K-means Clustering and Extensions

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Introduction to Machine Learning (CS771A)

September 13, 2018



Announcement: Mid-Sem Exam

- September 20 (Thursday) 13:00-15:00
- Venue: L17, L18, L19 (all OROS)
- Syllabus: Up to what we see today (only 1-2 small questions from today's lecture)
- Open notes, open slides (please only print in 4-up mode), books not allowed
- No electronic items allowed (please keep phones switched off)
- Important: ALL answers have to be written on the question paper itself (dedicated space)
- Important: You need to bring a few other things
 - A notebook for rough work (may use blank pages from your notes)
 - Pen, pensil and eraser (but final answers should be written with pen to avoid smudging)
- A review session: September 15/16/17 (timing/venue TBD)



Recap: Speeding Up Kernel Methods

• Can extract "good" features $\psi(\mathbf{x}) \in \mathbb{R}^L$ from a kernel k (with mapping ϕ) such that

$$\psi(\boldsymbol{x}_n)^{\top}\psi(\boldsymbol{x}_m) \approx \phi(\boldsymbol{x}_n)^{\top}\phi(\boldsymbol{x}_m)$$

- ullet Unlike the kernel's original mapping ϕ , the mapping ψ is low-dimensional (L is typically small)
- With these features $\psi(\mathbf{x}_n)$, we can apply a linear model (both train/test). No need to kernelize.
- ullet Looked at two main approaches to get such an approximate mapping ψ
- ullet Landmark based approach: Using landmark points z_1, \dots, z_L (selected or learned), compute

$$\psi(\mathbf{x}_n) = [k(\mathbf{z}_1, \mathbf{x}_n), k(\mathbf{z}_2, \mathbf{x}_n), \dots, k(\mathbf{z}_L, \mathbf{x}_n)]$$

• Kernel Random Features approach: Can be used for many kernels. For the RBF kernel

$$\psi(\mathbf{x}_n) = \frac{1}{\sqrt{L}} [\cos(\mathbf{w}_1^{\top} \mathbf{x}_n + b_1), \dots, \cos(\mathbf{w}_L^{\top} \mathbf{x}_n + b_L)]$$

$$\mathbf{w}_{\ell} \sim \mathcal{N}(0, \lambda^{-1} \mathbf{I}_D), \quad b_{\ell} \sim \mathsf{Unif}(0, 2\pi), \quad \ell = 1, \dots, L$$

• Some other approaches (that we didn't see): Nyström approx, other low-rank kernel matrix approx

Recap: K-means Algorithm

- Goal: Assign N inputs $\{x_1, \dots, x_N\}$, with each $x_n \in \mathbb{R}^D$, to K clusters (flat partitioning)
- Notation: $z_n \in \{1, ..., K\}$ or z_n is a K-dim one-hot vector($z_{nk} = 1$ and $z_n = k$ mean the same)

K-means Algorithm

- Initialize K cluster means μ_1, \ldots, μ_K
- **2** For n = 1, ..., N, assign each point x_n to the closest cluster

$$z_n = \operatorname{arg\,min}_{k \in \{1, \dots, K\}} ||x_n - \mu_k||^2$$

3 Suppose $C_k = \{x_n : z_n = k\}$. Re-compute the means

$$\mu_k = \mathsf{mean}(\mathcal{C}_k), \quad k = 1, \dots, K$$

- Go to step 2 if not yet converged
- \bullet Note: The basic K-means models each cluster only by a mean μ_k . Ignores size/shape of clusters

The K-means Algorithm: Some Comments

- One of the most popular clustering algorithms
- Very widely used, guaranteed to converge (to a local minima; will see a proof)
- Can also be used as a sub-routine in graph clustering (in the Spectral Clustering algorithm)
- Has some shortcomings (as we will see) but can be improved upon
- Some of the many improvements (some of which we will see)
 - Can be kernelized (using kernels or using kernel-based landmarks/random features)
 - More flexible cluster sizes/shapes via probabilistic models (e.g., every cluster is a Gaussian)
 - Soft-clustering (fractional/probabilistic memberships): z_n is a probability vector
 - Overlapping clustering a point can belong to multiple clusters: z_n is a binary vector
 - .. even deep learning based K-means :-)
- .. so it is worth looking a bit deeply into what K-means is doing



K-means Loss Function: Several Forms, Same Meaning!

Notation: **X** is $N \times D$, **Z** is $N \times K$ (each row is a one-hot z_n), μ is $K \times D$ (each row is a μ_k)

$$\mathcal{L}(\mathbf{X},\mathbf{Z},\boldsymbol{\mu}) = \sum_{n=1}^{N} ||\boldsymbol{x}_n - \boldsymbol{\mu}_{z_n}||^2 \qquad \qquad \mathcal{L}(\mathbf{X},\mathbf{Z},\boldsymbol{\mu}) = \sum_{k=1}^{K} \sum_{\substack{n:z_n = k \\ \text{within cluster variance}}} ||\boldsymbol{x}_n - \boldsymbol{\mu}_k||^2$$

$$\mathcal{L}(\mathbf{X},\mathbf{Z},\boldsymbol{\mu}) = \sum_{n=1}^{N} \sum_{k=1}^{K} z_{nk} ||\boldsymbol{x}_n - \boldsymbol{\mu}_k||^2 \qquad \qquad \mathcal{L}(\mathbf{X},\mathbf{Z},\boldsymbol{\mu}) = \underbrace{||\mathbf{X} - \mathbf{Z}\boldsymbol{\mu}||_F^2}_{\text{as matrix factorization}}$$

$$\{\hat{\mathbf{Z}},\hat{\boldsymbol{\mu}}\} = \arg\min_{\mathbf{Z},\boldsymbol{\mu}} \mathcal{L}(\mathbf{X},\mathbf{Z},\boldsymbol{\mu}) \qquad \text{Total "distortion" or reconstruction error}$$

Note: Replacing ℓ_2 squared (Euclidean) distance by absolute (ℓ_1) distance gives the K-medians algorithm (more robust to outliers)

Note: Most unsup. learning algos try to minimize the distortion or reconstruction error of X from Z

Optimizing the *K***-means Loss Function**

• So the K-means problem is

$$\{\hat{\mathbf{Z}}, \hat{\boldsymbol{\mu}}\} = \arg\min_{\mathbf{Z}, \boldsymbol{\mu}} \mathcal{L}(\mathbf{X}, \mathbf{Z}, \boldsymbol{\mu}) = \arg\min_{\mathbf{Z}, \boldsymbol{\mu}} \sum_{n=1}^{N} \sum_{k=1}^{K} z_{nk} ||\boldsymbol{x}_n - \mu_k||^2$$

ullet Can't optimize it jointly for **Z** and μ . Let's try alternating optimization for **Z** and μ

Alternating Optimization for K-means Problem

f 1 Fix μ as $\hat{\mu}$ and find the optimal f Z as

$$\hat{\mathbf{Z}} = \arg\min_{\mathbf{Z}} \mathcal{L}(\mathbf{X},\mathbf{Z},\hat{\boldsymbol{\mu}})$$
 (still not easy - next slide)

② Fix **Z** as $\hat{\mathbf{Z}}$ and find the optimal μ as

$$\hat{m{\mu}} = rg \min_{m{\mu}} \mathcal{L}(\mathbf{X}, \hat{\mathbf{Z}}, m{\mu})$$

3 Go to step 1 if not yet converged

Solving for Z

• Solving for **Z** with μ fixed at $\hat{\mu}$

$$\hat{\mathbf{Z}} = \arg\min_{\mathbf{Z}} \mathcal{L}(\mathbf{X}, \mathbf{Z}, \hat{\boldsymbol{\mu}}) = \arg\min_{\mathbf{Z}} \sum_{n=1}^{N} \sum_{k=1}^{K} z_{nk} ||\boldsymbol{x}_n - \hat{\mu}_k||^2$$

- Still not easy. Since **Z** is discrete, it is an NP-hard problem
 - Combinatorial optimization: K^N possibilities for **Z** ($N \times K$ matrix with one-hot rows)
- A greedy approach: Optimize **Z** one row (z_n) at a time keeping all others z_n 's (and μ) fixed

$$\hat{z}_n = \arg\min_{z_n} \sum_{k=1}^K z_{nk} ||x_n - \hat{\mu}_k||^2 = \arg\min_{z_n} ||x_n - \hat{\mu}_{z_n}||^2$$

- Easy to see that this is minimized by assigning x_n to the closest mean
 - This is exactly what the K-means algo does!



Solving for μ

• Solving for μ with **Z** fixed at $\hat{\mathbf{Z}}$

$$\hat{\boldsymbol{\mu}} = \arg\min_{\boldsymbol{\mu}} \mathcal{L}(\mathbf{X}, \hat{\mathbf{Z}}, \boldsymbol{\mu}) = \arg\min_{\boldsymbol{\mu}} \sum_{k=1}^K \sum_{n: \hat{z}_n = k} ||\boldsymbol{x}_n - \mu_k||^2$$

- This is not that hard to solve (μ_k 's are real-valued vectors, can optimize easily)
- Note that each μ_k can be optimized independently

$$\hat{\mu}_k = \arg\min_{\mu_k} \sum_{n: 2_n = k} ||\boldsymbol{x}_n - \mu_k||^2$$

- (Verify) This is minimized by setting $\hat{\mu}_k$ to be mean of points currently in cluster k
 - This is exactly what the K-means algo does!



Convergence of K-means Algorithm

- Each step (updating **Z** or μ) can **never increase** the K-means loss
- When we update **Z** from $\mathbf{Z}^{(t-1)}$ to $\mathbf{Z}^{(t)}$

$$\mathcal{L}(\mathsf{X}, \mathsf{Z}^{(t)}, \boldsymbol{\mu}^{(t-1)}) \leq \mathcal{L}(\mathsf{X}, \mathsf{Z}^{(t-1)}, \boldsymbol{\mu}^{(t-1)})$$

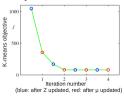
because the new $\mathbf{Z}^{(t)} = \arg\min_{\mathbf{Z}} \mathcal{L}(\mathbf{X}, \mathbf{Z}, \boldsymbol{\mu}^{(t-1)})$

ullet When we update μ from $\mu^{(t-1)}$ to $\mu^{(t)}$

$$\mathcal{L}(\mathbf{X}, \mathbf{Z}^{(t)}, \boldsymbol{\mu}^{(t)}) \leq \mathcal{L}(\mathbf{X}, \mathbf{Z}^{(t)}, \boldsymbol{\mu}^{(t-1)})$$

because the new $\mu^{(t)} = \arg\min_{\mu} \mathcal{L}(\mathbf{X}, \mathbf{Z}^{(t)}, \mu)$

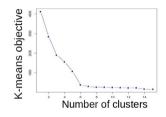
• Thus the K-means algorithm monotonically decreases the objective





K-means: Choosing *K*

• One way to select K for the K-means algorithm is to try different values of K, plot the K-means objective versus K, and look at the "elbow-point"



- For the above plot, K = 6 is the elbow point
- Can also information criterion such as AIC (Akaike Information Criterion)

$$AIC = 2\mathcal{L}(\hat{\boldsymbol{\mu}}, \mathbf{X}, \hat{\mathbf{Z}}) + KD$$

- .. and choose the K that has the smallest AIC (discourages large K)
- Several other approaches when using probabilistic models for clustering, e.g., comparing marginal likelihood $p(\mathbf{X}|K)$, using nonparametric Bayesian models, etc.

K-means: Hard vs Soft Assignments

- Makes hard assignments of points to clusters
 - A point either completely belongs to a cluster or doesn't belong at all
 - No notion of a soft assignment (i.e., probability of being assigned to each cluster: say K=3 and for some point x_n , $p_1=0.7$, $p_2=0.2$, $p_3=0.1$)





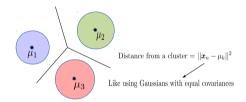
A heuristic to get soft assignments: Transform distances from clusters into probabilities

$$\gamma_{nk} = \frac{\exp(-||x_n - \mu_k||^2)}{\sum_{\ell=1}^K \exp(-||x_n - \mu_\ell||^2)} \quad \text{(prob. that } x_n \text{ belongs to cluster } k\text{)}$$

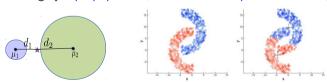
- These heuristics are used in "fuzzy" or "soft" K-means algorithms
- Soft K-means μ_k updates are slightly different: $\mu_k = \frac{\sum_{n=1}^N \gamma_{nk} \mathbf{x}_n}{\sum_{n=1}^N \gamma_{nk}}$ (all points used, but fractionally)

K-means: Decision Boundaries and Cluster Sizes/Shapes

- K-mean assumes that the decision boundary between any two clusters is linear
- Reason: The K-means loss function implies assumes equal-sized, spherical clusters



• Assumes clusters to be roughly equi-populated, and convex-shaped. Otherwise, may do badly



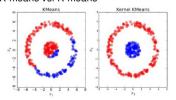
• Kernel K-means can help address some of these issues. Probabilistic models is another option

Kernel *K*-means

Basic idea: Replace the Euclidean distances in K-means by the kernelized versions

$$||\phi(\mathbf{x}_n) - \phi(\boldsymbol{\mu}_k)||^2 = ||\phi(\mathbf{x}_n)||^2 + ||\phi(\boldsymbol{\mu}_k)||^2 - 2\phi(\mathbf{x}_n)^{\top}\phi(\boldsymbol{\mu}_k)$$
$$= k(\mathbf{x}_n, \mathbf{x}_n) + k(\boldsymbol{\mu}_k, \boldsymbol{\mu}_k) - 2k(\mathbf{x}_n, \boldsymbol{\mu}_k)$$

- Here k(.,.) denotes the kernel function and ϕ is its (implicit) feature map
- Note: $\phi(\mu_k)$ is the average of ϕ 's the data points assigned to cluster kKernel K-means vs. K-means



Pyclust: Open Source Data Clustering Pokage

- Can also use landmark or random features approach to make it faster
 - Can then simply run the basic K-means on those features!



Going the Probabilistic Way..

- Assume a generative model for the inputs. Suppose ⊖ denotes all the unknown parameters
- Clustering then boils down to computing $p(z_n|x_n,\Theta)$ for each x_n , where z_n is a latent variable
- Using the Bayes rule, we can write $p(z_n|x_n,\Theta)$ as

$$p(z_n = k | x_n, \Theta) = \frac{p(z_n = k | \Theta)p(x_n | z_n = k, \Theta)}{p(x_n | \Theta)}$$

• Assuming $p(z|\Theta)$ as multinoulli(π) and each cluster as Gaussian $p(x|z=k,\Theta) = \mathcal{N}(x|\mu_k,\Sigma_k)$

$$p(\boldsymbol{z}_n = k | \boldsymbol{x}_n, \boldsymbol{\Theta}) \propto \pi_k \times \mathcal{N}(\boldsymbol{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$$
 Cluster assignment prob now depends on the number of points in cluster k points in cluster k (covariances)
$$(\text{here } \boldsymbol{\Theta} = \{\pi_k, \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k\}_{k=1}^K)$$

- We know how to estimate Θ for such problems.. if z_n is known (recall generative classification)
- The tricky part here is that we don't know z_n . How do we estimate Θ then?
- A solution: Take an alternating approach (like K-means)

Going the Probabilistic Way..

• At a high-level, a probabilistic clustering algorithm would look somewhat like this

Sketch of a Probabilistic Clustering Algorithm

- 1 Initialize the model parameters Θ somehow
- ② Given the current Θ , estimate **Z** (cluster assignments) in a soft/hard way

$$p(z_n = k | x_n, \Theta) = \gamma_{nk} = \frac{p(z_n = k | \Theta)p(x_n | z_n = k, \Theta)}{p(x_n | \Theta)}, \quad k = 1, \dots, K$$

$$OR \quad \hat{z}_n = \operatorname{arg max}_{k \in \{1, \dots, K\}} \gamma_{nk}$$

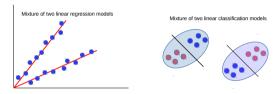
- Use $\{\hat{\mathbf{z}}_n\}_{n=1}^N$ (hard cluster labels) or $\{\gamma_{nk}\}_{n,k=1}^{N,K}$ (soft labels) to update Θ via MLE/MAP (similar to how we do for gen. classification where the labels are known)
- lacktriangle Note: The soft-label based Θ updates slightly more involved (wait until we see EM)
- Go to step 2 if not converged yet.
- The above algorithm is an instance of a more general Expectation Maximization (EM) algorithm for latent variable models (we will see this post mid-sem)

Clustering vs Classification

- Any clustering model typically learns two type of quantities
 - ullet Parameters Θ of the clustering model (e.g., cluster means $oldsymbol{\mu} = \{\mu_1, \dots, \mu_K\}$ in K-means)
 - Cluster assignments $\mathbf{Z} = \{z_1, \dots, z_N\}$ for the points
- If the cluster assignments Z are known, learning the parameters Θ is just like learning the parameters of a classification model (typically generative classification) using labeled data
- Therefore it helps to think of clustering as (generative) classification with unknown labels
- This equivalence is very important and makes it possible to solve clustering problems
- Therefore many clustering problems are typically solved in the following fashion
 - Initialize \(\text{\text{\text{o}}}\) somehow
 - **2** Predict **Z** given current estimate of Θ
 - \bullet Use the predicted **Z** to improve the estimate of Θ (like learning a generative classification model)
 - Go to step 2 if not converged yet

Clustering can help supervised learning, too

- Often "difficult" supervised learning problems can be seen as mixture of simpler models
- Example: Nonlinear regression or nonlinear classification as mixture of linear models



- An alternative to kernel methods and deep learning :-)
- Don't know which point belongs to which linear model ⇒ Clustering problem
- Can therefore solve such problems as follows
 - 1 Initialize each linear model somehow (maybe randomly)
 - Cluster the data by assigning each point to its "closest" linear model
 - (Re-)Learn a linear model for each cluster's data. Go to step 2 if not converged.
- Often called Mixture of Experts models. Will look at these more formally after mid-sem

