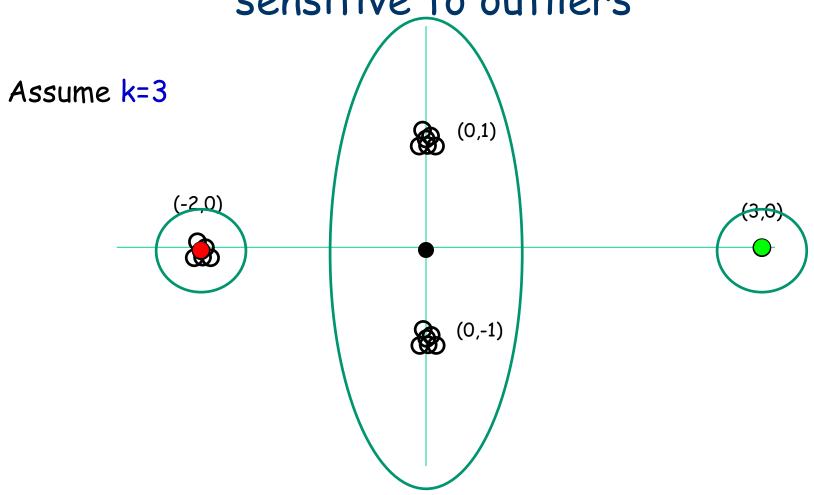




Furthest point initialization heuristic sensitive to outliers



Furthest point initialization heuristic sensitive to outliers



K-means++ Initialization: D² sampling [AV07]

- Interpolate between random and furthest point initialization
- Let D(x) be the distance between a point x and its nearest center. Chose the next center proportional to $D^2(x)$.
 - Choose c_1 at random.
 - For j = 2, ..., k
 - Pick c_j among $x^1, x^2, ..., x^d$ according to the distribution

$$Pr(c_j = x^i) \propto \min_{j' < j} \left| \left| x^i - c_{j'} \right| \right|^2 D^2(x^i)$$

Theorem: K-means++ always attains an O(log k) approximation to optimal k-means solution in expectation.

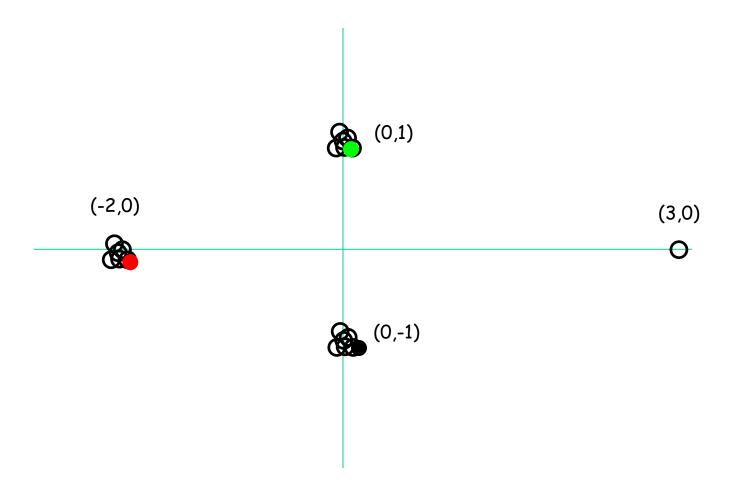
Running Lloyd's can only further improve the cost.

K-means++ Idea: D² sampling

- Interpolate between random and furthest point initialization
- Let D(x) be the distance between a point x and its nearest center. Chose the next center proportional to $D^{\alpha}(x)$.
 - $\alpha = 0$, random sampling
 - $\alpha = \infty$, furthest point (Side note: it actually works well for k-center)
 - $\alpha = 2$, k-means++

Side note: $\alpha = 1$, works well for k-median

K-means ++ Fix



K-means++/ Lloyd's Running Time

- K-means ++ initialization: O(nd) and one pass over data to select next center. So O(nkd) time in total.
- · Lloyd's method

Repeat until there is no change in the cost.

- For each j: $C_i \leftarrow \{x \in S \text{ whose closest center is } c_i\}$
 - For each $j: c_i \leftarrow mean of C_i$

Each round takes time O(nkd).

- Exponential # of rounds in the worst case [AV07].
- Expected polynomial time in the smoothed analysis (non worst-case) model!

K-means++/ Lloyd's Summary

- K-means++ always attains an O(log k) approximation to optimal k-means solution in expectation.
- Running Lloyd's can only further improve the cost.
- Exponential # of rounds in the worst case [AV07].
- Expected polynomial time in the smoothed analysis model!
- Does well in practice.

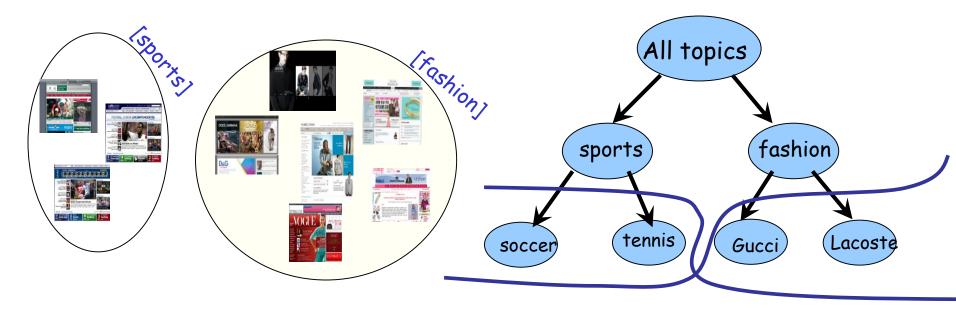
What value of k???

 Heuristic: Find large gap between k -1-means cost and k-means cost.

 Hold-out validation/cross-validation on auxiliary task (e.g., supervised learning task).

Try hierarchical clustering.

Hierarchical Clustering

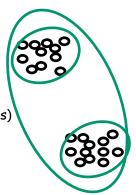


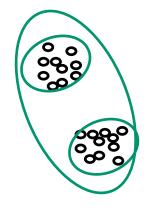
- · A hierarchy might be more natural.
- Different users might care about different levels of granularity or even prunings.

Hierarchical Clustering

Top-down (divisive)

- Partition data into 2-groups (e.g., 2-means)
- Recursively cluster each group.



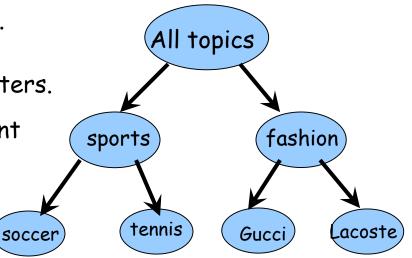


Bottom-Up (agglomerative)

Start with every point in its own cluster.

Repeatedly merge the "closest" two clusters.

 Different defs of "closest" give different algorithms.

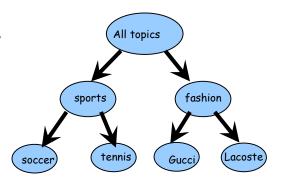


Bottom-Up (agglomerative)

Have a distance measure on pairs of objects.

d(x,y) - distance between x and y

E.g., # keywords in common, edit distance, etc



- Single linkage: $dist(A, B) = \min_{x \in A, x' \in B'} dist(x, x')$
- Complete linkage: $dist(A, B) = \max_{x \in A, x' \in B'} dist(x, x')$
- Average linkage: $dist(A, B) = \underset{x \in A, x' \in B'}{avg} dist(x, x')$

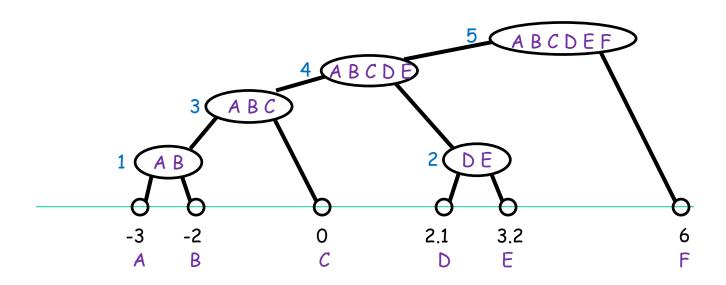
Single Linkage

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Single linkage:
$$dist(A, B) = \min_{x \in A, x' \in B} dist(x, x')$$

Dendogram



Single Linkage

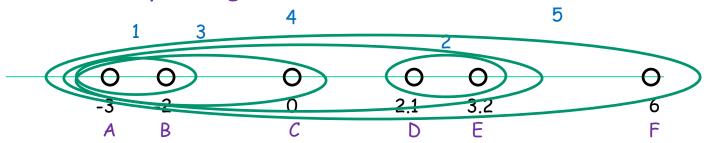
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Single linkage:
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One way to think of it: at any moment, we see connected components of the graph where connect any two pts of distance < r.

Watch as r grows (only n-1 relevant values because we only we merge at value of r corresponding to values of r in different clusters).



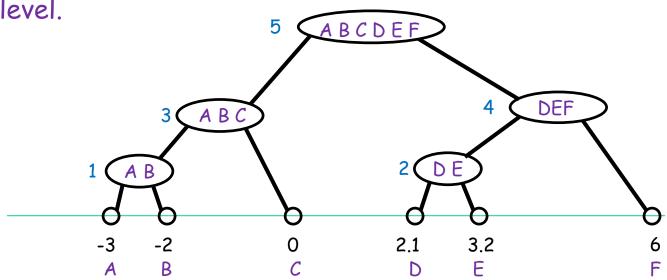
Complete Linkage

Bottom-up (agglomerative)

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Complete linkage:
$$dist(A, B) = \max_{x \in A, x' \in B} dist(x, x')$$

One way to think of it: keep max diameter as small as possible at any level.



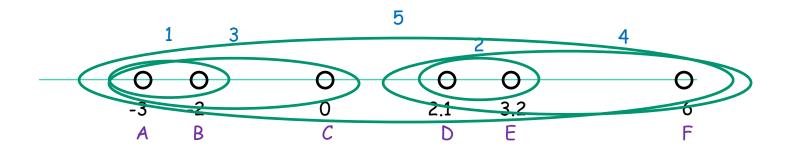
Complete Linkage

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Complete linkage:
$$dist(A, B) = \max_{x \in A, x' \in B} dist(x, x')$$

One way to think of it: keep max diameter as small as possible.



Running time

- Each algorithm starts with N clusters, and performs N-1 merges.
- For each algorithm, computing dist(C, C') can be done in time $O(|C| \cdot |C'|)$. (e.g., examining dist(x, x') for all $x \in C, x' \in C'$)
- Time to compute all pairwise distances and take smallest is $O(N^2)$.
- Overall time is $O(N^3)$.

In fact, can run all these algorithms in time $O(N^2 \log N)$.

If curious, see: Christopher D. Manning, Prabhakar Raghavan and Hinrich Schütze, Introduction to Information Retrieval, Cambridge University Press. 2008. http://www-nlp.stanford.edu/IR-book/

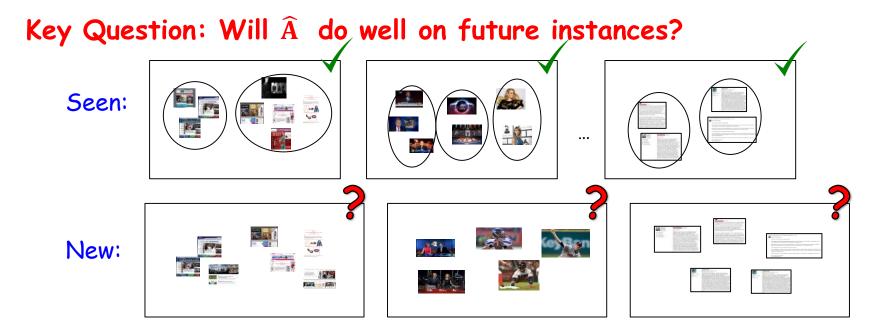
Goal: given family of algos \mathbf{F} , sample of typical instances from domain (unknown distr. \mathbf{D}), find algo that performs well on new instances from \mathbf{D} .

[Balcan-Nagarajan-Vitercik-White, COLT'17] [Balcan-Dick-White, NIPS'18]

Dynamic Programming Large family F of algorithms Farthest Location Greedy Sample of typical inputs Input m: Input 2: Input 1: Clustering: Input 2: Input m: Input 1:

Goal: given family of algos F, sample of typical instances from domain (unknown distr. D), find algo that performs well on new instances from D.

Approach: find \widehat{A} near optimal algorithm over the set of samples.



Sample Complexity: How large should our sample of typical instances be in order to guarantee good performance on new instances?

Goal: given family of algos F, sample of typical instances from domain (unknown distr. D), find algo that performs well on new instances from D.

Approach: find \widehat{A} near optimal algorithm over the set of samples.

Key Question: Will \widehat{A} do well on future instances?

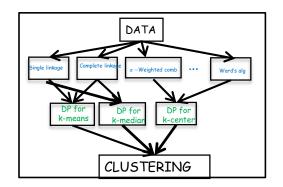
Key tool: uniform convergence, for any algo in F, average performance over samples "close" to its expected performance.

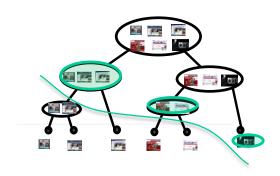
• Imply that $\widehat{\mathbf{A}}$ has high expected performance.

Learning theory: $N = O(\dim(\mathbf{F})/\epsilon^2)$ instances suffice for ϵ -close.

• Clustering: Linkage + Dynamic Programming

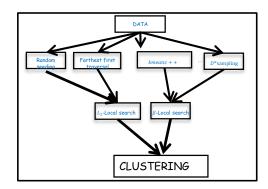
[Balcan-Nagarajan-Vitercik-White, COLT'17]





• Clustering: Greedy Seeding + Local Search
[Balcan-Dick-White, NIPS'18]

Parametrized Lloyds methods



What You Should Know

- Partitional Clustering. k-means and k-means ++
 - Lloyd's method
 - Initialization techniques (random, furthest traversal, k-means++)

- Hierarchical Clustering.
 - Single linkage, Complete linkage