New Developments In The Theory Of Clustering that's all very well in practice, but does it work in theory ?

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What we will cover

A few of the recent theory results on clustering:

- Practical algorithms that have strong theoretical guarantees
- Models to explain behavior observed in practice

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What we will not cover

The rest:

- Recent strands of theory of clustering such as metaclustering and privacy preserving clustering
- Clustering with distributional data assumptions
- Proofs

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I Euclidean Clustering and k-means algorithm

II Bregman Clustering and k-means

III Stability

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Outline

I Euclidean Clustering and k-means algorithm

- What to do to select initial centers (and what not to do)
- How long does k-means take to run in theory, practice and theoretical practice
- How to run k-means on large datasets
- II Bregman Clustering and k-means

III Stability

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- II Bregman Clustering and k-means
 - Bregman Clustering as generalization of k-means
 - Performance Results

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- **III** Stability
 - How to relate closeness in cost function to closeness in clusters.

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Euclidean Clustering and k-means

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What does it mean to cluster?

Given *n* points in \mathbb{R}^d find the best way to split them into *k* groups.

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How do we define "best" ? Example:



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How do we define "best" ?

Minimize the maximum radius of a cluster



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How do we define "best" ?

Maximize the average inter-cluster distance



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How do we define "best" ?

Minimize the variance within each cluster.



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How do we define "best" ?

Minimize the variance within each cluster.

Minimizing total variance

For each cluster $C_i \in \mathcal{C}$, $c_i = \frac{1}{|C_i|} \sum_{x \in C_i} x$ is the expected location of a point in a cluster. Then the variance of each cluster is:

$$\sum_{x \in C_i} = \|x - c_i\|^2$$

And the total objective is:

$$\phi = \sum_{c_i} \sum_{x \in C_i} \|x - c_i\|^2$$

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Given *X* and *k*, find a clustering $\mathscr{C} = \{C_1, C_2, \dots, C_k\}$ that minimizes: $\phi(X, \mathscr{C}) = \sum_{c_i} \sum_{x \in C_i} ||x - c_i||^2$

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Definition

Let ϕ^* denote the value of the optimum solution above. We say that a clustering C' is α -approximate if:

$$\phi^* \leq \phi(X, \mathscr{C}') \leq \alpha \cdot \phi^*$$

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Solving this problem

This problem is NP-complete, even when the pointset *X* lies in two dimensions...

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Solving this problem

This problem is NP-complete, even when the pointset *X* lies in two dimensions...

...but we've been solving it for over 50 years! [S56][L57][M67]

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Example

Given a set of data points



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Select initial centers at random



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Assign each point to nearest center



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Recompute optimum centers given a fixed clustering



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Example	
Repeat	



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Example	
Repeat	



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Example	
Repeat	



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Until the clustering doesn't change



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This algorithm terminates!

Recall the total error:

$$\phi(X, \mathscr{C}) = \sum_{c_i} \sum_{x \in C_i} ||x - c_i||^2$$

In every iteration ϕ is reduced:

- Assigning each point to the nearest center reduces ϕ
- Given a fixed cluster, the mean is the optimal location for the center (requires proof)

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The algorithm finds a local minimum ...



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... that's potentially arbitrarily worse than optimum solution







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Performance

But does this really happen?



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Performance

But does this really happen? YES!



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Finding a good set of initial points is a black art

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Finding a good set of initial points is a black art

- Try many times with different random seeds
 - Most common method
 - Has limited benefit even in case of Gaussians

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 - Hundreds of heuristics
 - Including pre & post processing ideas

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There exists a fast and simple initialization scheme with provable performance guarantees

Random Initializations on Gaussians

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Random Initializations on Gaussians

Some Gaussians are combined



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But the Gaussian case has an easy fix: use a furthest point heuristic

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But this fix is overly sensitive to outliers







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k-means++

What if we interpolate between the two methods?

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k-means++

What if we interpolate between the two methods?

Let D(x) be the distance between a point x and its nearest cluster center. Chose the next point proportionally to $D^{\alpha}(x)$.

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k-means++

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Let D(x) be the distance between a point x and its nearest cluster center. Chose the next point proportionally to $D^{\alpha}(x)$.

• $\alpha = 0 \longrightarrow$ Random initialization

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- $\alpha = 2 \longrightarrow k means + +$

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- $\alpha = 0 \longrightarrow$ Random initialization
- $\alpha = \infty \longrightarrow$ Furthest point heuristic
- $\alpha = 2 \longrightarrow k-means++$

More generally

Set the probability of selecting a point proportional to its contribution to the overall error.

- If minimizing $\sum_{c_i} \sum_{x \in C_i} ||x c_i||$, sample according to *D*.
- If minimizing $\sum_{c_i} \sum_{c \in C_i} ||x c_i||_{\infty}$, sample according to D^{∞} (take the furthest point).

If the data set looks Gaussian...







If the data set looks Gaussian...







If the data set looks Gaussian...







If the data set looks Gaussian...







If the data set looks Gaussian...







If the outlier should be its own cluster ...



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What can we say about performance of k-means++?

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What can we say about performance of k-means++?

Theorem (AV07)

This algorithm always attains an $O(\log k)$ approximation in expectation

What can we say about performance of k-means++?

Theorem (AV07)

This algorithm always attains an $O(\log k)$ approximation in expectation

Theorem (ORSS06)

A slightly modified version of this algorithm attains an O(1) approximation if the data is 'nicely clusterable' with k clusters.

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What do we mean by 'nicely clusterable'?

Intuitively, *X* is nicely clusterable if going from k - 1 to *k* clusters drops the total error by a constant factor.

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What do we mean by 'nicely clusterable'?

Intuitively, X is nicely clusterable if going from k - 1 to k clusters drops the total error by a constant factor.

Definition

A pointset X is (k, ϵ) -separated if $\phi_k^*(X) \le \epsilon^2 \phi_{k-1}^*(X)$.

Why does this work?

Intuition

Look at the optimum clustering. In expectation:

- If the algorithm selects a point from a new OPT cluster, that cluster is covered pretty well
- If the algorithm picks two points from the same OPT cluster, then other clusters must contribute little to the overall error
Why does this work?

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Look at the optimum clustering. In expectation:

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As long as the points are reasonably well separated, the first condition holds.

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- If the algorithm picks two points from the same OPT cluster, then other clusters must contribute little to the overall error

As long as the points are reasonably well separated, the first condition holds.

Two theorems

- Assume the points are (k, ϵ) -separated and get an O(1) approximation.
- Make no assumptions about separability and get an $O(\log k)$ approximation.

k-means++ Summary:

- To select the next cluster, sample a point in proportion to its current contribution to the error
- Works for *k*-means, *k*-median, other objective functions
- Universal *O*(log *k*) approximation, *O*(1) approximation under some assumptions
- Can be implemented to run in O(nkd) time (same as a single k-means step)

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But does it actually work?

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Large Evaluation

Data set	ϕ : k-means++ vs k-means			
	(k=10,25,50,100)			
1	1.011	1.036	1.115	1.544
2	1.052	1.344	1.523	1.421
3	1.122	2.083	2.987	3.481
4	1.007	1.206	1.207	1.421
5	1.038	1.303	1.367	1.861
6	1.010	1.013	1.018	1.043
7	1.027	1.160	1.560	2.056
8	0.997	1.009	1.044	1.186
9	1.090	1.181	1.314	1.297
10	13.87	132.7	684.9	3728
11	1.002	1.003	1.011	1.046
12	2.420	5.027	11.08	22.76
13	2.202	9.532	11.96	30.17
14	1.020	1.039	7.942	1.008
15	1.013	554.2	1.003	1.004
Range	[0.997, 3728]			
Median	1.207			
Mean	87.93			

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KM++ v. KM v. KM-Hybrid

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How fast does k-means converge?

It appears the algorithm converges in under 100 iterations (even faster with smart initialization).

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How fast does k-means converge?

It appears the algorithm converges in under 100 iterations (even faster with smart initialization).

Theorem (V09)

There exists a pointset X in \mathbb{R}^2 and a set of initial centers \mathscr{C} so that k-means takes $2^{\Omega(k)}$ iterations to converge when initialized with \mathscr{C} .

Finding the disconnect

In theory:

• k-means might run in exponential time

In practice:

• k-means converges after a handful of iterations

It works in practice but it does not work in theory!

Robustness of worst case examples

Perhaps the worst case examples are too precise, and can never arise out of natural data

Quantifying the robustness

If we slightly perturb the points of the example:

- The optimum solution shouldn't change too much
- Will the running time stay exponential?

Small Perturbations



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Perturbation

To each point $x \in X$ add independent noise drawn from $N(0, \sigma^2)$.

Definition

The smoothed complexity of an algorithm is the maximum expected running time after adding the noise:

 $\max_{X} \mathbb{E}_{\sigma}[Time(X + \sigma)]$

Theorem (AMR09)

The smoothed complexity of k-means is bounded by

$$O\left(\frac{n^{34}k^{34}d^8D^6\log^4 n}{\sigma^6}\right)$$

Notes

- While the bound is large, it is not exponential (2^k ≫ k³⁴ for large enough k)
- The $(D/\sigma)^6$ factor shows the bound is scale invariant

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Comparing bounds

The smoothed complexity of k-means is polynomial in n, k and D/σ where D is the diameter of X, whereas the worst case complexity of k-means is exponential in k

Implications

The pathological examples:

- Are very brittle
- Can be avoided with a little bit of random noise

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Running Time

- Exponential worst case running time
- Polynomial typical case running time

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Running Time

- Exponential worst case running time
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Solution Quality

- Arbitrary local optimum, even with many random restarts
- Simple initialization leads to a good solution

Implementing k-means++

Initialization:

- Takes *O*(*nd*) time and one pass over the data to select the next center
- Takes O(nkd) time total
- Overall running time:
 - Each round of k-means takes O(nkd) running time
 - Typically finish after a constant number of rounds

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Overall running time:

- Each round of k-means takes O(nkd) running time
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Large Data

What if O(nkd) is too much, can we parallelize this algorithm?

Approach

Partition the data:

• Split X into X_1, X_2, \ldots, X_m of roughly equal size.

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Approach

Partition the data:

• Split X into X_1, X_2, \ldots, X_m of roughly equal size.

In parallel compute a clustering on each partition:

• Find $\mathscr{C}^{j} = \{C_{1}^{j}, \dots, C_{k}^{j}\}$: a good clustering on each partition, and denote by w_{i}^{j} the number of points in cluster C_{i}^{j} .

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Partition the data:

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Find \$\mathcal{C}^j = {C_1^j, ..., C_k^j}\$: a good clustering on each partition, and denote by \$w_i^j\$ the number of points in cluster \$C_i^j\$.

Cluster the clusters:

Let Y = ∪_{1≤j≤m} 𝔅^j. Find a clustering of Y, weighted by the weights W = {w^j_i}.







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Partition the dataset



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Cluster each partition separately









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Cluster each partition separately









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Cluster the clusters



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Cluster the clusters



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Final clustering:



Final clustering:





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Quality of the solution

What happens when we approximate the approximation?

- Suppose the algorithm in phase 1 gave a β-approximate solution to its input
- Algorithm in phase 2 gave a γ-approximate solution to its (smaller) input

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Quality of the solution

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Theorem (GNMO00, AJM09)

The two phase algorithm gives a $4\gamma(1 + \beta) + 2\beta$ approximate solution.

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Running time

Suppose we partition the input across m different machines.

- First phase running time: $O(\frac{nkd}{m})$.
- Second phase running time $O(mk^2d)$.

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Approximation Guarantees

Using k-means++ sets $\beta = \gamma = O(\log k)$ and leads to a $O(\log^2 k)$ approximation.

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Approximation Guarantees

Using k-means++ sets $\beta = \gamma = O(\log k)$ and leads to a $O(\log^2 k)$ approximation.

Improving the Approximation

Must improve the approximation guarantee of the first round, but can use a larger k to ensure every cluster is well summarized.

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Theorem (ADK09)

Running k-means++ initialization for O(k) rounds leads to a O(1) approximation to the optimal solution (but uses more centers than OPT).

Two round k-means++

Final Algorithm

Partition the data:

• Split X into X_1, X_2, \ldots, X_m of roughly equal size.

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Compute a clustering using $\ell = O(k)$ centers each partition:

 Find 𝔅^j = {C^j₁,...,C^j_ℓ} using k-means++ on each partition, and denote by w^j_i the number of points in cluster C^j_i.

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- Find 𝔅^j = {C^j₁,...,C^j_ℓ} using k-means++ on each partition, and denote by w^j_i the number of points in cluster C^j_i.
 Cluster the clusters.
 - Let $Y = \bigcup_{1 \le j \le m} \mathscr{C}^j$ be a set of $O(\ell m)$ points. Use k-means++ to cluster Y, weighted by the weights $W = \{w_i^j\}$.

Theorem

The algorithm achieves an O(1) approximation in time $O(\frac{nkd}{m} + mk^2d)$

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Before...

k-means used to be a prime example of the disconnect between theory and practice – it works well, but has horrible worst case analysis

...and after

Smoothed analysis explains the running time and rigorously analyzed initializations routines help improve clustering quality.

Outline

Outline

I Euclidean Clustering and k-means algorithm

- What to do to select initial centers (and what not to do)
- How long does k-means take to run in theory, practice and theoretical practice
- How to run k-means on large datasets

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- II Bregman Clustering and k-means
 - Bregman Clustering as generalization of k-means
 - Performance Results

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 - Bregman Clustering as generalization of k-means
 - Performance Results
- **III** Stability
 - How to relate closeness in cost function to closeness in clusters.

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Clustering With Non-Euclidean Metrics

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Theory of Clustering

Application I: Clustering Documents



Kullback-Leibler distance:

$$D(p,q) = \sum_{i} p_i \log \frac{p_i}{q_i}$$

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Application II: Image Analysis



Kullback-Leibler distance:

$$D(p,q) = \sum_{i} p_i \log \frac{p_i}{q_i}$$

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Itakuro-Saito distance:

$$D(p,q) = \sum_{i} \frac{p_i}{q_i} - \log \frac{p_i}{q_i} - 1$$

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Definition

Let $\phi : \mathbb{R}^d \to \mathbb{R}$ be a strictly convex function. The Bregman divergence d_{ϕ} is defined as

$$D_{\phi}(x \parallel y) = \phi(x) - \phi(y) - \langle \nabla \phi(y), x - y \rangle$$

Examples:

Kullback-Leibler: $\phi(x) = \sum x_i \ln x_i - x_i, D_{\phi}(x \parallel y) = \sum x_i \ln \frac{x_i}{y_i}$ Itakura-Saito: $\phi(x) = -\sum \ln x_i, D_{\phi}(x \parallel y) = \sum_i \frac{x_i}{y_i} - \log \frac{x_i}{y_i} - 1$ ℓ_2^2 : $\phi(x) = \frac{1}{2} ||x||^2, D_{\phi}(x \parallel y) = ||x - y||^2$

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k-means clustering \equiv Bregman clustering

- The algorithm works the same way.
- Same (bad) worst-case behavior
- Same (good) smoothed behavior
- Same (good) quality guarantees, with correct initialization

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$$D_{\phi}(x \parallel y) = \phi(x) - \phi(y) - \langle \nabla \phi(y), x - y \rangle$$

- Asymmetry: In general, $D_{\phi}(p \parallel q) \neq D_{\phi}(q \parallel p)$
- No triangle inequality: $D_{\phi}(p \parallel q) + D_{\phi}(q \parallel r)$ can be less than $D_{\phi}(p \parallel r)$!

How can we now do clustering $?_{\text{B}} \rightarrow \text{A} = \text{A} \rightarrow \text{A}$

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Initialize cluster centers while not converged **do** Assign points to nearest cluster center

Find new cluster center by averaging points assigned together end while

Key Point Setting cluster center as centroid minimizes the average **squared distance** to center

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Initialize cluster centers **while** not converged **do** Assign points to nearest cluster center

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Key Point

Setting cluster center as centroid minimizes the average **squared distance** to center

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Problem

Given points $x_1, \ldots x_n \in \mathbb{R}^d$, find c such that

 $\sum_i D_{\phi}(x_i \parallel c)$

is minimized.

Answer

$$c = \frac{1}{n} \sum x_i$$

Independent of ϕ [BMDG05] !

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Initialize cluster centers while not converged do Assign points to nearest cluster center (by measuring $D_{\phi}(x \parallel c)$) Find new cluster center by averaging points assigned together end while

Key Point

Setting cluster center as centroid minimizes average **Bregman divergence** to center

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Lemma ([BMDG05])

The (Bregman) k-means algorithm converges in cost.

Euclidean distance:

The quantity

$$\sum_{C} \sum_{x \in C} \|x - \operatorname{center}(C)\|^2$$

decreases with each iteration of k-means

Bregman divergence: Bregman Information:

$$\sum_{C} \sum_{x \in C} D_{\phi}(x \parallel \text{center}(C))$$

decreases with each iteration of the Bregman k-means algorithm.

Expectation maximization:

- Initialize density parameters and means for k distributions while not converged **do**
- For distribution *i* and point *x*, compute *conditional probability* p(i|x) that *x* was drawn from *i* (by Bayes rule) For each distribution *i*, recompute new density parameters and means (via maximum likelihood) end while
- This yields a *soft clustering* of points to "clusters"
- Originally used for mixtures of Gaussians.

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Exponential Families And Bregman Divergences

Definition (Exponential Family)

Parametric family of distributions $p_{\Psi,\theta}$ is an exponential family if each density is of the form

$$p_{\Psi,\theta} = \exp(\langle x, \theta \rangle - \Psi(\theta))p_0(x)$$

with Ψ convex.

Let $\phi(t) = \Psi^*(t)$ be the Legendre-Fenchel dual of $\Psi(x)$:

$$\phi(t) = \sup_{x} (\langle x, t \rangle - \Psi(x))$$

Theorem ([BMDG05])

$$p_{\Psi,\theta} = \exp(-D_{\phi}(x \parallel \mu))b_{\phi}(x)$$

where μ is the expectation parameter $\nabla \Psi(\theta)$

Expectation maximization:

Initialize density parameters and means for k distributions while not converged **do**

For distribution *i* and point *x*, compute *conditional probability* p(i|x) that *x* was drawn from *i* (by Bayes rule) For each distribution *i*, recompute new density parameters and means (via maximum likelihood) end while

Choosing the corresponding Bregman divergence $D_{\phi}(\cdot \| \cdot), \phi = \Psi^*$ gives mixture density estimation for any exponential family $p_{\Psi,\theta}$.

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Performance Analysis

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Two questions:

Problem (Rate of convergence)

Given an arbitrary set of n points in d dimensions, how long does it take for (Bregman) k-means to converge ?

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Problem (Rate of convergence)

Given an arbitrary set of n points in d dimensions, how long does it take for (Bregman) k-means to converge ?

Problem (Quality of Solution)

Let OPT denote the optimal clustering that minimizes the average sum of (Bregman) distances to cluster centers. How close to OPT is the solution returned by (Bregman) k-means ?

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Two questions:

Problem (Rate of convergence)

Given an arbitrary set of n points in d dimensions, how long does it take for (Bregman) k-means to converge ?

Parameters: *n*,*k*,*d*.

ⓒ Good news

k-means always converges in $O(n^{kd})$ time.

🙂 Bad news

k-means can take time $2^{\Omega(k)}$ to converge:

Parameters: *n*,*k*,*d*.

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k-means can take time $2^{\Omega(k)}$ to converge:

• Even if d = 2, i.e in the plane

Parameters: *n*,*k*,*d*.

③ Good news

k-means always converges in $O(n^{kd})$ time.

🙂 Bad news

k-means can take time $2^{\Omega(k)}$ to converge:

- Even if d = 2, i.e in the plane
- Even if centers are chosen from the initial data

Euclidean distance:

k-means can take time $2^{\Omega(k)}$ to converge:

- Even if *d* = 2, i.e in the plane
- Even if centers are chosen from the initial data

Bregman divergence:

For some Bregman divergences, *k*-means can take time $2^{\Omega(k)}$ to converge[MR09]:

- Even if *d* = 2, i.e in the plane
- Even if centers are chosen from the initial data

Proof Idea

"Well behaved" Bregman divergences look "locally Euclidean":



Take a bad Euclidean instance and shrink it to make it local.



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Smoothed Analysis

Real inputs aren't worst-case!



Analyze expected run-time over perturbations.

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Smoothed Analysis

Real inputs aren't worst-case!



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Theorem

Smoothed complexity of k-means using Gaussian noise with variance σ is **polynomial** in n and $1/\sigma$.

Compare this to worst-case lower bound of $2^{\Theta(n)}$

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Bregman Smoothing

Normal smoothing doesn't work !





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More general notion of smoothing:

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Bregman smoothing

More general notion of smoothing:



• perturbation should stay close to a hyperplane

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Bregman smoothing

More general notion of smoothing:



- perturbation should stay close to a hyperplane
- density of perturbation is proportional to $1/\sigma^d$

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Theorem ([MR09])

For "well-behaved" Bregman divergences, smoothed complexity is bounded by $poly(n^{\sqrt{k}}, 1/\sigma)$ and $k^{kd}poly(n, 1/\sigma)$.

This is in comparison to worst-case bound of $2^{\Omega(n)}$.

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Two questions:

Problem (Rate of convergence)

Given an arbitrary set of n points in d dimensions, how long does it take for (Bregman) k-means to converge ?

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Optimality and Approximations

Problem

Given x_1, \ldots, x_n , and parameter k, find k centers c_1, \ldots, c_k such that

$$\sum_{x=1}^{n} \min_{j=1}^{k} d(x_i, c_j)$$

is minimized.

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Problem

Given x_1, \ldots, x_n , and parameter k, find k centers c_1, \ldots, c_k such that

$$\sum_{x=1}^{n} \min_{j=1}^{k} d(x_i, c_j)$$

is minimized.

Problem (*c*-approximation)

Let OPT be the optimal solution above. Fix c > 0. Find centers $c'_1, \ldots c'_k$ such that if $A = \sum_{x=1}^n \min_{j=1}^k d(x_i, c'_j)$, then

 $OPT \le A \le c \cdot OPT$

Initialization

- Let distance from x to nearest cluster center be D(x)
- Pick *x* as new center with probability

 $p(x) \propto D^2(x)$

Properties of solution:

- For arbitrary data, this gives $O(\log n)$ -approximation
- For "well-separated data", this gives constant (*O*(1))-approximation.

Informally, data is (k, α) -well separated if the best clustering that uses k - 1 clusters has cost that is $\geq 1/\alpha \cdot \text{OPT}$.

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What is 'well-separated'

Informally, data is (k, α) -well separated if the best clustering that uses k - 1 clusters has cost that is $\geq 1/\alpha \cdot \text{OPT}$.



What is 'well-separated'

Informally, data is (k, α) -well separated if the best clustering that uses k - 1 clusters has cost that is $\geq 1/\alpha \cdot \text{OPT}$.



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Initialization

- Let Bregman divergence from *x* to nearest cluster center be D(x)
- Pick *x* as new center with probability

 $p(x) \propto D(x)$

Run algorithm as before.

Theorem ([AB09, AB10])

- O(1)-approximation for (k, α) -separated sets.
- O(log n) approximation in general.

Stability in clustering

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Target and Optimal clustering



Two measures of cost:

• Distance between clusterings $\mathscr{C}, \mathscr{C}^*$:

 $d(\mathcal{C}, \mathcal{C}^*)$ = fraction of points on which they disagree

• (Quality) distance from *C* to OPT:

$$d_q(\mathscr{C}, OPT) = \frac{\cot(\mathscr{C})}{\cot(OPT)}$$

Can closeness in d_q imply closeness in d?

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NP-hardness is an obstacle to finding good clusterings.

- *k*-means and *k*-median are NP-hard, and hard to approximate in general graphs
- *k*-means, *k*-median can be approximated in \mathbb{R}^d but seem to need time exponential in *d*
- Same is true for Bregman clustering[CM08]

Target And Optimal Clusterings

What happens if target clustering and optimal clustering are not the same ?



The two distance functions might be incompatible.

Target And Optimal Clusterings

What happens if target clustering and optimal clustering are not the same ?



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The two distance functions might be incompatible.

An instance is *stable* if approximating the cost function gives us a solution close to the target clustering.

View 1: If we perturb inputs, the output should not change.

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An instance is *stable* if approximating the cost function gives us a solution close to the target clustering.

- View 1: If we perturb inputs, the output should not change.
- View 2: If we change the distance function, output should not change.
- View 3: If we change the cost quality of solution, then output should not change.

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Stability I: Perturbing Inputs

Well separated sets:

Data is (k, α) -well separated if the best clustering that uses k - 1 clusters has cost that is $\geq 1/\alpha \cdot \text{OPT}$.



Two interesting properties[ORSS06]:

- All optimal clusterings mostly look the same: d_q small $\Rightarrow d$ small.
- Small perturbations of the data don't change this property.

Computationally, well-separatedness makes *k*-means work well

Definition (α -perturbations[BL09])

A clustering instance (P,d) is α -perturbation-resilient if the optimal clustering is identical to the optimal clustering for any (P,d'), where

 $d(x,y)/\alpha \le d'(x,y) \le d(x,y) \cdot \alpha$

- The smaller the *α*, the more resilient the instance (and the more "stable")
- Center-based clustering problems (*k*-median, *k*-means, *k*-center) can be solved optimally for √3-perturbation-resilient inputs[ABS10]

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Definition ((c, ϵ)-property[BBG09])

Given an input, all clusterings that are c-approximate are also ϵ -close.

Surprising facts:

- Finding a *c*-approximation in general might be NP-hard.
- Finding a *c*-approximation here is easy !



- If near-optimal clusters are close to true answer, then clusters must be well-separated.
- If clusters are well-separated, then choosing the right threshold separates them cleanly.
- Important that **ALL** near-optimal clusterings are close to true answer.

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- If clusters are well-separated, then choosing the right threshold separates them cleanly.
- Important that **ALL** near-optimal clusterings are close to true answer.

Theorem

In polynomial time, we can find a clustering that is $O(\epsilon)$ -close to the target clustering, even if finding a c-approximation is NP-hard.

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Strong assumption: **ALL** near-optimal clusterings are close to true answer.

Variant[ABS10]: Only consider *Voronoi*-based clusterings, where each point is assigned to *nearest* cluster center.



Same results hold as for previous case.

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Generalization

Strong assumption: **ALL** near-optimal clusterings are close to true answer.

Variant[ABS10]: Only consider *Voronoi*-based clusterings, where each point is assigned to *nearest* cluster center.

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Same results hold as for previous case.

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Wrap Up

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Theory of Clustering
- We understand much more about the behavior of *k*-means, and why it does well in practice.
- A simple initialization procedure for *k*-means is both effective and gives provable guarantees
- Much of the theoretical machinery around *k*-means works for the generalization to Bregman divergences.
- New and interesting questions on the relationship between the target clustering and cost measures used to get near it: ways of subverting NP-hardness.

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Slides for this tutorial can be found at

http://www.cs.utah.edu/~suresh/web/2010/05/08/ new-developments-in-the-theory-of-clustering-tutorial/

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