Soft Clustering

- So far we have assumed hard clustering:
 - Data point is deterministically assigned to one and only one cluster
- In reality clusters may overlap
- Soft-clustering:
 - Data points are assigned to clusters with probabilities
- To obtain probabilities, we must have some probabilistic models
 - Sometimes referred to as model-based clustering
 - One of the most commonly used model is Gaussian

Side track: Gaussian Bayes Classifier

- We have k classes in our data
- Each class contains data generated from a particular Gaussian distribution
- Data is generated by
 - first randomly select one of the classes according to class prior
 - draw random samples from the Gaussian distribution of that particular class

$$P(\mathbf{x}, y) = P(\mathbf{x} | y)P(y)$$

$$P(\mathbf{x} | y = i) = \frac{1}{(2\pi)^{d/2} |\Sigma_i|^{1/2}} e^{-\frac{1}{2}(\mathbf{x} - \mu_i)^T \sum_i^{-1} (\mathbf{x} - \mu_i)}$$

Supervised vs Unsupervised Learning

- Now assume we know our data is generated in this way
- And we know the labels (y's), we can estimate the mean and covariance matrix of each class using MLE
- What if the labels are hidden?
- How can we learn the correct model from the incomplete data?

Gaussian Mixture Model (GMM)

$$P(\mathbf{x}) = \sum_{i=1}^{k} P(\mathbf{x}, y = i)$$

$$= \sum_{i=1}^{k} P(\mathbf{x} | y = i) P(y = i)$$

$$= \sum_{i=1}^{k} \alpha_{i} P(\mathbf{x} | \theta_{i})$$

$$\theta_{i} = \{\mu_{i}, \Sigma_{i}\}$$

 $\underline{\alpha_i} = P(y=i)$: the class prior Called the mixing parameter

Goal of unsupervised learning:

- Given a set of x's, estimate $\{\alpha_1, \dots, \alpha_k, \theta_1, \dots, \theta_k\}$
- Once the model is identified, we can compute p(y=i|x) for $i=1, \dots, k$, which gives the soft-clustering we want

Maximum Marginal Likelihood

$$\arg \max_{\theta} \prod_{j} P(\mathbf{x}^{j}) = \arg \max_{\theta} \prod_{j=1}^{k} \sum_{i=1}^{k} P(\mathbf{x}^{j}, y^{j} = i)$$

$$= \arg \max_{\theta} \sum_{j=1}^{n} \log \sum_{i=1}^{k} P(\mathbf{x}^{j}, y^{j} = i)$$

log sum is nasty to optimize!

Expectation Maximization (EM)

- A highly used approach for dealing with in-complete data
- It is an iterative algorithm that starts with some initial guesses of the model parameters
- Iteratively performs two main steps:
 - Expectation (E-step): given current model parameters, compute the expectation for the hidden (missing) data
 - Maximization (M-step): re-estimate the parameters assuming that the expected values computed in the E-step are the true values

E-Step for GMM

- If we know $\theta = \{\alpha_1, \dots, \alpha_k, \theta_1, \dots, \theta_k\}$
- We can easily compute the probability that point x^j belong to class y=i

$$p(y = i | \mathbf{x}^{j}; \theta^{(t)}) = \frac{p(\mathbf{x}^{j} | \theta_{i}^{(t)}) \cdot \alpha_{i}^{(t)}}{p(\mathbf{x}^{j})} \\ \propto \alpha_{i}^{(t)} \cdot \frac{1}{(2\pi)^{d/2} |\Sigma_{i}^{(t)}|^{1/2}} \cdot e^{-\frac{1}{2}(x - \mu_{i}^{(t)})^{T} \Sigma_{i}^{(t)^{-1}} (x - \mu_{i}^{(t)})}$$

if
$$p(y=1|x, \theta) \propto 0.3$$

 $p(y=2|x, \theta) \propto 0.2$
then $p(y=1|x, \theta) = 0.6$
 $p(y=2|x, \theta) = 0.4$

Same as classification!

M-Step

• If we know the probability of point x^j belongs to y=i, we can then re-estimate θ

$$\alpha_{i} = \frac{\sum_{j} P(y^{j} = i | x^{j})}{n}$$

$$\alpha_{i} = \frac{\sum_{j} P(y^{j} = i | x^{j}) \cdot \mathbf{x}^{j}}{\sum_{i} P(y^{j} = i | x^{j})}$$

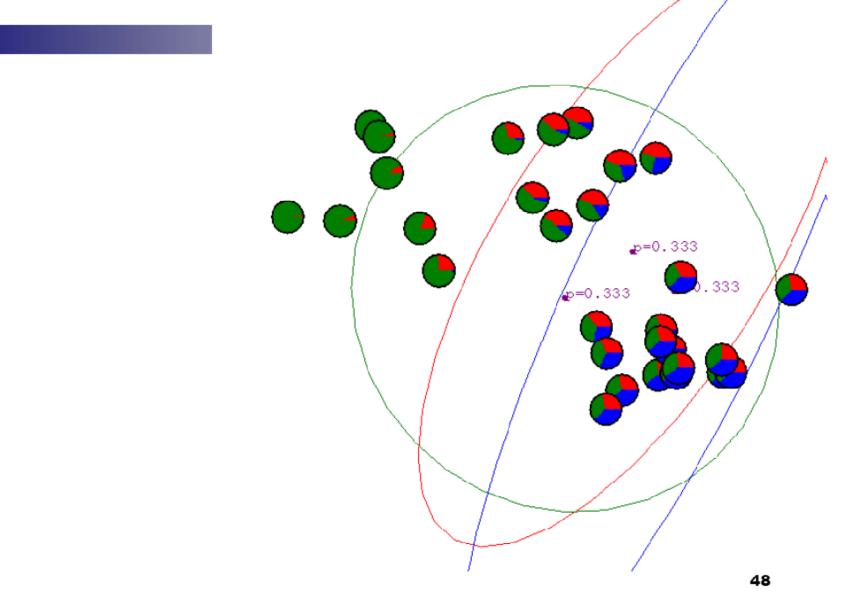
$$\hat{\Sigma}_{i} = \frac{1}{n_{i}} \sum_{y^{j} = i} (\mathbf{x}^{j} - \hat{\mu}_{i}) \cdot (\mathbf{x}^{j} - \hat{\mu}_{i})^{T}$$

$$\sum_{i} P(y^{j} = i | x^{j}) \cdot (\mathbf{x}^{j} - \mu_{i}^{(t+1)}) \cdot (\mathbf{x}^{j} - \mu_{i}^{(t+1)})^{T}$$

$$\sum_{i} P(y^{j} = i | x^{j})$$

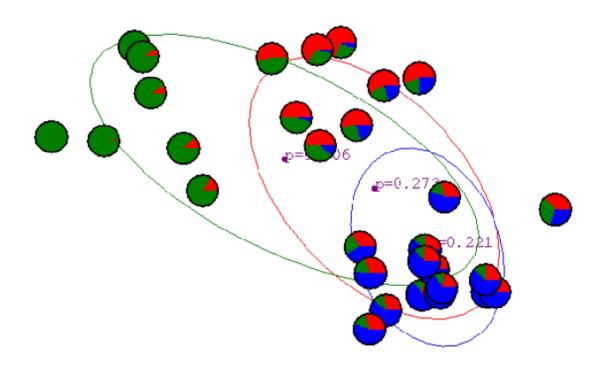
Imagine k copies of each x^{j} , each with weight $P(y^{j}=i|x^{j})$:

Gaussian Mixture Example: Start



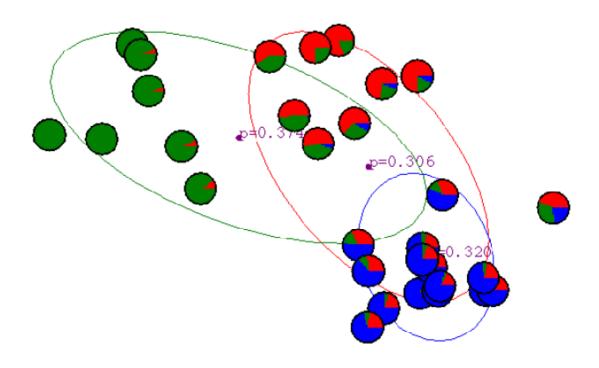
After first iteration





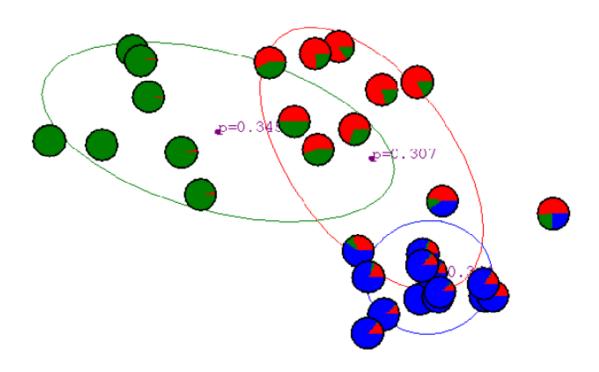
After 2nd iteration





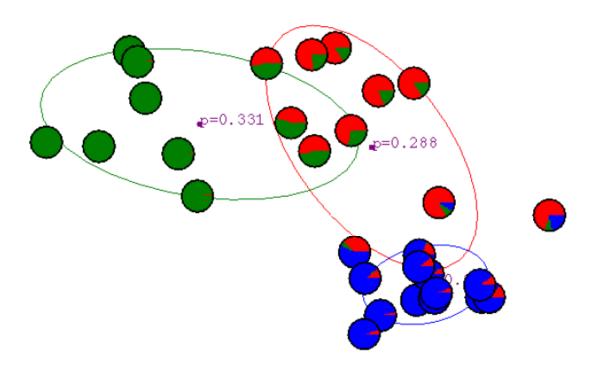
After 3rd iteration





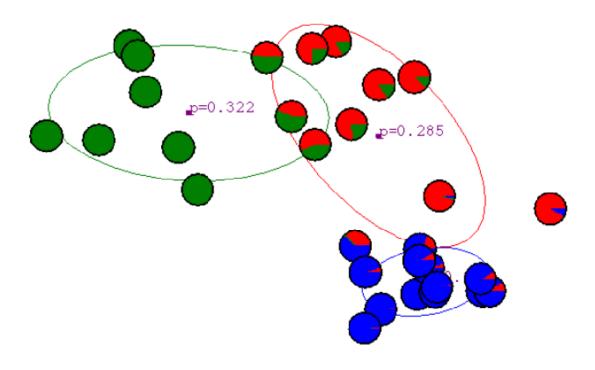
After 4th iteration





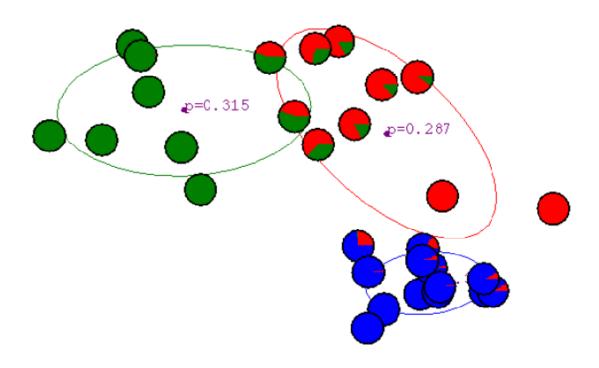
After 5th iteration





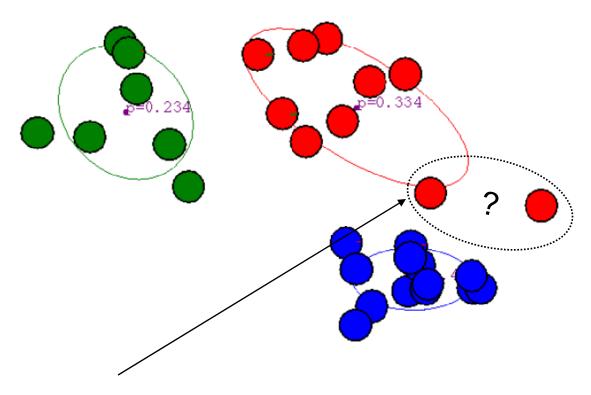
After 6th iteration





After 20th iteration





Q: Why are these two points red?

K-Means vs GMM

- we get K-Means if we make following restrictions:
 - Contain only spherical Gaussian (because all dimensions are equally contributing to the Euclidean distance function)
 - Same covariance matrix for all Gaussians
 - Use hard assignment

Behavior of EM

- It is guaranteed to converge
 - Convergence proof is based on the fact that $P(x|\theta)$ must increase or remain same between iterations (not obvious)
 - But $P(x|\theta)$ can never exceed 1 (obvious)
 - So it should always converge
 - In practice it may converge slowly, one can stop early if the change in log-likelihood is smaller than a threshold
- It converges to a local optimum
 - Multiple restart is recommended
 - Output the one with the best log-likelihood