

# CS 236756 - Technion - Intro to Machine Learning

#### **Tal Daniel**

# **Tutorial 09 - Expectation Maximization**

• Based on Pattern Recognition and Machine Learning by Christopher Bishop (chapter 9) and slides by Shai Fine



## Agenda

- Demonstrations
  - K-means
  - Gaussian Mixture Models (GMM)
- · Expectation-Maximization
  - Formalization
- Gaussian Mixture Models (GMM)
  - Overview
- · Binomial Mixture Model (BMM)
- · K-means as "Hard GMM"



# **Great Video Explaining EM Algorithm**

Just in case you need some more explanations and visualizations, a great series of videos covering GMMs and the EM algorithm:

- Part 1 (7:53 min) (https://www.youtube.com/watch?v=REypj2sy\_5U)
- Part 2 (10:39 min) (https://www.youtube.com/watch?v=iQoXFmbXRJA)
- Part 3 (3:05 min) (https://www.youtube.com/watch?v=TG6Bh-NFhA0)
- Part 4 (3:29 min) (https://www.youtube.com/watch?v=zL MHtT56S0)
- Part 5 (10:53 min) (https://www.youtube.com/watch?v=BWXd5dOkuTo)

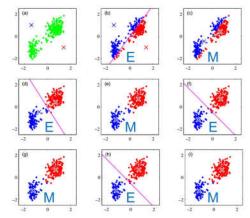
# Demonstrations

- · Definitions:
  - Hard Clustering clusters do not overlap, an element either belongs to a cluster or does not.
  - Soft Clustering clusters may overlap. Strength of association between clusters and instances (confidence level).

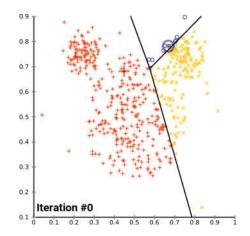


#### K-Means

- · K-means clustering is one of the simplest and popular unsupervised machine learning algorithms.
  - Typically, unsupervised algorithms make inferences from datasets using only input vectors without referring to known, or labeled, outcomes.
- The objective of K-means: group similar data points together and discover underlying patterns. To achieve this objective, K-means looks for a fixed number (k) of clusters in a dataset.
  - This is also called hard clustering as the association of a data point to a cluster is definitive, that is, it belongs to one cluster only and cannot be included in other cluster.
- · A cluster refers to a collection of data points aggregated together because of certain similarities.
- Illustration:



- E and M stand for the E-step and M-step
- Algorithm: Starts with a first group of randomly selected centroids, which are used as the beginning points for every cluster, and then performs
  iterative (repetitive) calculations to optimize the positions of the centroids. It halts creating and optimizing clusters when either:
  - The centroids have stabilized—there is no change in their values because the clustering has been successful.
  - The defined number of iterations has been achieved.
- Animation:



Explanations and example from <u>Understanding K-Means Clustering in Machine Learning - towardsdatascience.com</u> (<a href="https://towardsdatascience.com/understanding-k-means-clustering-in-machine-learning-6a6e67336aa1">https://towardsdatascience.com/understanding-k-means-clustering-in-machine-learning-6a6e67336aa1</a>)

In [1]: # imports for the tutorial
import numpy as np

import pandas as pd

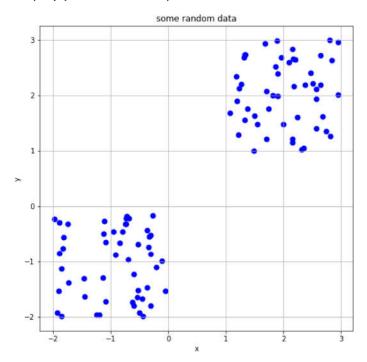
import matplotlib.pyplot as plt

%matplotlib notebook

```
In [3]: # k-means example
    from sklearn.cluster import KMeans
# generate random data

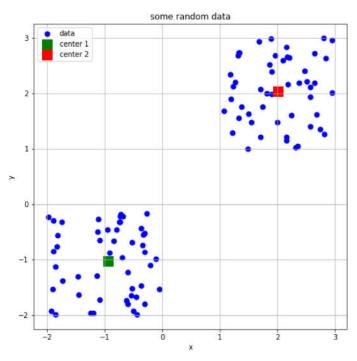
X = -2 * np.random.rand(100,2)
X[50:100, :] = 1 + 2 * np.random.rand(50,2)
    fig = plt.figure(figsize=(8,8))
    ax = fig.add_subplot(1,1,1)
    ax.scatter(X[:,0], X[:,1], s = 50, c = 'b')
    ax.grid()
    ax.set_xlabel("x")
    ax.set_ylabel("y")
    ax.set_title("some random data")
```

Out[3]: Text(0.5,1,'some random data')



```
In [6]: k_mean = KMeans(n_clusters=2)
                                                         k_{mean.fit(X)}
                                                         print(k_mean)
                                                         # plot the centroids
                                                         fig = plt.figure(figsize=(8,8))
                                                        ax = fig.add_subplot(1,1,1)
ax.scatter(X[:, 0], X[:, 1], s = 50, c = 'b', label="data")
                                                        ax. scatter (k\_mean.cluster\_centers\_[0][0], \ k\_mean.cluster\_centers\_[0][1], \ s=200, \ c='g', \ marker='s', \ label="center 1" \ label="center 
                                                        ax.scatter(k\_mean.cluster\_centers\_[1][0], \ k\_mean.cluster\_centers\_[1][1], \ s=200, \ c='r', \ marker='s', \ label="center 2" \ label="center 2"
                                                        ax.grid()
                                                        ax.legend()
                                                         ax.set_xlabel("x")
                                                         ax.set_ylabel("y")
                                                        ax.set_title("some random data")
                                                        KMeans(algorithm='auto', copy_x=True, init='k-means++', max_iter=300,
                                                                                  n_clusters=2, n_init=10, n_jobs=1, precompute_distances='auto',
                                                                                   random_state=None, tol=0.0001, verbose=0)
```

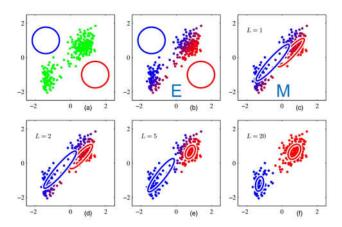
#### Out[6]: Text(0.5,1,'some random data')



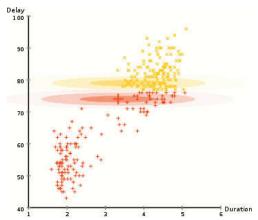
# A

## **Gaussian Mixture Models (GMMs)**

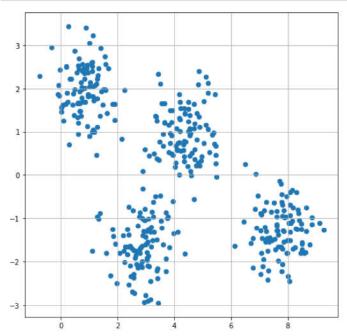
- A Gaussian mixture model is a probabilistic model that assumes all the data points are generated from a mixture of a finite number of Gaussian distributions with unknown parameters.
- One can think of mixture models as generalizing k-means clustering to incorporate information about the covariance structure of the data as well as the centers of the latent Gaussians.
- The parameters of the model are the mean and covariance, which are unknown and will be learned by the EM algorithm.
- Goal: Soft clustering the data under the assumption that it is generated by a mixture of Gaussians.
  - The optimization method is called Expectation Maximization (EM) and will be used to achieve this goal.
- Illustration:



• Animation:

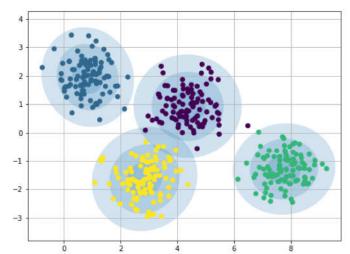


• Example: the following example is taken from <u>The Python Data Science Handbook (https://jakevdp.github.io/PythonDataScienceHandbook/05.12-gaussian-mixtures.html)</u>



```
In [7]: # some helper functions for plotting
         from matplotlib.patches import Ellipse
         def draw_ellipse(position, covariance, ax=None, **kwargs):
              """Draw an ellipse with a given position and covariance"""
             ax = ax or plt.gca()
             # Convert covariance to principal axes
             if covariance.shape == (2, 2):
                  U, s, Vt = np.linalg.svd(covariance)
                 angle = np.degrees(np.arctan2(U[1, 0], U[0, 0]))
width, height = 2 * np.sqrt(s)
                  angle = 0
                  width, height = 2 * np.sqrt(covariance)
             # Draw the Ellipse
             for nsig in range(1, 4):
                  ax.add_patch(Ellipse(position, nsig * width, nsig * height,
                                        angle, **kwargs))
         def plot_gmm(gmm, X, label=True, ax=None):
             ax = ax or plt.gca()
             labels = gmm.fit(X).predict(X)
             if label:
                 ax.scatter(X[:, 0], X[:, 1], c=labels, s=40, cmap='viridis', zorder=2)
             else:
                 ax.scatter(X[:, 0], X[:, 1], s=40, zorder=2)
             ax.axis('equal')
             w_factor = 0.2 / gmm.weights_.max()
             for pos, covar, w in zip(gmm.means_, gmm.covariances_, gmm.weights_):
    draw_ellipse(pos, covar, alpha=w * w_factor)
```

```
In [12]: gmm = GaussianMixture(n_components=4, random_state=42)
    fig = plt.figure(figsize=(8,6))
    ax = fig.add_subplot(1,1,1)
    ax.grid()
    plot_gmm(gmm, X, ax=ax)
```





# **Expectation Maximization (EM)**

- Probabilistic method for soft clustering.
  - The "soft" version of K-means
- Assumes a probabilistic model of clusters that allows computing  $Pr(c_i|x)$  for each cluster  $c_i$  for a given example x.
  - If we had known for each data instance from what distribution it came from, we could have used a parametric estimation.
- We introduce unobservable (latent) variables which indicate source distribution.
- · We run an iterative process:
  - Estimate latent variables from the data and the current estimation of distribution parameters.
  - Use current values of latent variables to refine parameter estimation.



## **Formalization**

• Log likelihood for a mixture model (under the i.i.d assumption):

$$\mathcal{L}(X|\Theta) = \log \prod_{i} Pr(x_i|\Theta) = \sum_{i} \log \sum_{j=1}^{k} Pr(x_i|C_j;\Theta) Pr(C_j;\Theta)$$

- Assume  $\mathit{latent}$  variables z, which when known make the optimization simpler
  - Complete likelihood,  $\mathcal{L}_c(X, Z|\Theta)$ , in terms of x and z
  - **Incomplete** likelihood,  $\mathcal{L}(X|\Theta)$ , in terms of x
- However, z is *latent*, so we can't compute  $\mathcal{L}_c(X, Z|\Theta)$

- But we can compute its **conditional expected value**, given 
$$X$$
 and old  $\theta^t$ : 
$$Q(\Theta;\Theta^t) = \mathbb{E}_Z[\mathcal{L}_c(X,Z|\Theta)|X,\Theta^t)] = \sum_Z Pr(Z|X,\Theta^t) \log Pr(X,Z;\Theta)$$

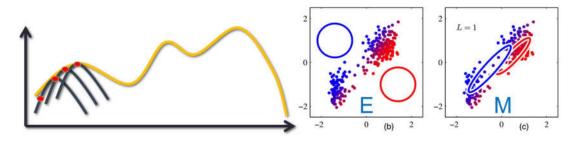
- From a computation viewpoint:
  - The E-Step: computes the posterior probability  $Pr(Z|X,\Theta^t)$  using the *current* estimates (probability point i belongs to model j)
  - The M-Step: updates the parameter estimates to get  $\Theta^{t+1}$  by maximizing  $Q(\Theta;\Theta^t)$
- The EM Algorithm requires an **initial guess**  $\Theta^0$  for the parameters.
- Each iteration of E-step and M-step is guaranteed to increase the log-likelihood of the observed data,  $\log Pr(X|\Theta)$  until a local maximum is reached.



# The Steps

- · Iterate the two steps:
  - ullet E-step: Estimate Z given X and current  $\Theta$ 
    - $\circ \ Q(\Theta|\Theta^t) = \mathbb{E}[\mathcal{L}(X, Z|\Theta)|X, \Theta^t]$
  - M-step: Find new  $\Theta$  given Z,X and old  $\Theta$ 
    - $\bullet \ \Theta^{t+1} = argmax_{\Theta}Q(\Theta; \Theta^t)$
- An increase in  ${\cal Q}$  increases the incomplete likelihood

$$\mathcal{L}(X|\Theta^t) \geq Q(\Theta|\Theta^t)$$





# E Step - EM Receipe

Estimate Z given X and current  $\Theta$ :

$$Pr(z_i = j | x_i, \Theta) = rac{Pr(x_i, z_i = j | \Theta)}{Pr(x_i | \Theta)} = rac{Pr(x_i, z_i = j | \Theta)}{\sum_{j'} Pr(x_i, z_i = j' | \Theta)}$$

· Substitute the probabilities with the desiered distribution.



# M Step (Derive Q) - EM Receipe

$$Q(\Theta|\Theta^t) = \mathbb{E}[\mathcal{L}(X,Z|\Theta)|X,\Theta^t] = \sum_{Z} Pr(Z|X,\Theta^t) \log Pr(X,Z;\Theta) = \sum_{i} \sum_{j=1}^k Pr(z_i=j|x_i,\Theta^t) \log Pr(x_i,z_i=j,\Theta) = \sum_{j=1}^k Pr(z_i=j|x_j,\Theta^t) \log Pr(x_j,z_j=j,\Theta) = \sum_{j=1}^k Pr(z_j=j|x_j,\Theta^t) \log Pr(x_j=j,\Theta^t) = \sum_{j=1}^k Pr(z_j=j|x_j,\Theta^t) = \sum_{j=1}^k P$$

$$\sum_i \sum_{j=1}^k r_{ij} [\log Pr(z_i=j|\Theta) + \log Pr(x_i|z_i=j,\Theta)]$$

- $r_{ij}=Pr(z_i=j|x_i,\Theta)$  (from E step)
   Substitute  $Pr(x_i|z_i=j,\Theta)$  with the desired probability
   Find MLE (differentiate and compare to 0)

### Gaussian Mixture Models (GMMs) as EM



· One Gaussian:

$$\mathcal{N}(x|\mu,\Sigma) = Pr(x|\mu_j,\Sigma_j) = rac{1}{(2\pi)^{rac{d}{2}}|\Sigma_j|^{rac{1}{2}}} e^{-rac{1}{2}(x-\mu_j)^T\Sigma_j^{-1}(x-\mu_j)}$$

Gaussian Mixture:

$$Pr(x) = \sum_{i=1}^k lpha_j \mathcal{N}(x|\mu_j, \Sigma_j)$$

- $\sum_{j=1}^k lpha_j = 1$  Tha parameters of the model are:  $lpha_j, \mu_j, \Sigma_j, orall j \in \{1,\dots,k\}$

$$\mathcal{L}(X|\Theta) = \log \prod_i Pr(x_i|\Theta) = \sum_i \log \sum_{j=1}^k lpha_j \mathcal{N}(x|\mu_j, \Sigma_j)$$

- · No closed form solution and not convex!
- We introduce a latent random variable z
  - $z \in \{0,1\}^k$  a one-hot random variable indicating the source Gaussian the sample belongs to
  - $lacksquare Pr(z_k) = lpha_k$  the probability of that source
  - lacksquare Reminder:  $\sum_{j=1}^k lpha_k = 1$
- · The marginal probability:

$$Pr(x) = \sum_z p(z) p(x|z) = \sum_{j=1}^k lpha_j \mathcal{N}(x|\mu_j, \Sigma_j)$$

#### GMM - E-Step

• The E-step computes the posterior probability of the missing data

$$Pr(z_{i} = j | x_{i}, \Theta) = \frac{Pr(x_{i}, z_{i} = j | \Theta)}{\sum_{j'} Pr(x_{i}, z_{i} = j' | \Theta)} = \frac{\alpha_{j} Pr(x_{i} | \mu_{j}, \Sigma_{j})}{\sum_{j'} \alpha_{j'} Pr(x_{i} | \mu_{j'}, \Sigma_{j'})} = \frac{\alpha_{j} e^{-\frac{1}{2}(x_{i} - \mu_{j})^{T} \Sigma_{j}^{-1}(x_{i} - \mu_{j})}}{\sum_{j'} \alpha_{j'} e^{-\frac{1}{2}(x_{i} - \mu_{j})^{T} \Sigma_{j}^{-1}(x_{i} - \mu_{j})}}$$

• Denote:  $r_{ij} = Pr(z_i = j | x_i, \Theta)$ 

GMM - Calculate  $Q(\Theta; \Theta^t)$ 

$$\begin{split} Q(\Theta|\Theta^t) &= \mathbb{E}[\mathcal{L}(X,Z|\Theta)|X,\Theta^t] = \sum_{Z} Pr(Z|X,\Theta^t) \log Pr(X,Z;\Theta) = \sum_{i} \sum_{j=1}^k Pr(z_i=j|x_i,\Theta^t) \log Pr(x_i,z_i=j,\Theta) = \\ &\sum_{i} \sum_{j=1}^k r_{ij} [\log Pr(z_i=j|\Theta) + \log Pr(x_i|z_i=j,\Theta)] = \\ &\sum_{i} \sum_{j=1}^k r_{ij} [\log \alpha_j + \log Pr(x_i|\mu_j,\Sigma_j)] = \\ &\sum_{i} \sum_{j=1}^k r_{ij} \log \alpha_j - \frac{1}{2} \sum_{j=1}^k \log |\Sigma_j| \sum_{i} r_{i,j} - \frac{1}{2} \sum_{i} \sum_{j=1}^k r_{ij} (x_i - \mu_j)^T \Sigma_j^{-1} (x_i - \mu_j) + Const \end{split}$$

#### GMM - M-Step

- To maximize  $Q(\Theta;\Theta^t)$  with respect to  $\mu_j$  , we set the gradient to zero.
- Reminder:  $\frac{\partial}{\partial s}(x-As)^TW(x-As)=-2A^TW(x-As)$
- · Derive:

$$rac{\partial}{\partial \mu_j}Q(\Theta;\Theta^t) = \sum_{i=1}^n r_{ij}\Sigma_j^{-1}(x_i-\mu_j) = 0 
ightarrow \hat{\mu}_j = rac{\sum_{i=1}^n r_{ij}x_i}{\sum_{i=1}^n r_{ij}}$$

Similarly:

$$\hat{\Sigma}_{j} = rac{\sum_{i=1}^{n} r_{ij} (x_{i} - \mu_{j}) (x_{i} - \mu_{j})^{T}}{\sum_{i=1}^{n} r_{ij}}$$

- To maximize  $Q(\Theta; \Theta^t)$  with respect to  $\alpha_i$ :
  - Use Lagrange multiplier:

$$egin{array}{l} \bullet & \max Q(\Theta;\Theta^t) ext{ s.t. } \sum_j lpha_j = 1 & \Longleftrightarrow \ \bullet & \mathcal{L} = Q(\Theta;\Theta^t) + \lambda (1 - \sum_i lpha_i) \end{array}$$

ullet Substituting  $\lambda$  back in the  $\emph{Lagrangian}$  derivative:

$$rac{\partial \mathcal{L}}{\partial lpha_j} = \sum_i rac{r_{ij}}{lpha_j} - \lambda = 0 
ightarrow \hat{lpha}_j = rac{\sum_{i=1}^n r_{ij}}{n}$$

• To sum up:

$$\hat{\mu}_{j} = rac{\sum_{i=1}^{n} r_{ij} x_{i}}{\sum_{i=1}^{n} r_{ij}} \ \hat{\Sigma}_{j} = rac{\sum_{i=1}^{n} r_{ij} (x_{i} - \mu_{j}) (x_{i} - \mu_{j})^{T}}{\sum_{i=1}^{n} r_{ij}} \ \hat{lpha}_{j} = rac{\sum_{i=1}^{n} r_{ij}}{n}$$

# Bernoulli Mixture Models (BMMs) as EM

- We have k coins such that:
  - The probability of observing *heads* with the  $j^{th}$  coin is  $p_j$ .
  - We do not observe which coin was used.
  - We only observe  $x_i \in \{0,1\}$ , which records whether we see a *heads* or *tails*.
- Let  $z_i \in \{1,\dots,k\}$  be the **missing information** of which coin was used on each flip (in other words, the *source* like in the GMM case).
  - The probability of using the  $j^{th}$  coin is  $Pr(z_i=j)=lpha_j$  (which is a *parameter*)
- The complete data is given by (X, Z)
  - Using the **law of total probability**, the (marginal) probability of the observed data X:

$$Pr(X) = \sum_{j} Pr(X|Z=j) Pr(Z=j)$$

$$Pr(X) = \sum_{j} Pr(X|Z=j) Pr(Z=j)$$

$$\blacksquare \text{ Thus, the } \textit{likelihood} \text{ of the full data set (incomplete likelihood) is:}$$

$$\pounds(X|\Theta) = \prod_{i} \sum_{j} Pr(x_{i}|z_{i}=j) Pr(z_{i}=j) = \prod_{i} \sum_{j} \alpha_{j} p_{j}^{x_{i}} (1-p_{j})^{1-x_{i}}$$

$$\bullet \Theta = (\alpha, p)$$

#### BMM - E-Step

• The E-step computes the posterior probability of the missing data

$$Pr(z_i = j | x_i, \Theta) = rac{Pr(x_i, z_i = j | \Theta)}{\sum_{j'} Pr(x_i, z_i = j' | \Theta)} = rac{lpha_j Pr(x_i | p_j)}{\sum_{j'} lpha_j Pr(x_i | p_{j'})} = rac{lpha_j p_j^{x_i} (1 - p_j)^{1 - x_i}}{\sum_{j'} lpha_j p_{j'}^{x_i} (1 - p_j)^{1 - x_i}}$$

• Denote:  $r_{ij} = Pr(z_i = j | x_i, \Theta)$ 

 $\Theta = (\alpha, p)$ 

#### BMM - Calculate $Q(\Theta; \Theta^t)$

$$egin{aligned} Q(\Theta|\Theta^t) &= \mathbb{E}[\mathcal{L}(X,Z|\Theta)|X,\Theta^t] = \sum_{Z} Pr(Z|X,\Theta^t) \log Pr(X,Z;\Theta) = \sum_{i} \sum_{j=1}^k Pr(z_i=j|x_i,\Theta^t) \log Pr(x_i,z_i=j,\Theta) = \\ &\sum_{i} \sum_{j=1}^k r_{ij} [\log Pr(z_i=j|\Theta) + \log Pr(x_i|z_i=j,\Theta)] = \\ &\sum_{i} \sum_{j=1}^k r_{ij} [\log lpha_j + \log Pr(x_i|p_j)] = \\ &\sum_{i} \sum_{j=1}^k r_{ij} \log lpha_j + \sum_{i} \sum_{j=1}^k r_{ij} \log \left(p_j^{x_i}(1-p_j)^{1-x_i}\right) = \\ &\sum_{i} \sum_{j=1}^k r_{ij} \log lpha_j + \sum_{i} \sum_{j=1}^k r_{ij} x_i \log(p_j) + r_{ij}(1-x_i) \log(1-p_j) \end{aligned}$$

#### BMM - M-Step

- To maximize  $Q(\Theta; \Theta^t)$  with respect to  $p_i$ , we set the gradient to zero.

$$rac{\partial}{\partial p_j}Q(\Theta;\Theta^t)=\sum_{i=1}^n r_{ij}ig(rac{x_i}{p_j}-rac{1-x_i}{1-p_j}ig)=0
ightarrow\hat{p}_j=rac{\sum_{i=1}^n r_{ij}x_i}{\sum_{i=1}^n r_{ij}}$$

- To maximize  $Q(\Theta; \Theta^t)$  with respect to  $\alpha_i$ :
  - Use Lagrange multiplier:

$$\begin{array}{l} \bullet \text{ ose Lagrange Hulliphen.} \\ \bullet \max Q(\Theta;\Theta^t) \text{ s.t. } \sum_j \alpha_j = 1 \iff \\ \bullet \ \mathcal{L} = Q(\Theta;\Theta^t) + \lambda(1-\sum_j \alpha_j) \\ \bullet \ \frac{\partial \mathcal{L}}{\partial \alpha_j} = \sum_i \frac{r_{ij}}{\alpha_j} - \lambda = 0 \end{array}$$

$$\bullet$$
  $\frac{\partial \mathcal{L}}{\partial x_i} = \sum_{i} \frac{r_{ij}}{\partial x_i} - \lambda = 0$ 

Find an expression for 
$$\lambda$$
 by summing all partial derivatives of  $\alpha_j$ : 
$$\sum_i r_{ij}^{(t)} = \lambda \alpha_j \rightarrow \sum_j \sum_i r_{ij}^{(t)} = \lambda \sum_j \alpha_j \rightarrow \lambda = n$$
 
$$\circ \ \sum_j \sum_i r_{ij}^{(t)} = \sum_j \sum_i Pr(z_i = j | x_i, \Theta^t) = \sum_i \frac{\sum_j Pr(x_i, z_i = j | \Theta)}{\sum_j Pr(x_i, z_i = j | \Theta)} = \sum_{i=1}^n 1 = n$$

ullet Substituting  $\lambda$  back in the Lagrangian derivative

$$rac{\partial \mathcal{L}}{\partial lpha_{i}} = \sum_{i} rac{r_{ij}}{lpha_{i}} - \lambda = 0 
ightarrow \hat{lpha}_{j} = rac{\sum_{i=1}^{n} r_{ij}}{n}$$

· To sum up:

$$\hat{p}_{j} = rac{\sum_{i=1}^{n} r_{ij} x_{i}}{\sum_{i=1}^{n} r_{ij}}$$

$$\hat{lpha}_j = rac{\sum_{i=1}^n r_{ij}}{n}$$

- If all  $r_{ij} = \{0,1\}$ , that is, deterministic, then:
  - The component labels,  $z_i$ , are **known**
  - The above update equations reduce to the standard formulas for binomial distribution.



# Relation to K-Means: K-Means as "Hard GMM"

• Recall the E-Step for GMM:

$$Pr(z_i = j | x_i, \Theta) = \frac{Pr(x_i, z_i = j | \Theta)}{\sum_{j'} Pr(x_i, z_i = j' | \Theta)} = \frac{\alpha_j Pr(x_i | \mu_j, \Sigma_j)}{\sum_{j'} \alpha_{j'} Pr(x_i | \mu_{j'}, \Sigma_{j'})} = \frac{\alpha_j e^{-\frac{1}{2}(x_i - \mu_j)^T \Sigma_j^{-1}(x_i - \mu_j)}}{\sum_{j'} \alpha_{j'} e^{-\frac{1}{2}(x_i - \mu_j)^T \Sigma_j^{-1}(x_i - \mu_j)}}$$

- Let's assume all the Gaussians have the same  $\Sigma = \epsilon I$  :

$$Pr(z_i=j|x_i,\Theta) = \frac{Pr(x_i,z_i=j|\Theta)}{\sum_{j'} Pr(x_i,z_i=j'|\Theta)} = \frac{\alpha_j e^{-\frac{1}{2\epsilon}||(x_i-\mu_j)||_2^2)}}{\sum_{j'} \alpha_j e^{-\frac{1}{2\epsilon}||(x_i-\mu_j)||_2^2}}$$
 • At the limit  $\epsilon \to 0$ :  $Pr(z_i=j|x_i,\Theta) = 1$  for  $j=argmin\{x_i-\mu_j\}$  and  $Pr(z_i=j|x_i,\Theta) = 0$  for all others. • Thus:

$$\left\{egin{aligned} r_{ij} = 1 & & ext{if } j = argmin_j ||x_i - \mu_j||_2^2 \ r_{ij} = 0 & & ext{else} \end{aligned}
ight.$$

• The GMM equations are now identical to the K-Means' eqautions:

$$\hat{\mu}_j = rac{\sum_{i=1}^n r_{ij} x_i}{\sum_{i=1}^n r_{ij}}$$

$$\hat{lpha}_j = rac{\sum_{i=1}^n r_{ij}}{n}$$

• The  $\alpha$ s are not really required.



# **Credits**

- Icons from Icon8.com (https://icons8.com/) https://icons8.com (https://icons8.com)
- Datasets from Kaggle (https://www.kaggle.com/) https://www.kaggle.com/ (https://www.kaggle.com/)
- Examples and code snippets were taken from "Hands-On Machine Learning with Scikit-Learn and TensorFlow" (http://shop.oreilly.com/product/0636920052289.do)