Mixture Models & EM



Lecture outline

- Brief review
 - what k-means CANNOT do
- How to model each cluster?
 - e.g., spherical Gaussians (brief review)
- Mixture Models
 - motivation, formulation
 - estimation, the EM algorithm
 - selecting the number of mixture components



Recall: K-means

- K-means clustering
 - initialize K different means
 - assign each data point to the closest cluster mean
 - re-estimate cluster means based on the points assigned to them
 - iterate until convergence



What K-means cannot do?

- It cannot handle overlapping clusters
- It cannot represent clusters with different "spreads"
- It cannot properly deal with clusters that have different numbers of points

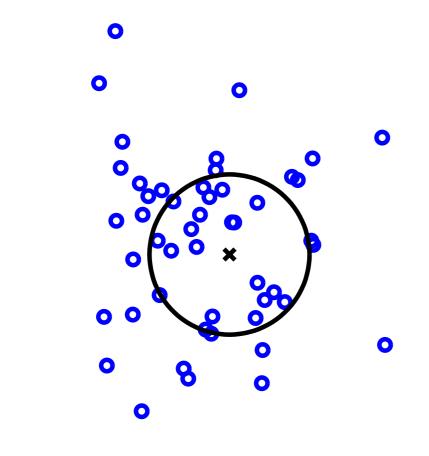


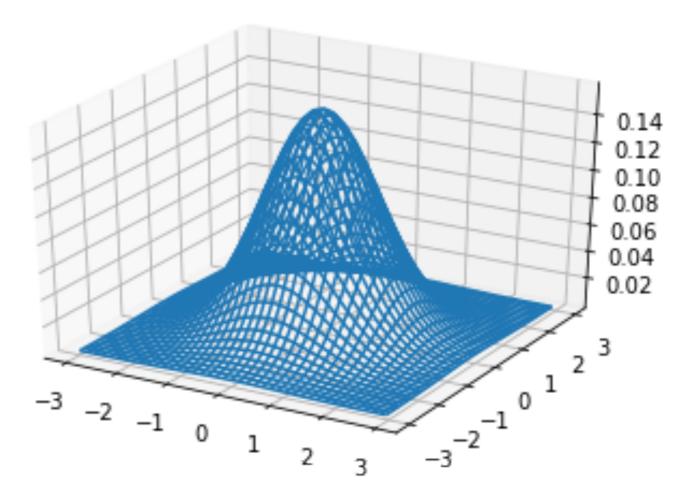
Recall: spherical Gaussian

The pdf of a spherical Gaussian

$$N(x; \mu, \sigma^2 I) = \frac{1}{(2\pi\sigma^2)^{d/2}} \exp\left(-\frac{1}{2\sigma^2} ||x - \mu||^2\right)$$

 Graphical representation (when dim d = 2) in terms of mean and stdv

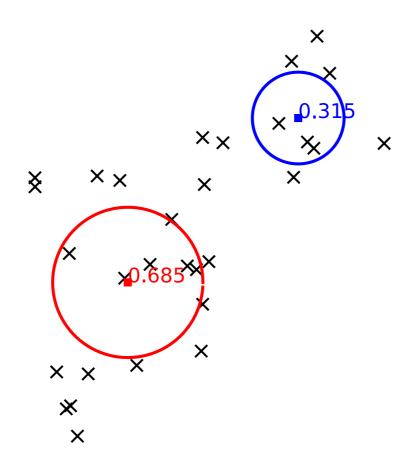






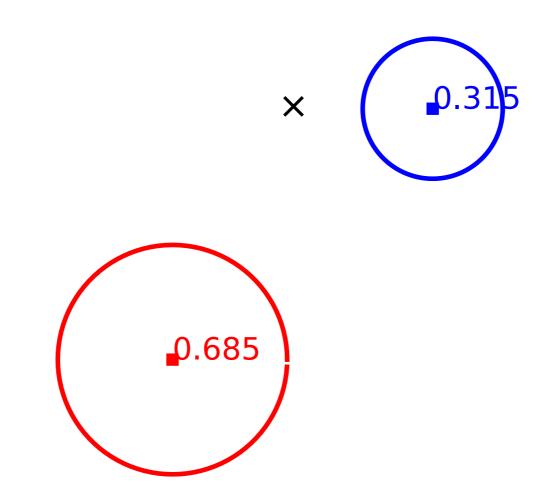
Mixture model: overview

- We use a spherical Gaussian to model each cluster
- These cluster models can have different means, variances (spreads), as well as "sizes"
- A mixture model combines these "components" into an overall probability model $P(x;\theta)$



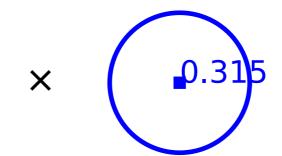
Mixture model: data generation

 We consider alternative ways that each data point x could have been created



Mixture model: data generation

- We consider alternative ways that each data point x could have been created:
 - select a cluster
 - select x from the cluster model





$$P(x;\theta) = p_1 N(x; \mu^{(1)}, \sigma_1^2 I) + p_2 N(x; \mu^{(2)}, \sigma_2^2 I)$$



Mixture model: posterior

 We can also infer (after the fact) which cluster likely generated each point by evaluating the posterior probability

$$P(j=1|x,\theta)$$
 $P(j=2|x,\theta)$

$$P(j=1|x,\theta) = \frac{p_1 N(x;\mu^{(1)},\sigma_1^2 I)}{p_1 N(x;\mu^{(1)},\sigma_1^2 I) + p_2 N(x;\mu^{(2)},\sigma_2^2 I)}$$



Mixture models: estimation

 The goal is to find the parameters of the mixture model that maximize the log-likelihood that the data points came from the mixture distribution

$$l(D; \theta) = \sum_{i=1}^{n} \log P(x^{(i)}; \theta)$$

$$= \sum_{i=1}^{n} \log \left[\sum_{j=1}^{K} p_{j} N(x^{(i)}; \mu^{(j)}, \sigma_{j}^{2} I) \right]$$

 The difficulty lies in the fact that the Gaussian cluster models cannot be estimated independently (since we don't know which data points they should generate)



The EM algorithm: overview

- The EM algorithm solves the problem by iteratively reassigning points to clusters (softly) and re-estimating the corresponding cluster models (cf. K-means)
- Initialize mixture
- E-step (complete the data)
 - evaluate the posterior probability that each data point came from a particular cluster
- M-step (maximize expected log-likelihood)
 - use the posterior probabilities (now fixed) as cluster specific weights on data points to separately reestimate each cluster model

The EM algorithm (iterative)

Initialize:

- e.g. means as randomly selected points, all variances set to overall variance, uniform mixing proportions
- **E-step:** calculate posterior assignments

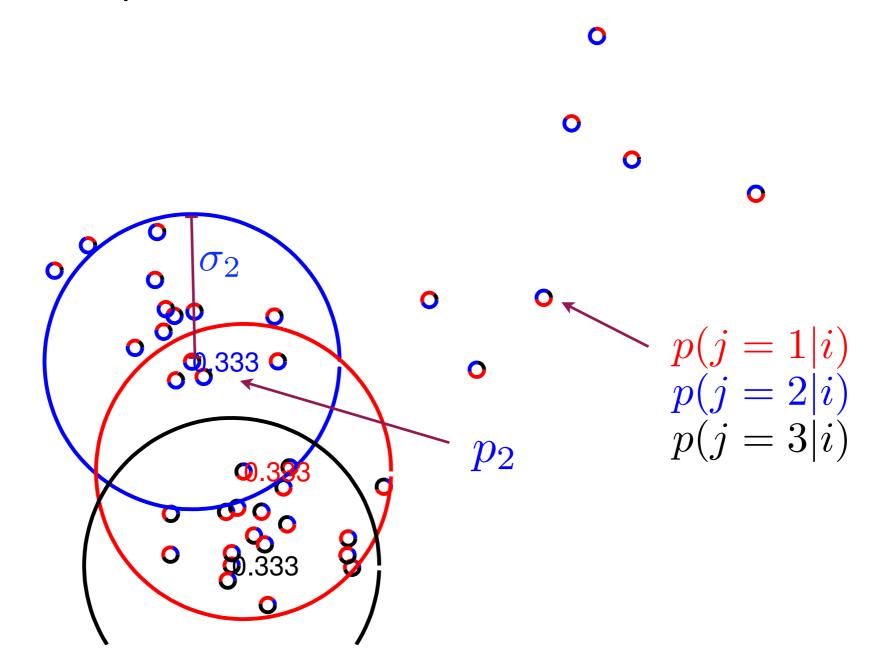
$$p(j|i) = \frac{p_j N(x^{(i)}; \mu^{(j)}, \sigma_j^2 I)}{P(x^{(i)}|\theta)}, \quad j = 1, \dots, K, \quad i = 1, \dots, n$$

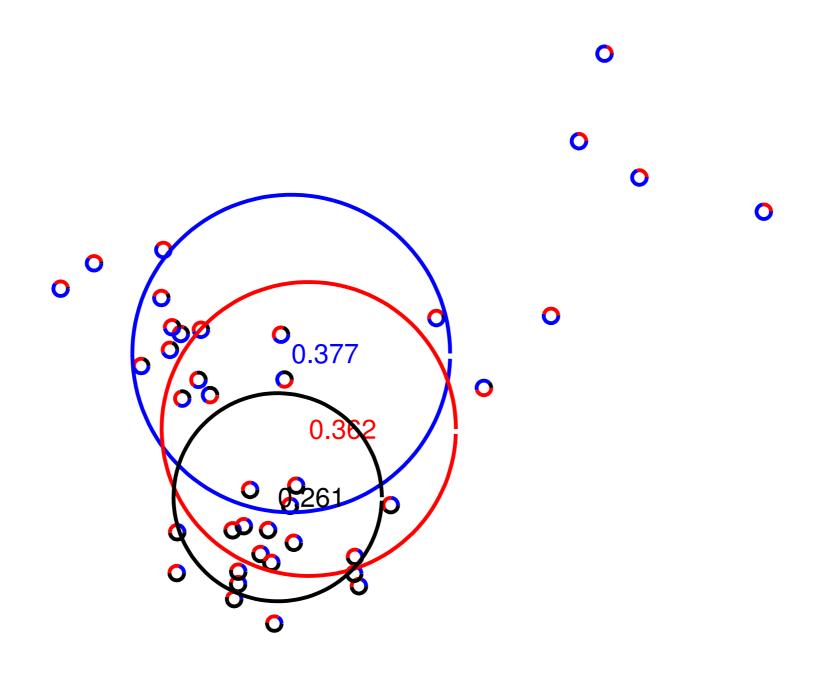
M-step: maximize

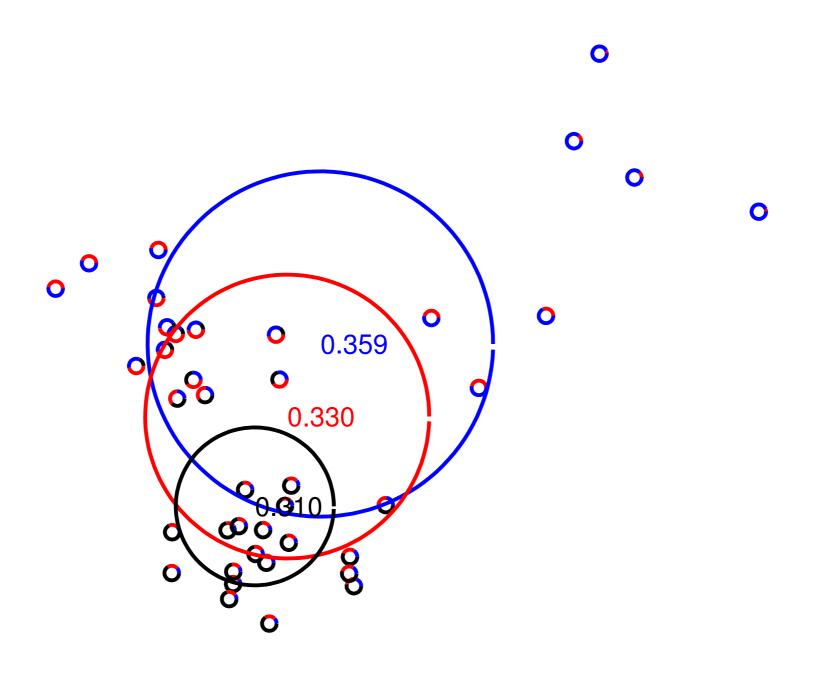
$$\tilde{l}(D; \theta) = \sum_{j=1}^{K} \sum_{i=1}^{n} p(j|i) \log \left[\frac{p_j N(x^{(i)}; \mu^{(j)}, \sigma_j^2 I)}{p(j|i)} \right]$$

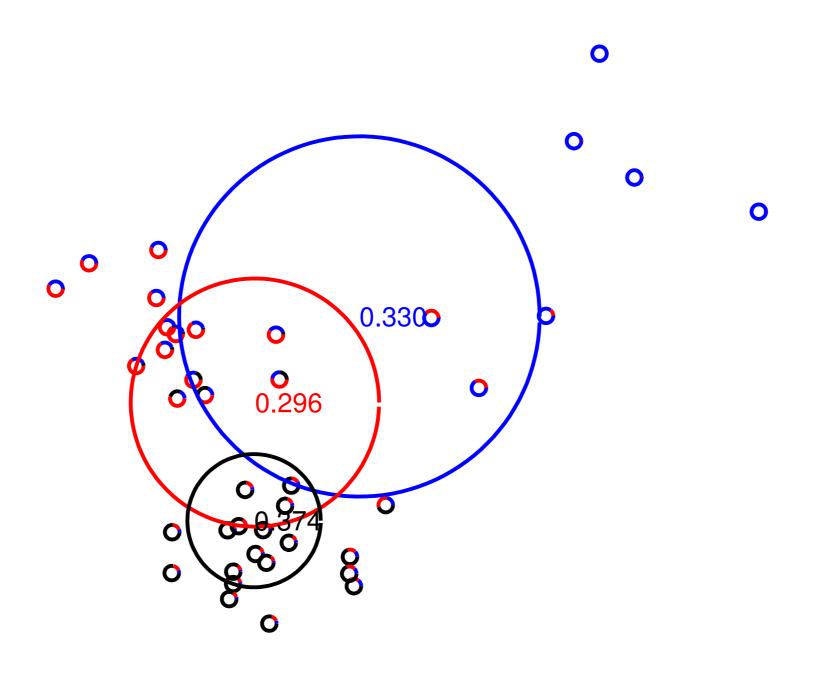
with respect to mixture parameters while keeping p(j|i)fixed

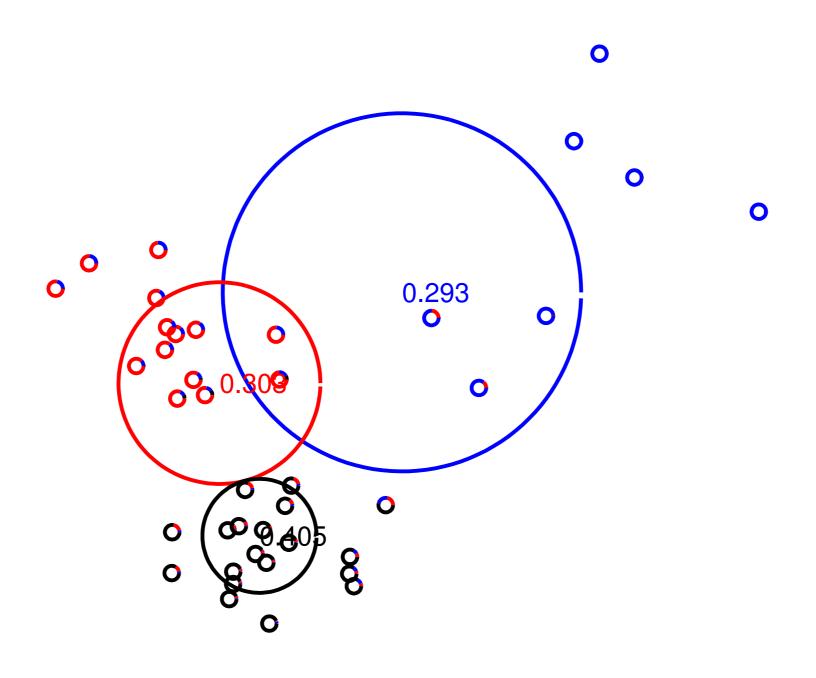
• initial 3-component mixture

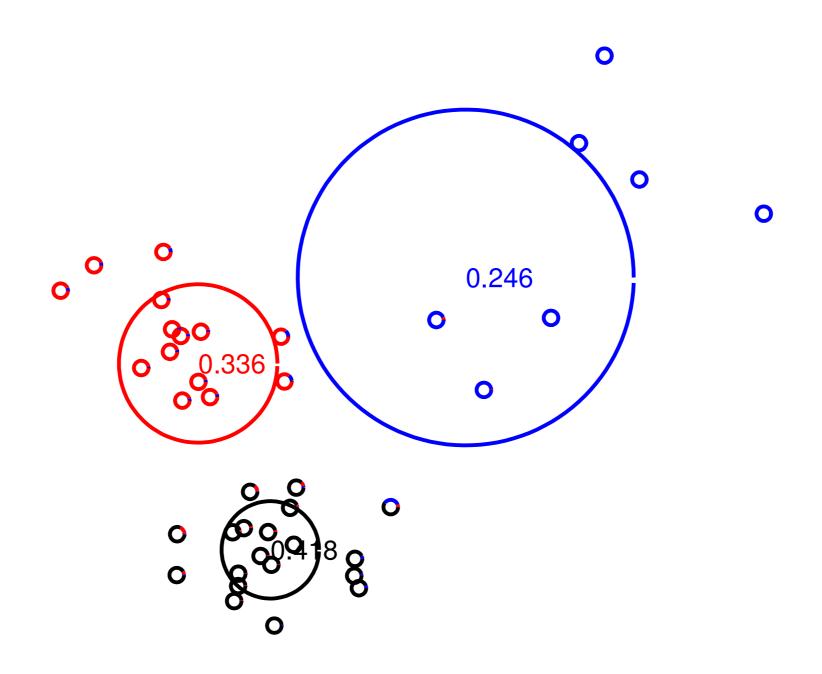


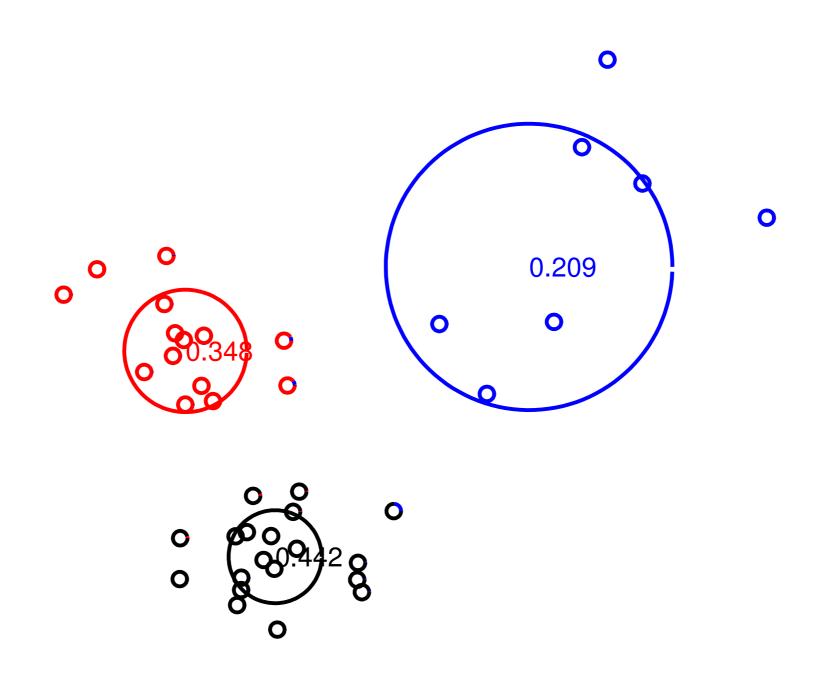


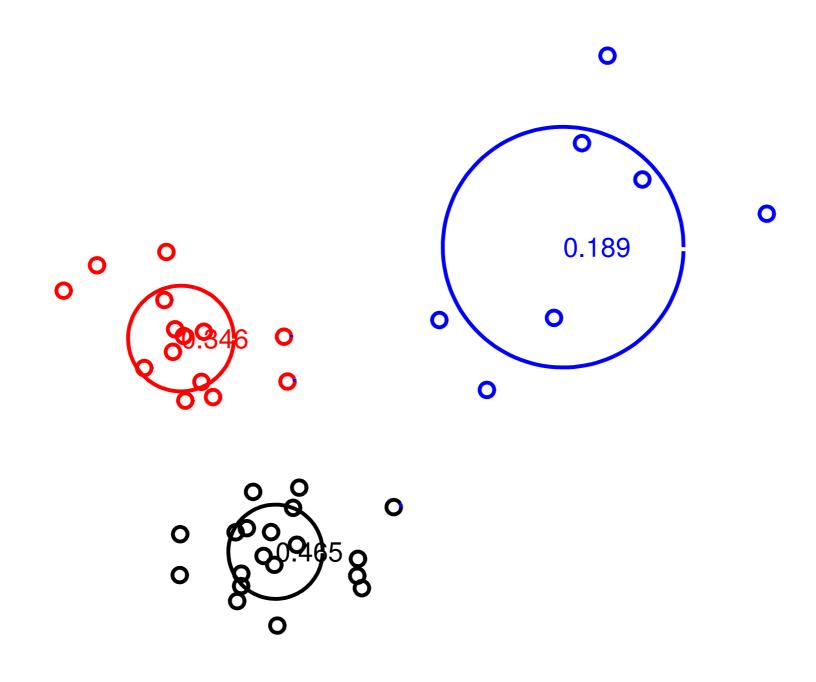






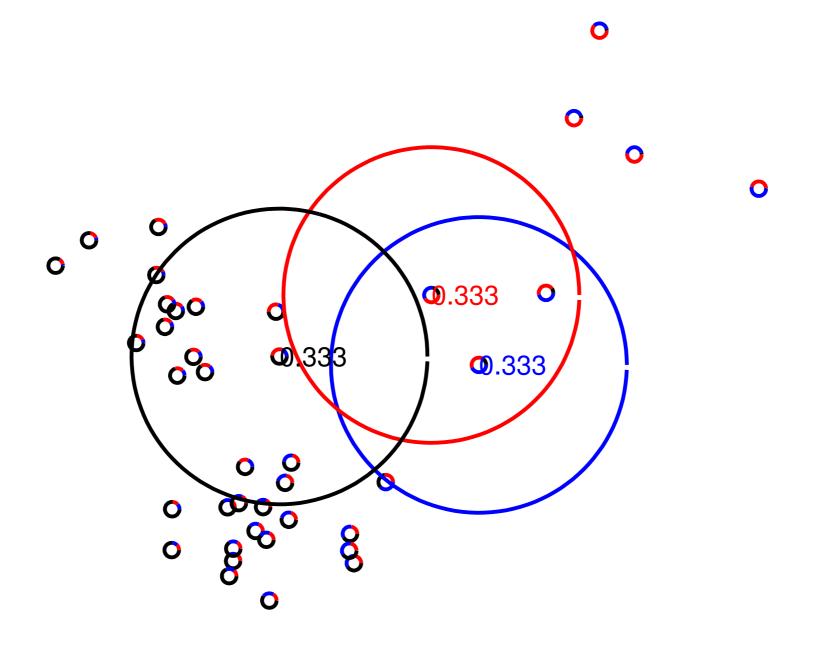


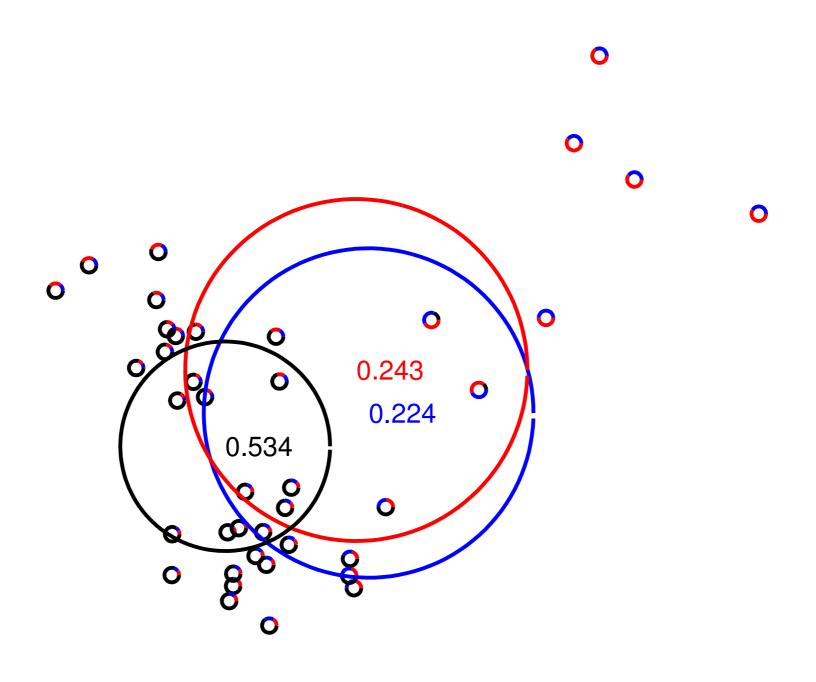


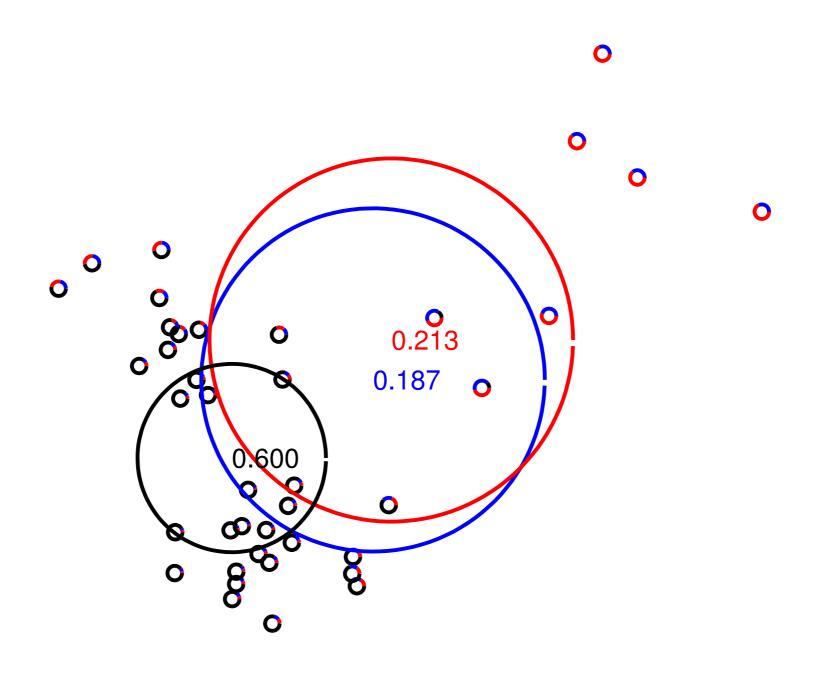


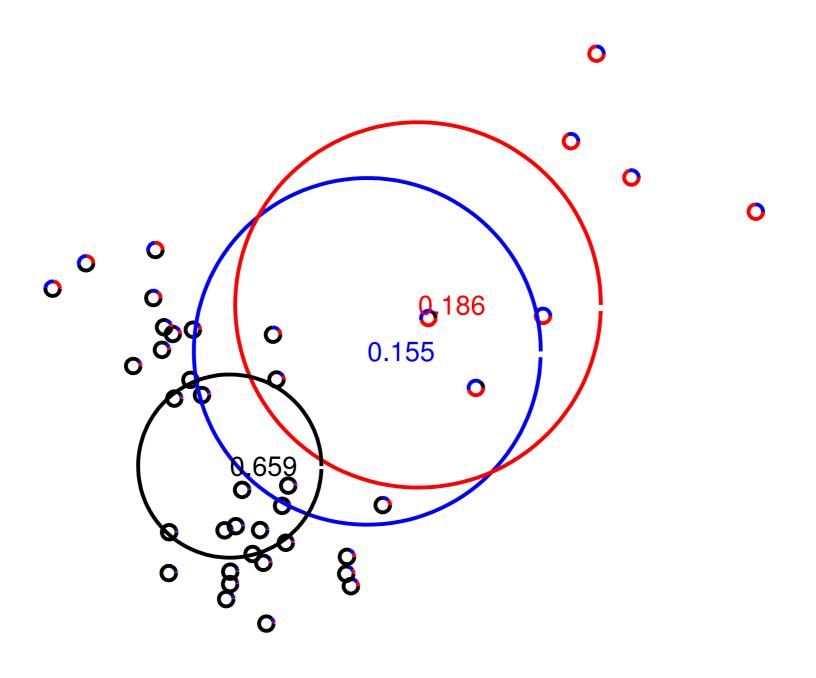


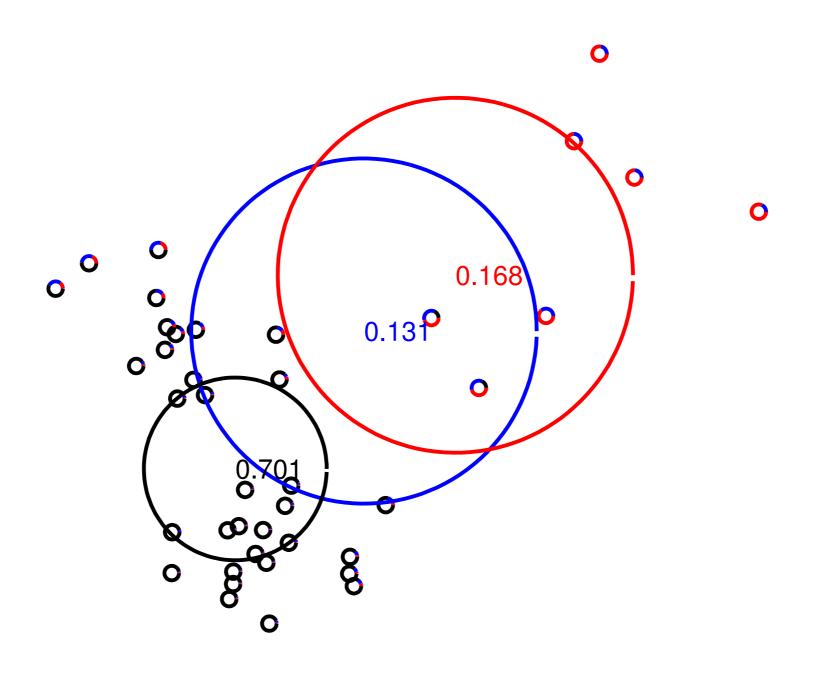
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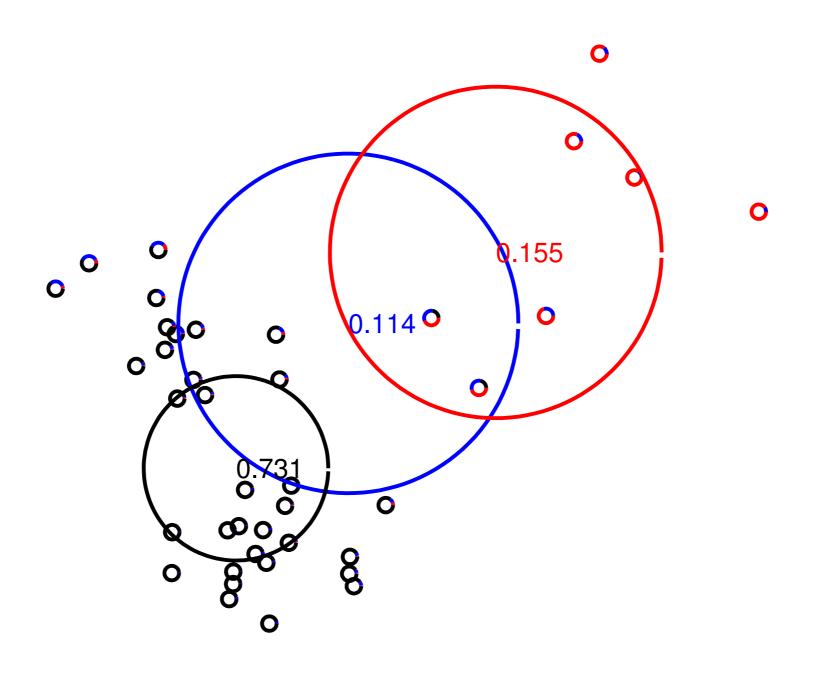


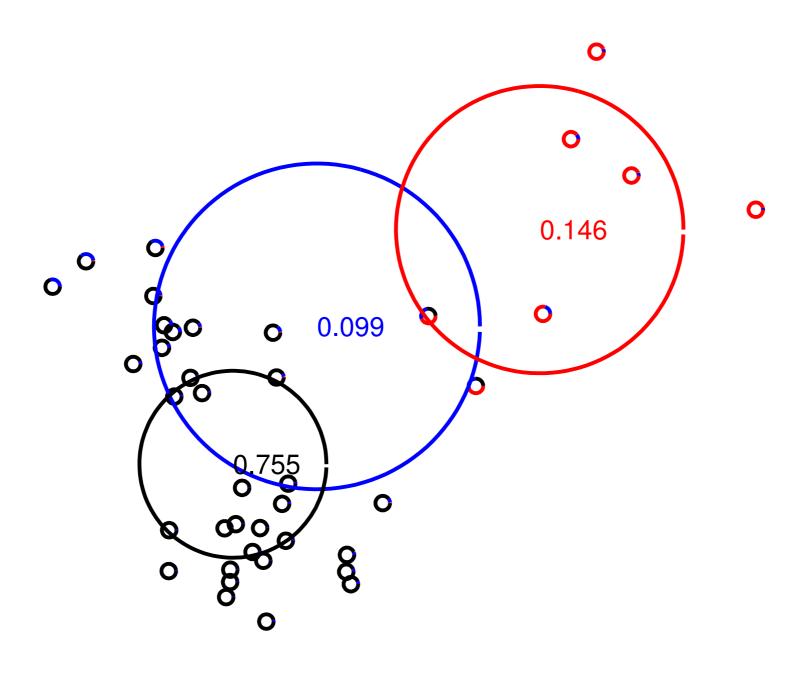


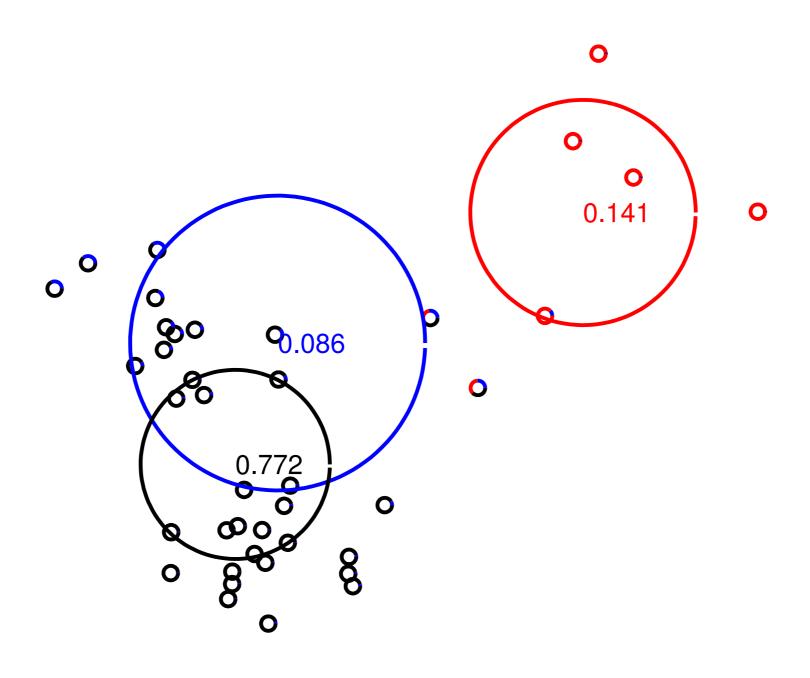


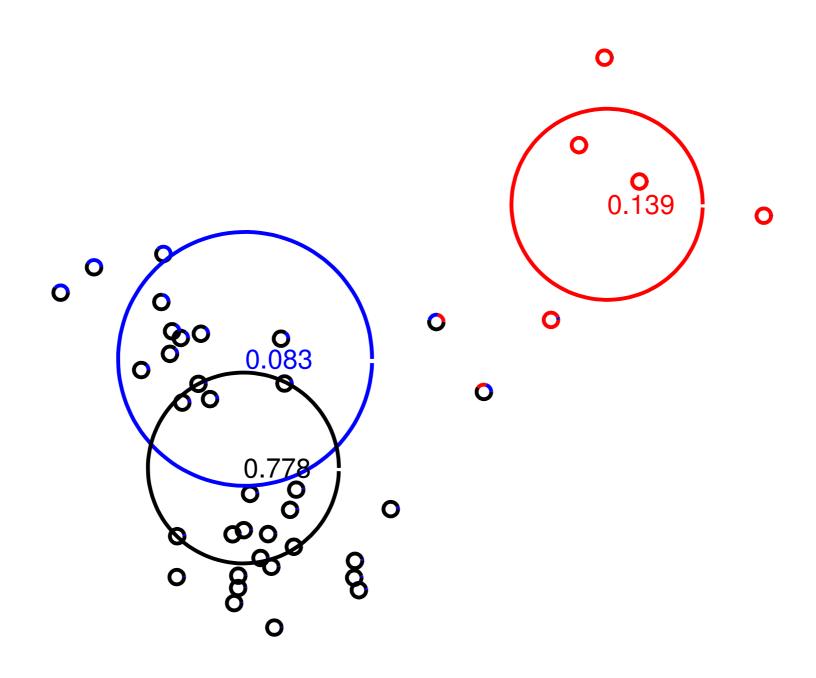


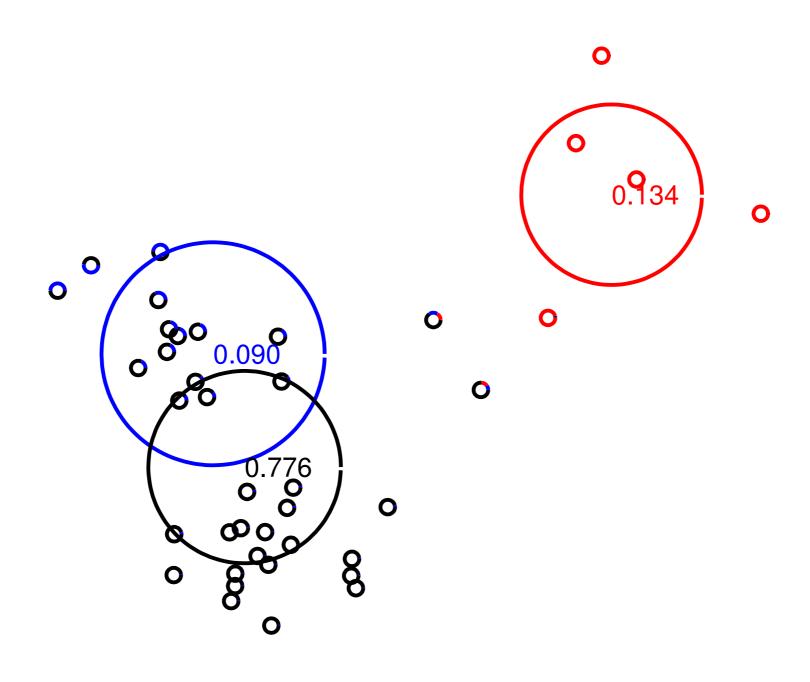


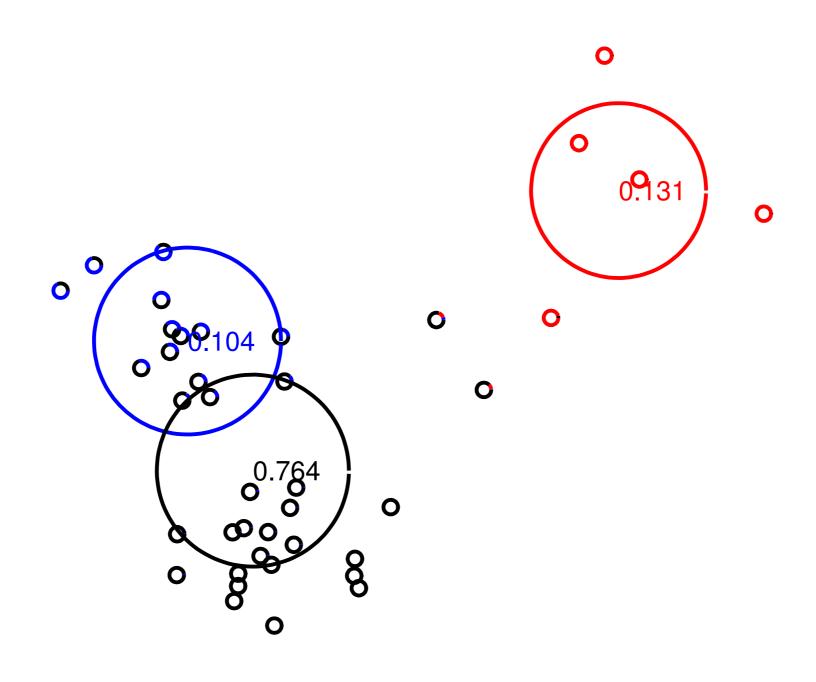


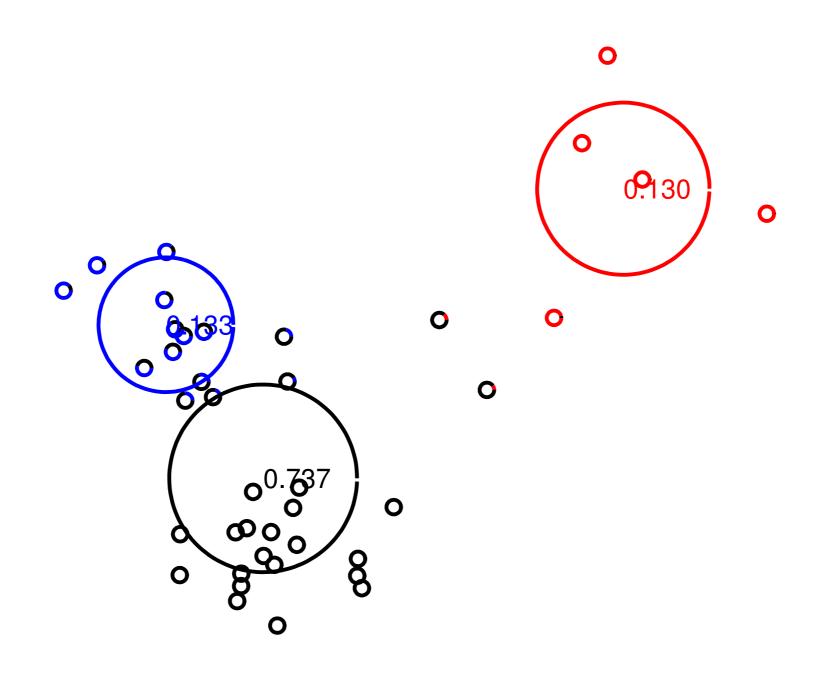


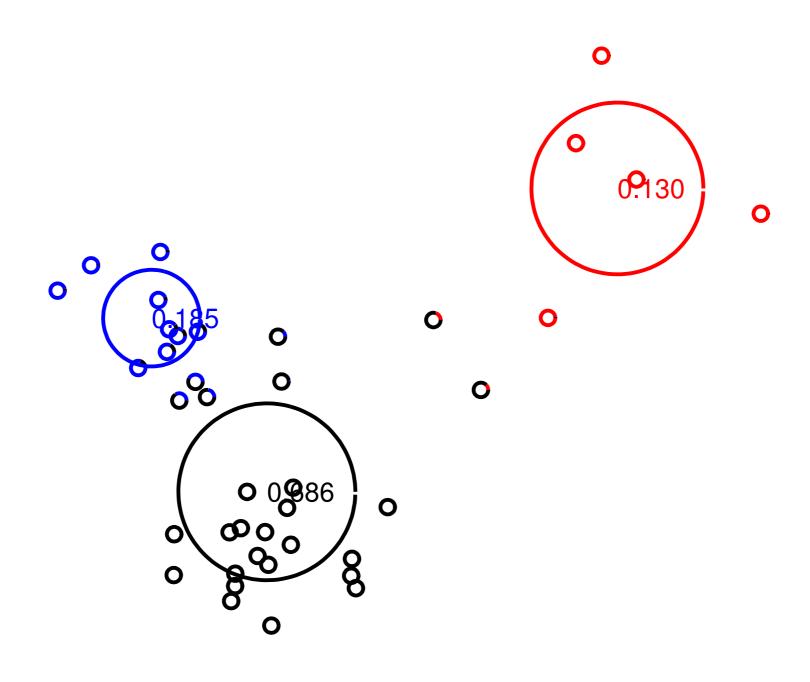








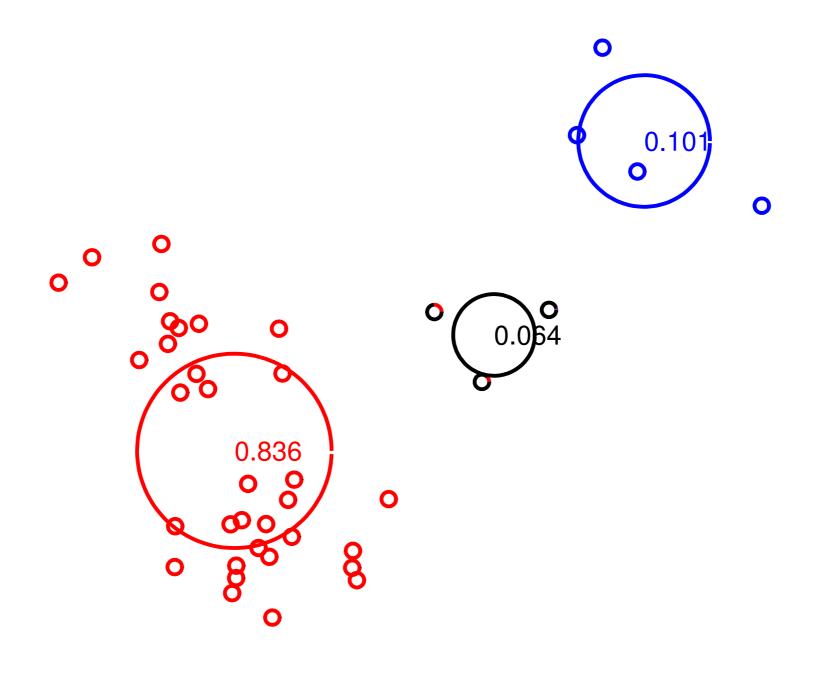




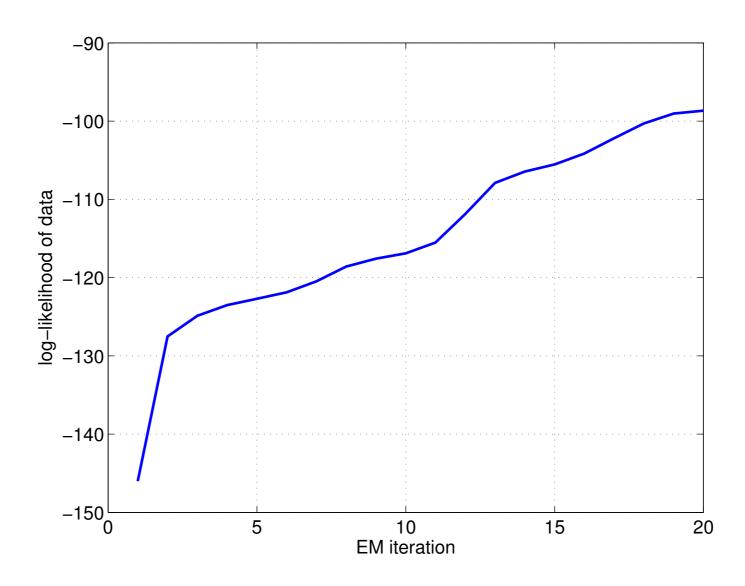




Doesn't always work... well



The EM algorithm



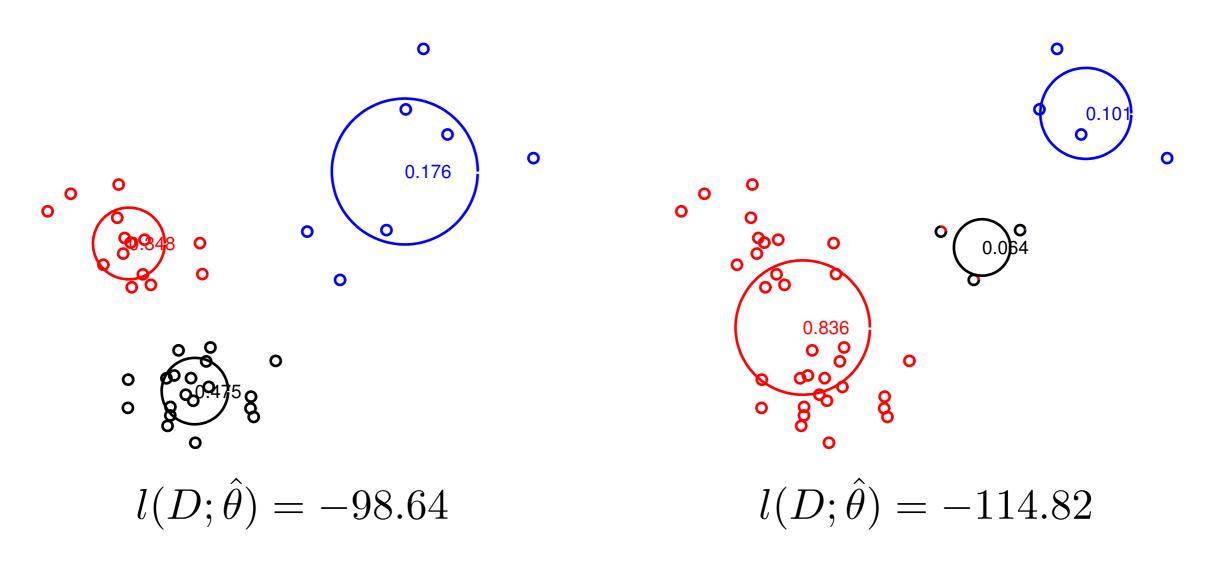
 The EM-algorithm monotonically increases the log-likelihood of the training data (cf. K-means)

$$l(D; \theta) < l(D; \theta') < l(D; \theta'') < \dots$$



Locally optimal solutions

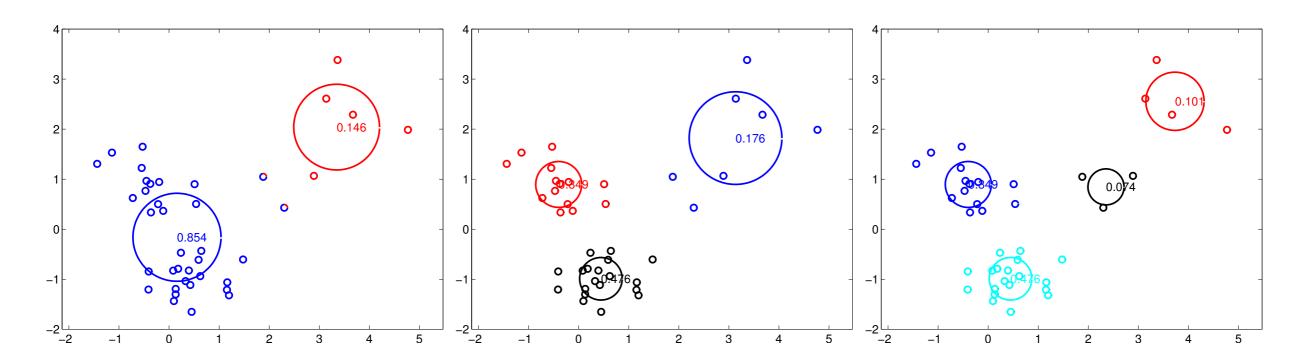
- The EM-algorithm is guaranteed to find a locally optimal solution by monotonically increasing the log-likelihood (the estimation problem with respect to θ is typically not convex)
- Whether the algorithm converges to the globally optimal solution depends on the initialization





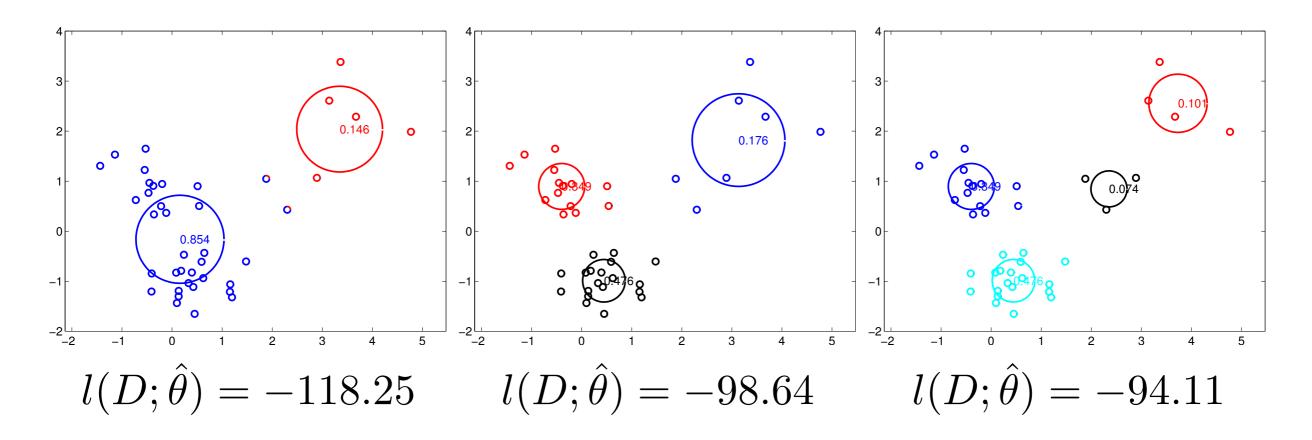
Model selection

 We can run the EM-algorithm with different numbers of components. Need to specify a criterion for selecting among the different models.



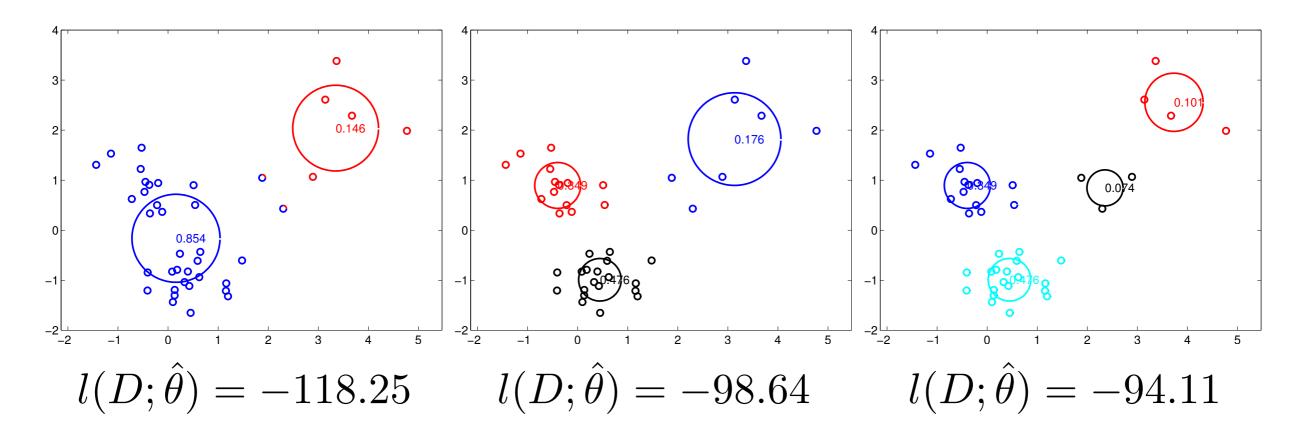
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Model selection

 We can run the EM-algorithm with different numbers of components. Need to specify a criterion for selecting among the different models.



 Basing the selecting on the value of log-likelihood would invariably lead to the largest number of components



Key things to know

- K-means failures
- Mixture model as a latent variable generative model
- Evaluating posterior probabilities
- Mixture estimation
 - ML criterion
 - complete data case
 - EM algorithm
- Why ML cannot be used to select the number of mixture components