# Tools & Models for Data Science Mixture Models

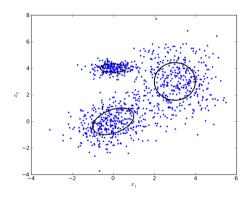
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#### Mixture Model Introduction

- At highest level:
  - Have a set of data
  - And a set of random variables
  - Don't know which one produced which point
- This is a mixture model!
- In one line:
  - MM: "hierarchical," stochastic, latent variable model

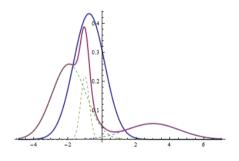


## Why Use Them?

- Sometimes, we want to segment the data
  - Observe a set of test scores
  - Want 3 types of students: good, average, bad
  - Associate each with a different Normal distribution
- Sometimes, we just want a very flexible model
  - A mixture can give a complicated, multi-modal distribution
  - We can construct these from simple distributions
  - In reality, not much data is purely normally distributed

## GMM Example

- Blue curve is what you get if you fit a single normal distribution to the purple data
- Note the shift in the mean
- The dashed curves are the mixture distributions
- Note there are actually 3 normal distributions in this plot



#### Mixture Model Introduction

- Choose the parameters for each normal distribution
- And a weighting for each mixture
- Use these to produce a set of data  $x_1, x_2, ..., x_n$ 
  - And a set of hidden (latent) indicators  $c_1, c_2, ..., c_n$  specifying which mixture was selected
- MM begins with a distribution function *f* 
  - Common *f*: Gaussian, Multinomial, Gamma, etc.
  - We have k sets of parameters for f:  $\theta_1, \theta_2, ..., \theta_k$
  - lacktriangle And a probability vector  $\pi$  that tells us how important each mixture component is
  - Note: we can have a mixture model where each component is from a different distribution
  - This is uncommon, though

#### Mixture Model Introduction

■ Pseudo-code to produce *n* observations is:

```
for i = 1 to n do:

c_i \sim \operatorname{Categorical}(\pi)

x_i \sim f(\theta_{c_i})
```

#### Mixture Model PDF

■ In general, PDF is:

$$P(x_1, x_2, ..., x_n) = \prod_{i}^{n} \left( \sum_{j}^{K} \pi_{j} f(x_i | \theta_j) \right)$$

- $\blacksquare$  Where *n* is the number of data points
- Where *K* is the number of mixtures
- Where *f* is the PDF for distribution *k*
- ? Why?

#### Mixture Model PDF

■ In general, PDF is:

$$P(x_1,x_2,...,x_n) = \prod_{i}^{n} \left( \sum_{j}^{K} \pi_{j} f(x_i | \theta_j) \right)$$

- Why?
  - Likelihood of choosing component j, then j producing x is:

$$\pi_j f(x|\theta_j)$$

lacksquare Since n data points are independent, we have a product over the likelihoods

#### Mixture Model PDF

For each data point, the PDF is:

$$\sum_{j}^{k} \pi_{j} f(x_{i} | \theta_{j})$$

- We can represent the choice, *c* as a 1-of-k valued vector, where one dimension is 1 and the rest are all 0
- The dimension is set to 1 if that mixture model generated the data point
- and set to 0 otherwise
- It follows that  $p(c_j = 1) = \pi_j$
- $\blacksquare$  and that  $\sum_{i=1}^{k} \pi_{i} = 1$
- To get the marginal distribution of *x* we sum the joint distribution over all possible states of *c*

## Learning

- Would be easy if we knew  $c_1, c_2, ..., c_n$
- Where  $c_j$  is the indicator of which mixture distribution produced data point j
  - In that case, we'd have *k* different MLE problems
  - That is, to learn, partition the data points according to *c* values
  - Then perform an MLE separately on each
  - For standard distributions, this is easy
  - Example, for 1-D Normal
    - MLE for  $\mu_j$  is mean of points with  $c_i = j$
    - MLE for  $\sigma_j$  is std. dev. of points with  $c_i = j$
- But there are complications in doing this
- $\blacksquare$  And we don't know the  $c_i$  values anyway!

## Learning

- But we don't!
- Standard methods:
  - MCMC (sample  $c_1, c_2, ..., c_n$ ): Stochastic, Bayesian approach, not covered in this course
  - EM

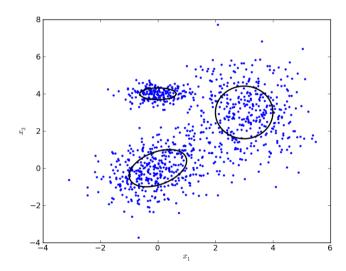
#### EM for Mixture Models...

■ The Patriarch: The Gaussian Mixture Model (GMM)

$$P(x_1, x_2, ..., x_n) = \prod_i \left( \sum_j \pi_j \text{Normal}(x_i | \mu_j, \sigma_j^2) \right)$$

- Make non-stupid guesses for the means, variances, and the mixing parameters
- E-step: Use the current parameter values to asign each data point to the most likely mixture component
- 3 M-step: Use the expected values to update the mixture parameters

## Example GMM



#### Mixture of Dirichlets

- Imagine that I have a large corpus of text documents
- And I want to understand the various types of documents present
- Basically, I want to cluster the documents
- We might view the TF (term frequency) vector as being produced as a sample from a Dirichlet distribution
- ? What's a Dirichlet distribution?

#### **Dirichlet Distribution**

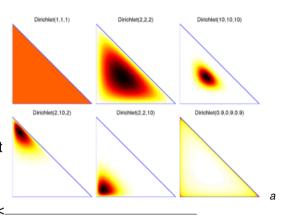
- Generalization of Beta to many dimensions
- Produces vectors on the "simplex"
  - That is, vectors whose entries sum to 1
  - Hence, used as a distribution to produce vectors of probabilities
- Takes params  $\langle v_1, v_2, ..., v_m \rangle$ 
  - Produces m values that sum to 1
  - "looks" like a probability vector
  - To generate, sample from m Gammas, each with shape  $v_j$
  - ith value is ith Gamma's fractional contribution to total

#### **Dirichlet Distribution**

- Expected value of *i*th is  $\frac{v_j}{\sum_{j'} v_{j'}}$ 
  - Then what's the difference between parameter vector  $\langle 1,2,3 \rangle$  and  $\langle 10,20,30 \rangle$ ?
- Both result in Dirichlets with same mean
  - But smaller parameters allow more variance
  - So the former has much more variance... resulting Dirichlet could produce  $\langle 0.2, 0.1, 0.7 \rangle$
  - Latter will always produce something like ⟨0.1666, 0.3333, 0.5⟩
  - If we normalize the parameter vectors (divide by their sum) we get  $6 \times \langle 0.166, 0.333, 0.5 \rangle$  and  $60 \times \langle 0.166, 0.333, 0.5 \rangle$
  - The higher the normalization constant is, the closer the samples will be to the mean

#### **Dirichlet Distribution**

- $1 \langle 1, 1, 1 \rangle$  Everything is equally likely
- 2  $\langle 2,2,2 \rangle$  As the parameters grow, on expectation, the samples are more concentrated
- 3  $\langle 10, 10, 10 \rangle$  As the numbers continue to get bigger, the samples are more likely to be  $\frac{1}{3}, \frac{1}{3}, \frac{1}{3}$
- 4  $\langle 2,10,2\rangle$  If a parameter is larger, that dimension get more weight
- 5  $\langle 2, 2, 10 \rangle$
- (0.9,0.9,0.9) When parameters are <...</li>
   1, the samples go to the outside (more extreme differences)



 ${\it a}_{\rm www.cs.cmu.edu/\sim epxing/Class/10701-08s/recitation/dirichlet.pdf}$ 

## Mixture of Dirichlets Application

- Imagine that I have a large corpus of text documents
- And I want to understand the various types of documents present
- Basically, I want to cluster the documents
- We might view the TF (term frequency) vector as being produced as a sample from a Dirichlet distribution
- TF vector is basically a vector of probabilities

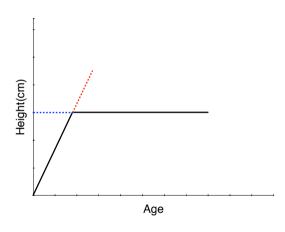
## Mixture of Dirichlets Example

■ If param vector for *j*th Dirichlet is  $\alpha_j$ , then PDF is:

$$P(x_1, x_2, ..., x_n) = \prod_i \left( \sum_j \pi_j \text{Dirichlet}(x_i | \alpha_j) \right)$$

- Take all the papers Chris has written, all the papers Luay has written
- Dictionary has words: 〈 biology, databases, phylogenetics, statistics 〉
- Might get  $\alpha_{\text{Chris}} = \langle .5, 12, 1.2, 8 \rangle$
- Means Chris' mean probability vector is  $\langle \frac{.5}{21.7}, \frac{12}{21.7}, \frac{1,2}{21.7} \frac{8}{21.7} \rangle = \langle 0.02, 0.55, 0.055, 0.37 \rangle$
- For most docs, Chris is most likely to write the word "databases"
- Might get  $\alpha_{\text{Luay}} = \langle 11, 2, 18, 3.2 \rangle$
- His mean probability vector is  $\langle 0.32, 0.06, 0.53, 0.09 \rangle$
- Means for most docs, he's most likely to write the word "phylogenetics"
- So, we have learned how likely each topic is to appear in each author's publications

- Used for regression/classification
- Uses a combination of simpler models to form a better model
- Each simpler model covers a range of input values
- We have a switch that determines which model is in place



- In "classical" GLMs
- Use dot product  $x \cdot r = \sum_{j} x_j \times r_j$  to get "natural" parameter for error distribution
- Ex: Normal (least squares) regression:

$$f(y|x) = \text{Normal}(y|x \cdot r, \sigma^2)$$

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- Now, let's have a mixture of these
  - Instead of r, we have  $r_1, r_2, ..., r_k$
  - Instead of  $\sigma^2$ , we have  $\sigma_1^2, \sigma_2^2, ..., \sigma_k^2$

- Mixture proportions computed by looking at x
  - **Each** component has a gating vector  $\eta$
  - We use a "softmax gating network" to get  $\pi$  for a given x... that is:

$$\pi_j = \frac{\exp(x \cdot \eta_j)}{\sum_{j'} \exp(x \cdot \eta_{j'})}$$

■ In other words, we normalize the mixture selections to sum to 1

■ If param vector for *j*th Dirichlet is  $\alpha_i$ , then PDF is:

$$P(x_1, x_2, ..., x_n) = \prod_i \left( \sum_j \pi_j \text{Dirichlet}(x_i | \alpha_j) \right)$$

So PDF is:

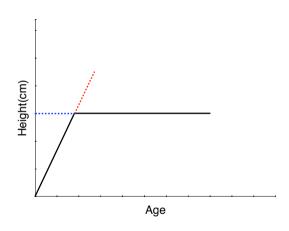
$$P(x_1, x_2, ..., x_n) = \prod_i \left( \sum_j \frac{\exp(x_i \cdot \eta_j)}{\sum_j' \exp(x_i \cdot \eta_{j'})} \operatorname{Normal}(x_i | x_i \cdot r_j, \sigma_j^2) \right)$$

## Mixture of Experts: What's It Good For?

- Allows more flexible regression/classification models
- Example: we want to predict height *y* in cm using age *x* 
  - We know people grow as children, then stop somewhere between age 16 and 21
  - Model as a mixture of two Normal regression models
  - Allow bias by making input 2-D ⟨age, 1⟩
  - Then second regression coef is the bias
  - $r_1 = \langle 6.5, 50 \rangle$  (person starts off at 50cm at birth, grows 6.5cm per year)
  - $ightharpoonup r_2 = \langle 0, 175 \rangle$  (person hits around 175cm and stops growing)
- Then gating function
  - $\eta_0 = \langle 0, 8 \rangle$ ,  $\eta_1 = \langle 1, -8 \rangle$ ... you can do the math...
  - At age 10, prob of first Normal is 0.9995
  - At age 14, prob of first Normal reduces to 0.88
  - At age 16, prob is 50/50
  - At age 18, down to 0.12
  - Down to 0.007 at age 21... means almost everyone has stopped growing

## Mixture of Experts Motivation

- In reality, there are really 2 or 3 linear relationships
- Typically, we know what part of the line you are on



## Mixture of Experts Summary

- Used for regression/classification
  - In "classical" GLMs
  - Use dot product  $x \cdot r = \sum_i x_i \times r_i$  to get "natural" parameter for error distribution
  - Ex: Normal (least squares) regression:

$$f(y|x) = \text{Normal}(y|x \cdot r, \sigma^2)$$

- Now, let's have a mixture of these
  - Instead of r, we have  $r_1, r_2, ..., r_k$
  - Instead of  $\sigma^2$ , we have  $\sigma_1^2, \sigma_2^2, ..., \sigma_k^2$
- And then we use a "softmax gating network" to get  $\pi$ ... that is:

$$\pi_j = \frac{\exp(x \cdot \eta_j)}{\sum_{j'} \exp(x \cdot \eta_{j'})}$$

## Mixture of Experts Summary

So PDF is:

$$P(x_1, x_2, ..., x_n) = \prod_{i} \left( \sum_{j} \frac{\exp(x_i \cdot \eta_j)}{\sum_{j}' \exp(x_i \cdot \eta_{j'})} \operatorname{Normal}(x_i | x_i \cdot r_j, \sigma_j^2) \right)$$

- $\blacksquare$   $\eta_i$  is the set of gating coefficients for mixture j
- The denominator normalizes the values, so we have probabilities
- Basically, we are using a softmax function to decide the mixture
- $\blacksquare$   $\pi$  was a vector before
- Now,  $\pi$  is a vector whose value is dependent on the vector of xs
- If we have a large dot product value it's highly likely that the *y* value (output) will be generated by the *j*th component

■ Smaller ages are to the left, larger are to the right

#### General Question: How to Choose Mixture Size

- In last example...
  - With four mixture components, could have two growth trends (women and men)
  - So probably, four is better than two here
- How to choose in general case?
  - Guess! Then see if it is useful
- Not good enough? Are some alternatives
  - Information theoretic methods (choose model that is able to encode data most efficiently)
  - Bayesian methods (put a prior on the number of components... "infinite mixture models")

#### Questions?

- What do we know now that we didn't know before?
  - We understand that data can be generated by /represented as a mixture of different distributions
  - We know about a Dirichlet distribution that can generate probability vectors
- How can we use what we learned today?
  - We can better model complex data by using mixture models
  - We can combine simpler models into more complex models to get better predictions