HW4: K-Means and Gaussian Mixture Models

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Due date: Thu Apr 06 at 11:59pm ET

How to turn in: Submit PDF to https://www.gradescope.com/courses/496674/assignments/2782332

Jump to: Problem 1 Problem 2 Problem 3 Problem 4

Questions?: Post to the hw4 topic on the Piazza discussion forums.

Instructions for Preparing your PDF Report

What to turn in: PDF of typeset answers via LaTeX. No handwritten solutions will be accepted, so that grading can be speedy and you get prompt feedback.

Please use provided LaTeX Template: https://github.com/tufts-ml-courses/cs136-23s-assignments/blob/main/unit4 HW/hw4_template.tex

Your PDF should include (in order):

- Cover page with your full name, estimate of hours spent, and Collaboration statement
- Problem 1a, 1b, 1c, 1d, 1e
- Problem 2a, 2b
- Problem 3a
- Problem 4a is OPTIONAL. Worth up to 6 points back on other parts (cannot go higher than 100%).

When you turn in the PDF to gradescope, mark each part via the in-browser Gradescope annotation tool)

Problem 1: K-means walk-through

Recall that K-means minimizes the following cost function:

$$J(x_{1:N}, r_{1:N}, \mu_{1:K}) = \sum_{n=1}^{N} \sum_{k=1}^{K} r_{nk} (x_n - \mu_k)^T (x_n - \mu_k)$$

where each assignment variable r_n is a one-hot vector of size $K\!$.

The K-means algorithm is specified in pseudocode as:

Inputs:

- $x_1, \dots x_N$: Training dataset
- $\mu_{1:K}^0$: Initial guess of cluster center locations

Procedure:

For iter t in 1, 2, ... until converged:

1.
$$r_{1:N}^t \leftarrow \arg\min_{r_{1:N}} J(x_{1:N}, r_{1:N}, \mu_{1:K}^{t-1})$$

2. $\mu_{1:K}^t \leftarrow \arg\min_{\mu_{1:K}} J(x_{1:N}, r_{1:N}^t, \mu_{1:K})$

Consider running K-means on the following dataset of N=7 examples, in this (N, D)-shaped array

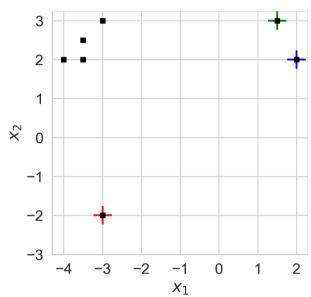
```
x_ND = array([
  [-3.0,-2.0],
  [-4.0, 2.0],
  [-3.5, 2.5],
  [-3.5, 2.0],
  [-3.0, 3.0],
  [ 1.5, 3.0],
  [ 2.0, 2.0]])
```

When the initial cluster locations are given by the following K=3 cluster locations, in this (K, D)-shaped array:

```
mu_KD = array([
  [-3.0,-2.0],
  [ 1.5, 3.0],
  [ 2.0, 2.0]])
```

We'll denote these initial locations mathematically as μ^0 . Here and below, we'll use *superscripts* to indicate the specific *iteration* of the algorithm at which we ask for the value, and we'll assume that iteration 0 corresponds to the initial configuration.

We've visualized the 7 data examples (black squares) and the 3 initial cluster locations (crosses) in this figure:



Plot of the N=7 toy data examples (squares) and K=3 initial cluster locations (crosses).

Problem 1a: Find the optimal one-hot assignment vectors r^1 for all N=7 examples, when given the initial cluster locations μ^0 . This corresponds to executing step 1 of K-means algorithm. Report the value of the cost function $J(x, r^1, \mu^0)$.

Problem 1b: Find the optimal cluster locations μ^1 for all K=3 clusters, using the optimal assignments r^1 you found in 2a. This corresponds to executing step 2 of K-means algorithm. Report the value of the cost function $J(x, r^1, \mu^1)$.

Problem 1c: Find the optimal one-hot assignment vectors \mathbf{r}^2 for all N=7 examples, using the cluster locations μ^1 from 2b. Report the value of the cost function $J(x, \mathbf{r}^2, \mu^1)$.

Problem 1d: Find the optimal cluster locations μ^2 for all K=3 clusters, using the optimal assignments r^2 you found in 2c. Report the value of the cost function $J(x, r^2, \mu^2)$.

Problem 1e: What interesting phenomenon do you see happening in this example regarding cluster 2? How could you set cluster 2's location after part d above to better fulfill the goals of K-means (find K clusters that reduce cost the most)?

Problem 2: Relationship between GMM and K-means

Bishop's PRML textbook Sec. 9.3.2 describes a technical argument for how a GMM can be related to the K-means algorithm. In this problem, we'll try to make this argument concrete for the same toy dataset as in Problem 1.

To begin, given any GMM parameters, we can use Bishop PRML Eq. 9.23 to compute the posterior probability of assigning each example x_n to cluster k via the formula:

$$\begin{aligned} \gamma_{nk} &\triangleq p(z_{nk} = 1 | x_n) \\ &= \frac{\pi_k \quad (x_n | \mu_k, \Sigma_k)}{\sum_{j=1}^K \ \pi_j \quad (x_n | \mu_j, \Sigma_j)} \end{aligned}$$

Now, imagine a GMM with the following *concrete* parameters:

- mixture weights $\pi_{1:K}$ set to the uniform distribution over K=3 clusters
- covariances $\Sigma_{1:K}$ set to εI_D for all clusters, for some $\varepsilon>0$

We can leave the locations $\mu_{1:K}$ at any valid values.

Problem 2a: Show (with math) that using the parameter settings defined above, the general formula for γ_{nk} will simplify to the following (inspired by PRML Eq. 9.42):

$$\gamma_{nk} = \frac{\exp(-\frac{1}{2\epsilon}(x_n - \mu_k)^T(x_n - \mu_k))}{\sum_{j=1}^{K} \exp(-\frac{1}{2\epsilon}(x_n - \mu_k)^T(x_n - \mu_k))}$$

Problem 2b: What will happen to the vector γ_n as $\epsilon \to 0$? How is this related to K-means?

Hint 2(i): Try it out concretely on the toy data from Problem 1 above.

Hint 2(ii): No need for a formal proof here. Just show you understand what happens in the limit, not why.

Problem 3: Covariances of mixtures

Background: Consider a continuous random variable x which is a vector in D-dimensional space: $x \in \mathbb{R}^{|D|}$.

We assume that x follows a mixture distribution with PDF p^{mix} , using K components indexed by integer k:

$$p^{mix}(x|\pi,\mu,\Sigma) = \sum_{k=1}^K \pi_k f_k(x|\mu_k,\Sigma_k)$$

The k-th component has a mixture "weight" probability of π_k . Across all K components, we have a parameter $\pi = [\pi_1 \ \pi_2 \ \dots \ \pi_K]$, whose entries are non-negative and sum to one.

The k-th component has a specific data-generating PDF f_k . We don't know the functional form of this PDF (it could be Gaussian, or something else), and the form could be different for every k. However, we do know that this PDF f_k takes two parameters, a vector $\mu_k \in \mathbb{R}^D$ and a matrix Σ_k which is a $D \times D$ symmetric, positive definite matrix. We further know that these parameters represent the mean and covariance of vector x under the pdf f_k :

$$\mathbb{E}_{(f_k(x|\mu_k,\Sigma_k)}[x] = \mu_k$$

$$Cov_{(f_k(x|\mu_k,\Sigma_k)}[x] = \Sigma_k$$

Problem 3a: Prove that the covariance of vector **x** under the mixture distribution is given by:

$$Cov_{p^{mix}(x)}[x] = \sum_{k=1}^{K} \pi_k(\Sigma_k + \mu_k \mu_k^T) - mm^T$$

where we define $m = \mathbb{E}_{p^{mix(x)}}[x]$.

Hint 3(i): We know a closed-form for m: $m = \sum_{k=1}^{K} \, \pi_{\!k} \mu_{\!k}.$

Hint 3(ii): For any random vector x, we know: $\mathbb{E}[xx^T] = \text{Cov}(x) + \mathbb{E}[x]\mathbb{E}[x]^T$

Problem 4: Jensen's Inequality and KL Divergence

Optional. Not required.

Background reading

Skim Bishop PRML's Sec. 1.6 ("Information Theory"), which introduces several key concepts useful for the EM algorithm, including:

- Entropy
- Jensen's inequality
- KL divergence

Background: Negative logarithms are convex

Consider the negative logarithm function: $f(a) = -\log a$, for inputs a > 0. Recall that f(a) is a *convex* function, because its second derivative is always positive:

$$f(a) = -\log a,$$

 $f'(a) = -a^{-1},$
 $f''(a) = a^{-2},$ therefore: $f''(a) > 0$ for all $a > 0$.

Background: Jensen's inequality for negative logarithms

Now, suppose we have a random variable A that takes one of K possible values.

Define each candidate value $a_k > 0$, and let its associated probability be $r_k \in [0,1]$. Writing the probabilities as a vector $\mathbf{r} = [r_1, \dots r_K]$, we know these non-negative values must sum to one: $\mathbf{r} \in \Delta^K$.

We are interested in the expected value of f(A), where f is the negative logarithm. We can derive the following bound using **Jensen's inequality** (see PRML textbook Eq. 1.115),

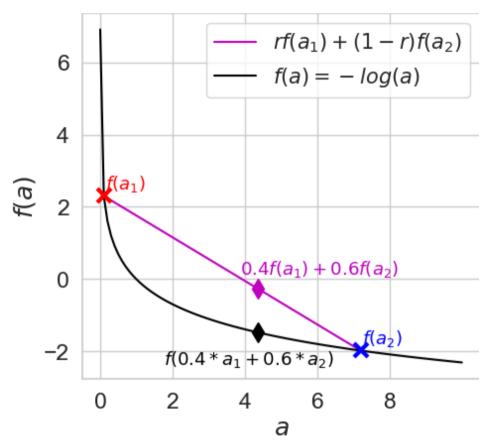
$$\mathbb{E}[f(A)] \ge f(\mathbb{E}[A])$$

Expanding out these expectations and invoking f 's definition as the negative log (which only takes positive inputs), we have:

$$\sum_{k=1}^{K} r_k [-\log a_k] \ge -\log \left[\sum_{k=1}^{K} r_k a_k \right]$$

This bound holds for any positive vector \mathbf{a} and any probability vector \mathbf{r} .

We can visualize this Jensen bound in the following figure, using two selected points $a_1 = 0.1$ and $a_2 = 7.2$.



Plot of our convex function of interest ("f", black) and its *linear interpolation* (magenta) between outputs that correspond to two inputs "a1" and "a2". Clearly, function f is a *lower bound* of its interpolation (magenta). In terms of probabilities, this means π \mathbb{E}[f(A)] \geq f(\mathbb{E}[A])\$

Notation setup

We'll use one-hot indicator vectors here. Let e_k denote the one-hot vector of size K where entry k is non-zero.

Define random variable z as a one-hot indicator vector of size K. So, the K possible values of z are $\{e_1, e_2, \dots e_K\}$.

Define two possible Categorical distributions over z, denoted q and p.

$$\begin{split} q(z) &= CatPMF(z|r_1,\ldots r_K) = \prod_{k=1}^K r_k^{z_k}, & q(z=e_k) = r_k, & r_k > 0 \ \forall k \\ \\ p(z) &= CatPMF(z|\pi_1,\ldots \pi_K) = \prod_{k=1}^K \pi_k^{z_k}, & p(z=e_k) = \pi_k, & \pi_k > 0 \ \forall k \end{split}$$

Each uses an all positive probability vector parameters $\mathbf{r} \in \Delta_+^K$ and $\pi \in \Delta_+^K$. Here Δ_+^K denotes the set of K-length vectors whose sum is one and whose entries are all *strictly positive*.

The KL divergence from q to p is defined as:

$$KL(q(z)||p(z)) \triangleq \mathbb{E}_{q(z)}[-\log \frac{p(z)}{q(z)}]$$

Problem statement

Problem 4a: Consider any two Categorical distributions q(z) and p(z) that assign *positive* probabilities over the same size-K sample space. Show that their KL divergence is non-negative.

That is, show that $KL(q(z)||p(z)) \ge 0$, or equivalently that

 $KL (CatPMF(z|\mathbf{r})||CatPMF(z|\pi)) \ge 0$

when $\mathbf{r} \in \Delta_+^K$ and $\pi \in \Delta_+^K$.

Hint: Expand the definition of KL as an expectation out so it is purely an evaluatable function of r and π , then use Jensen's inequality for negative logarithms.

Note: it is possible to prove the KL is non-negative even when some entries in r or π are exactly zero, but this requires taking some limits rather carefully, and we want you to avoid that burdensome detail. Thus, here we consider r and π as having all positive entries.

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