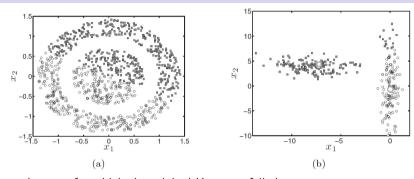
Clustering - Gaussian mixture model

Rohit Budhiraja

Machine Learning for Wireless Communications (EE798L)

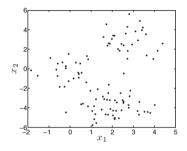
March 4, 2024

Recap and agenda of today's class



- Figure shows datasets for which the original K-means failed
- Problem with K-means algorithm is that its definition of a cluster was too crude
 - Characteristics of stretched clusters cannot be represented by a single point and squared distance
- Need to incorporate notion of shape
 - Statistical mixtures represent each cluster as a probability density
- Probabilistic mixture models helps in clustering a wide variety of shapes in almost any type of data
- Probabilistic mixture models helps in determine number of clusters

Mixture model – generative process



- Our earlier clustering dataset how could we generate data that looks like this?
- Above data does not look like samples from any density function that we have encountered
 - There appear to be three disjoint regions in which data are concentrated
 - None of the density functions that we have seen can produce data with this complex structure
- However, each of the three regions looks simple enough to generate on its own
 - In fact, they all look a bit like samples from two-dimensional Gaussians

Mixture model – generative process

- Assuming that data was generated by three separate Gaussians we propose
- Two-step procedure for sampling the nth data object \mathbf{x}_n :
 - Select one of the three Gaussians
 - 2 Sample x_n from this Gaussian
- Both steps are straightforward. Step 1 chooses one value from a discrete set, like rolling a die
 - ullet To do this, we just need to define the probability of each outcome π_k such that $\sum_k \pi_k = 1$
- Having chosen which Gaussian to sample from, the second step is straightforward
- As in K-means, we will use z_{nk} as an indicator variable
 - If we choose kth component as the source of nth object, we set $z_{nk}=1$, and $z_{nj}=0$ for all $j\neq k$
- We will use μ_k and Σ_k to denote the parameters of kth Gaussian
- Density function for \mathbf{x}_n given that it was produced by the kth component $(z_{nk}=1)$ is
 - ullet Gaussian with mean and covariance $oldsymbol{\mu}_k$ and $oldsymbol{\Sigma}_k$, respectively

$$p\left(\mathbf{x}_{n} \mid z_{nk} = 1, \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right) = \mathcal{N}\left(\boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right)$$

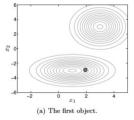


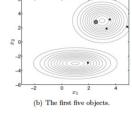
Mixture model – generative process (2)

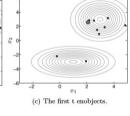
- ullet To illustrate this process, we will sample some data from a setup with K=2 Gaussians
- We will use the following means and covariances for the two components

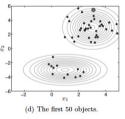
$$\boldsymbol{\mu}_1 = \begin{bmatrix} 3, 3 \end{bmatrix}^\mathsf{T}, \ \boldsymbol{\Sigma}_1 = \begin{bmatrix} 1 & 0 \\ 0 & 2 \end{bmatrix} \qquad \boldsymbol{\mu}_2 = \begin{bmatrix} 1, -3 \end{bmatrix}^\mathsf{T}, \ \boldsymbol{\Sigma}_2 = \begin{bmatrix} 2 & 0 \\ 0 & 1 \end{bmatrix}$$

ullet Finally, we need to define π_k : assume component 1 is more likely, we will use $\pi_1=0.7$ and $\pi_2=0.3$



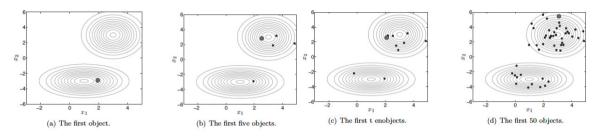






- Figure shows first 50 generated data objects and the density functions of the two Gaussians
- For first point in Figure (a), k = 2 is chosen and object sampled from second (lower) component
- Figure (b) shows the first five objects (the most recent is always denoted as a larger circle)
 - All but the first one have come from the first component because $\pi_1 > \pi_2$

Mixture model – generative process (3)



- Data in Fig. (d) looks similar to our earlier figure
 - Generative procedure generates data by sampling it from a mixture of individual density functions
- Mixture models are widely used in data modelling
 - Fitting a set of simple distributions is often more straightforward than fitting one more complex one
- Learning task: Infer, from observed data
 - Component parameters (μ_k, Σ_k) and assignments of objects to components

Mixture model - learning objective

- As with K- means, this is a circular argument:
 - component parameters would be easy to compute if we knew the assignments, and
 - assignments would be easy to compute if we knew the component parameters
- Without either, it is hard to know where to start
- Answer comes in form of Expectation-Maximisation (EM) algorithm parallel to K-means algorithm
- EM algorithm iteratively maximizes the likelihood, and used for a wide range of models
- Develop EM algorithm as general as possible: we will work with $p(\mathbf{x}_n \mid z_{nk} = 1, \Delta_k)$
 - Δ_k denotes parameters of the kth density (not necessarily Gaussian)
 - Δ will denote the collection of parameters of all mixture components $\Delta = \{\Delta_1, \cdots, \Delta_K\}$
 - Use $\pi = \{\pi_1, \dots, \pi_K\}.$
- Require likelihood of data objects \mathbf{x}_n under the whole model: $p(\mathbf{x}_n \mid \Delta, \pi)$
- We start with likelihood of a particular data object conditioned on $z_{nk} = 1$:

$$p(\mathbf{x}_n \mid z_{nk} = 1, \Delta) = p(\mathbf{x}_n \mid \Delta_k)$$



Mixture model likelihood

- To obtain $p(\mathbf{x}_n \mid \Delta, \pi)$, we need to get rid of z_{nk}
- To do this, we first multiply both sides by $p(z_{nk})$, which we have defined as π_k :

$$p(\mathbf{x}_{n} \mid z_{nk} = 1, \Delta) p(z_{nk} = 1) = p(\mathbf{x}_{n} \mid \Delta_{k}) p(z_{nk} = 1)$$

$$p(\mathbf{x}_{n}, z_{nk} = 1 \mid \Delta, \pi) = p(\mathbf{x}_{n} \mid \Delta_{k}) \pi_{k}$$

 \bullet Summing both sides over k (marginalising over the individual components) yields

$$\sum_{k=1}^{K} \rho(\mathbf{x}_{n}, z_{nk} = 1 \mid \Delta, \pi) = \sum_{k=1}^{K} \rho(\mathbf{x}_{n} \mid \Delta_{k}) \pi_{k}$$
$$\rho(\mathbf{x}_{n} \mid \Delta, \pi) = \sum_{k=1}^{K} \pi_{k} \rho(\mathbf{x}_{n} \mid \Delta_{k})$$

 \bullet Making standard independence assumption, we can extend this to likelihood of all N data objects:

$$p(\mathbf{X} \mid \Delta, \pi) = \prod \sum \pi_k p(\mathbf{x}_n \mid \Delta_k)$$

ullet Maximise log likelihood to calculate optimal parameter $\mu_k, oldsymbol{\Sigma}_k, \pi$

$$L = \log p(\mathbf{X} \mid \Delta, \pi) = \sum_{n=1}^{N} \log \sum_{k=1}^{N} \pi_{k} p(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k})) \tag{1}$$

Fundamental theorem of expectation

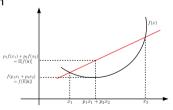
- Summation inside log term complicates maximization helps comes in form of Jensen inequality
- X is a random variable then $\mathbb{E}X$ is its expectation:

$$\mathbb{E}X = \sum_{x \in \mathcal{X}} x p(x)$$

- Let $X = x_1$ with prob. p_1 and $X = x_2$ with prob. p_2 , then $\mathbb{E}X = p_1x_1 + p_2x_2$ with $p_1 + p_2 = 1$
- Recall fundamental theorem of expectation

$$\mathbb{E}f(X) = \sum_{x \in \mathcal{X}} f(x)p(x) = p_1 f(x_1) + p_2 f(x_2)$$

Let's understand a convex function



$$p_1f(x_1) + p_2f(x_2) \ge f(p_1x_1 + p_2x_2)$$

Jensen inequality

• Jensen inequality: If f is a convex function and X is a random variable then

$$\mathbb{E}f(X) \geq f(\mathbb{E}X)$$

• Proof: Let $X = x_1$ with probability p_1 and $X = x_2$ with probability p_2 , then for convex f

$$p_1 f(x_1) + p_2 f(x_2) \ge f(p_1 x_1 + p_2 x_2)$$
 definition of convexity $\mathbb{E} f(X) \stackrel{\text{(a)}}{\ge} f(\mathbb{E} X)$

- Can be proved using induction for larger number of mass points
- Inequality sign will change for concave functions

$$f(\mathbb{E}X) \geq \mathbb{E}f(X)$$



Simplification of log likelihood using Jensen inequality (1)

• We shall now demonstrate the use of the EM algorithm to maximise the log likelihood

$$L = \log p(\mathbf{X} \mid \Delta, \boldsymbol{\pi}) = \sum_{n=1}^{N} \log \sum_{k=1}^{K} \pi_{k} p(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}))$$
 (2)

- Summation inside logarithm makes finding optimal parameter μ_k, Σ_k, π difficult
- EM algorithm overcomes this problem by deriving a lower bound on this likelihood
 Instead of maximising L directly, we instead maximise the lower bound
- To obtain a lower bound on L we use Jensen's inequality

$$\log \mathbb{E}_{p(z)}\{f(z)\} \geq \mathbb{E}_{p(z)}\{\log f(z)\}$$

- To use Jensen's inequality, we need to make RHS of (2) look like the log of an expectation.
- ullet To do this, we multiply and divide expression inside summation over k by a new variable q_{nk}

$$L = \sum_{n=1}^{N} \log \sum_{k=1}^{K} \pi_{k} p\left(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right) \frac{q_{nk}}{q_{nk}}$$

• If we restrict q_{nk} to be positive and satisfy the summation constraint $\sum_{k=1}^{K} q_{nk} = 1$

Simplification of log likelihood using Jensen inequality (2)

• q_{nk} is some probability distribution over the K components for the nth object

$$L = \sum_{n=1}^{N} \log \sum_{k=1}^{K} q_{nk} \frac{\pi_{k} p\left(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right)}{q_{nk}} = \sum_{n=1}^{N} \log \mathbb{E}_{q_{nk}} \left\{ \frac{\pi_{k} p\left(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right)}{q_{nk}} \right\}$$

• Applying Jensen's inequality, we can lower bound this expression:

$$L = \sum_{n=1}^{N} \log \mathbb{E}_{q_{nk}} \left\{ \frac{\pi_{k} p\left(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right)}{q_{nk}} \right\} \geq \sum_{n=1}^{N} \mathbb{E}_{q_{nk}} \left\{ \log \frac{\pi_{k} p\left(\mathbf{x}_{n} \mid \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}\right)}{q_{nk}} \right\}$$

- ullet RHS of this expression is the bound (we will denote it ${\cal B}$) that we shall optimise
- Expanding the expression gives us something more manageable:

$$\mathcal{B} = \sum_{n=1}^{N} \mathbb{E}_{q_{nk}} \left\{ \log \frac{\pi_k p\left(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k\right)}{q_{nk}} \right\} = \sum_{n=1}^{N} \sum_{k=1}^{K} q_{nk} \log \left(\frac{\pi_k p\left(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k\right)}{q_{nk}} \right)$$

$$= \sum_{n=1}^{N} \sum_{k=1}^{K} q_{nk} \log \pi_k + \sum_{n=1}^{N} \sum_{k=1}^{K} q_{nk} \log p\left(\mathbf{x}_n \mid \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k\right) - \sum_{n=1}^{N} \sum_{k=1}^{K} q_{nk} \log q_{nk}$$
(3)

• Calculate $q_{nk}, \mu_k, \Sigma_k, \pi$ to find a local maxima of this bound • Values will correspond to a local maxima of log-likelihood L

