



Clustering

Introduction to Data Science University of Colorado Boulder SLIDES ADAPTED FROM LAUREN HANNAH K-means associates data with cluster centers.

What if we actually modeled the data?

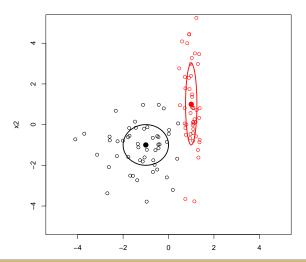
- real-valued data
- observation \mathbf{x}_i in cluster c_i
- have K clusters
- model each cluster with a Gaussian distribution

$$\mathbf{x}_i | \mathbf{c}_i = \mathbf{k} \sim N(\mu_k, \Sigma_k)$$

• μ_k is mean vector, Σ_k is covariance matrix

Mixture Models

Gaussian mixture model (K = 2):



Why mixture models?

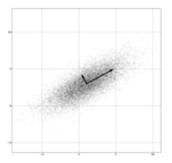
- more flexible: can account for clusters with different shapes
- have data model (will be useful for choosing K)
- less sensitive to data scaling

Multivariate Gaussian

Multivariate Gaussian distribution for $\mathbf{x} \in \mathbf{R}^d$:

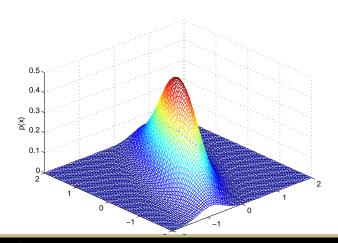
$$p(\mathbf{x}|\mu,\Sigma) = (2\pi)^{-\frac{d}{2}} |\Sigma|^{-\frac{1}{2}} e^{-\frac{1}{2}(\mathbf{x}-\mu)^{T}\Sigma^{-1}(\mathbf{x}-\mu)}$$

- µ is vector of means
- Σ is covariance matrix

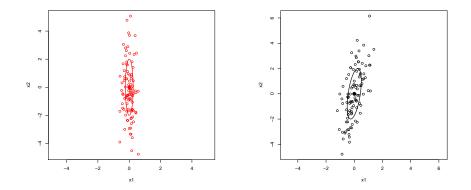


Multivariate Gaussian

pdf when
$$\mu = [0, 0]$$
 and $\Sigma = \begin{bmatrix} 0.9 & 0.4 \\ 0.4 & 0.3 \end{bmatrix}$:



Multivariate Gaussian



Mixture model:

- observation x_i in cluster c_i with K clusters
- model each cluster with a Gaussian distribution

$$\mathbf{x}_i | c_i = k \sim N(\mu_k, \Sigma_k)$$

How do we find c_1, \ldots, c_n (clusters) and $(\mu_1, \Sigma_1), \ldots, (\mu_K, \Sigma_K)$ (cluster centers)?

First, let's simplify the model:

· covariance matrices have only diagonal elements,

$$\Sigma = \begin{bmatrix} \sigma_1^2 & 0 & \dots & 0 \\ 0 & \sigma_2^2 & \dots & 0 \\ \dots & \dots & \dots & 0 \\ 0 & 0 & 0 & \sigma_K^2 \end{bmatrix}$$

• set
$$\sigma_1^2 = \cdots = \sigma_K^2$$
, suppose known

Next, use a method similar to K-means:

- start with random cluster centers
- associate observations to clusters by (log-)likelihood,

$$\ell(\mathbf{x}_{i} | c_{i} = k) = -\frac{d}{2} \log(2\pi) - \frac{1}{2} \log\left(\prod_{j=1}^{d} \sigma_{k,j}^{2}\right) - \frac{1}{2} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2} / \sigma_{k,j}^{2}$$
$$\propto -d \log(\sigma_{k}) - \frac{1}{2\sigma_{k}^{2}} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}$$
$$\propto -\sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}$$

• refit centers μ_1, \ldots, μ_K given clusters by

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i=k} x_{i,j}$$

recluster observations...

clustering with K-means

minimize distance

$$d(\mathbf{x}_{i}, \mu_{k}) = \sqrt{\sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}}$$

clustering with GMM maximize likelihood

$$\ell(\mathbf{x}_i | c_i = k) \propto -\sum_{j=1}^{a} (x_{i,j} - \mu_{k,j})^2$$

update means with K-means

use average

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i = k} x_{i,j}$$

update means with GMM use average

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i=k} x_{i,j}$$

OK, now what if

$$\Sigma = \left[egin{array}{ccccc} \sigma_1^2 & 0 & \dots & 0 \ 0 & \sigma_2^2 & \dots & 0 \ \dots & \dots & \dots & 0 \ 0 & 0 & 0 & \sigma_K^2 \end{array}
ight]$$

and $\sigma_1^2, \ldots, \sigma_K^2$ can take different values?

- use same algorithm
- update μ_k and σ_k^2 with maximum likelihood estimator,

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i = k} x_{i,j}$$

$$\sigma_{k,j}^2 = \frac{1}{n_k} \sum_{c_i = k} (x_{i,j} - \mu_{k,j})^2$$

Data:

| <i>x</i> ₁ | <i>x</i> ₂ | ۰ |
|-----------------------|-----------------------|-------------|
| -3.7 | -0.4 | ю — |
| 0.4 | 0.1 | 4 - |
| 0.4 | -1.7 | ю — |
| -0.4 | -1.0 | o |
| -1.3 | -1.7 | - ~ × |
| 1.0 | 3.3 | |
| 1.2 | 5.2 | ° – ° |
| 1.3 | 0.3 | ° _ ° |
| 1.1 | -0.8 | · · · · · · |
| 0.5 | 2.8 | |
| | | x1 |

- pick centers and variances, $\mu_1 = [-1, -1]$, $\sigma_1^2 = [1, 1]$, $\mu_1 = [1, 1]$, $\sigma_1^2 = [1, 1]$
- compute (proportional) log likelihoods,

$$\ell(\mathbf{x}_{i} | c_{i} = k) = -\sum_{j=1}^{d} \log(\sigma_{j}) - \frac{1}{2} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2} / \sigma_{k,j}^{2}$$

| <i>x</i> ₁ | <i>x</i> ₂ | <i>k</i> = 1 | k = 2 |
|-----------------------|-----------------------|--------------|-------|
| -3.7 | -0.4 | -3.8 | -12.1 |
| 0.4 | 0.1 | -1.6 | -0.6 |
| 0.4 | -1.7 | -1.2 | -3.8 |
| -0.4 | -1.0 | -0.2 | -3.0 |
| -1.3 | -1.7 | -0.3 | -6.3 |
| 1.0 | 3.3 | -11.2 | -2.6 |
| 1.2 | 5.2 | -22.0 | -9.0 |
| 1.3 | 0.3 | -3.6 | -0.3 |
| 1.1 | -0.8 | -2.2 | -1.6 |
| 0.5 | 2.8 | - 8.2 | -1.7 |

fit new means and variances:

$$\mu_1 = [-1.3, -1.2]$$

$$\sigma_1^2 = [3.1, 0.4]$$

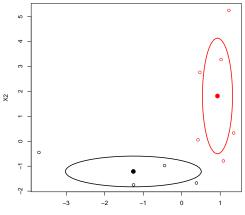
$$\mu_2 = [0.9, 1.8]$$

$$\sigma_2^2 = [0.2, 5.4]$$

compute new distances...

| <i>x</i> ₁ | <i>x</i> ₂ | <i>k</i> = 1 | k = 2 |
|-----------------------|-----------------------|--------------|-------|
| -3.7 | -0.4 | -1.8 | -70.8 |
| 0.4 | 0.1 | -2.7 | -1.0 |
| 0.4 | -1.7 | -0.8 | -2.0 |
| -0.4 | -1.0 | -0.3 | -6.8 |
| -1.3 | -1.7 | -0.5 | -16.6 |
| 1.0 | 3.3 | -27.4 | -0.1 |
| 1.2 | 5.2 | -55.9 | -1.3 |
| 1.3 | 0.3 | -4.3 | -0.7 |
| 1.1 | -0.8 | -1.2 | -0.6 |
| 0.5 | 2.8 | -21.3 | -0.7 |

No change, so clusters are final



k-means is fast and simple, but ...

- What if your data are discrete?
- What if each data point has more than one cluster? (digits vs. documents)
- What if you don't know the number of clusters?

- Clustering helps discover patterns
- *k*-means is a simple approach
- Gaussian mixture models more probabilistic foundation