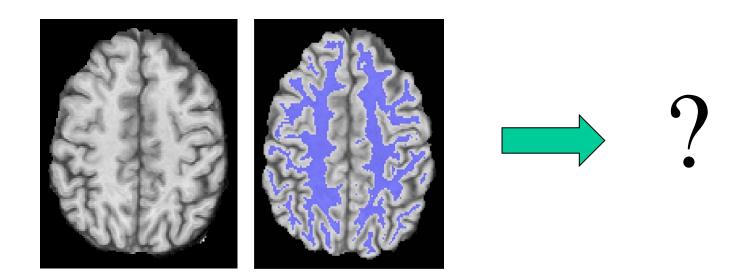
ECSE-626 Statistical Computer Vision

Gaussian Mixture Models, EM, Kernel Density Estimation

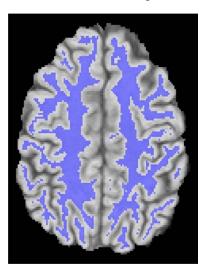
Introduction

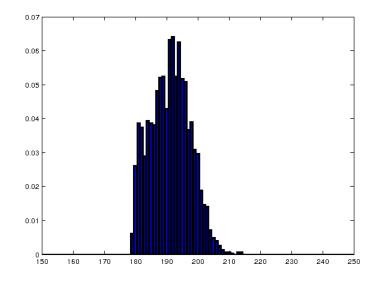
 We often want to estimate a probability density function from a set of observations.



Modeling Densities

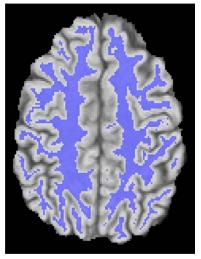
 Given a set of observations, how do we model a probability density?

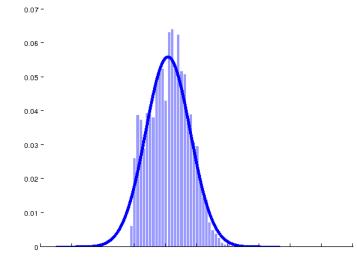




Modeling Densities

 Given a set of observations, how do we model a probability density?





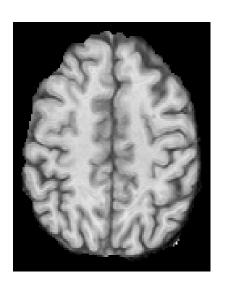
• Often we model a set of samples as having a Gaussian density

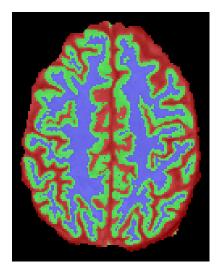
- Gaussian densities can be compactly represented by their mean and variance
- Sample mean and variance can be computed in linear time

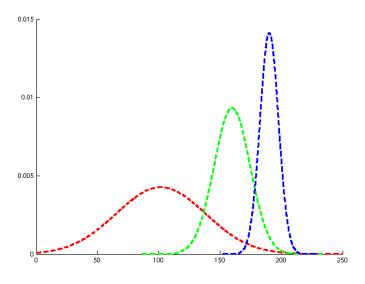
$$\bar{x} = \frac{1}{N} \sum_{i=1}^{N} x_i$$

$$s^2 = \frac{1}{N-1} \sum_{i=1}^{N} (x_i - \bar{x})^2$$

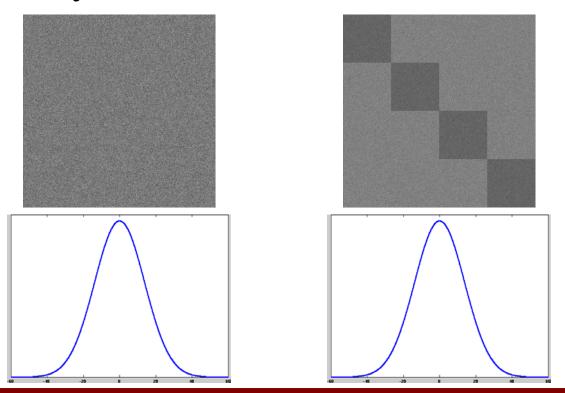
• Gaussian densities have many convenient mathematical properties and serve as a reasonable approximation for many true densities.







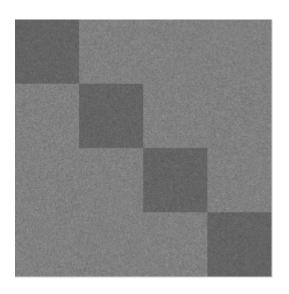
• However, many real densities are not well modeled by Gaussians.

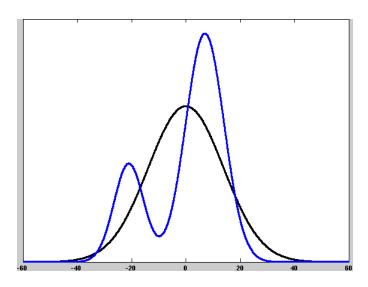


- Parametric densities such as Gaussians, and all exponential densities, are unimodal (have a single local maximum).
- Many practical problems involve multi-modal densities.

Multi-modal Densities

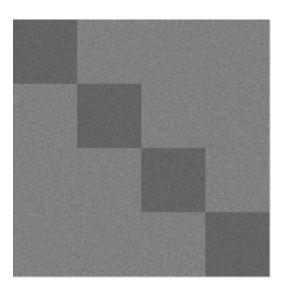
• Gaussian is poor representation of densities that have multiple modes.

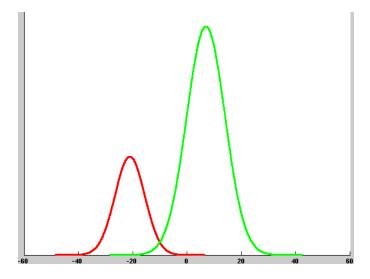




Gaussian Mixture Model (GMM)

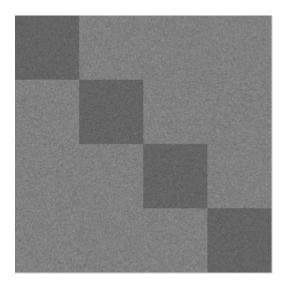
• Allow us to represent densities as weighted sum (or mixture) of multiple Gaussians

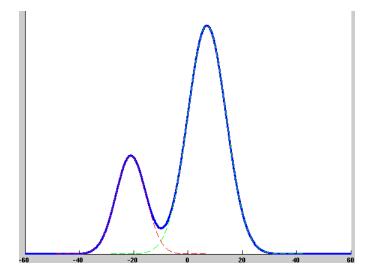




Gaussian Mixture Model (GMM)

• Allow us to represent densities as weighted sum (or mixture) of multiple Gaussians





Gaussian Mixture Models (GMM)

- Mixtures of Gaussian functions are well-suited to modeling clusters of points.
- Each cluster is assigned a Gaussian, with its mean in the middle of the cluster and with a standard deviation measuring its spread.

Gaussian Mixture Models (GMM)

K: number of gaussian components

 μ_k : mean of k^{th} gaussian component

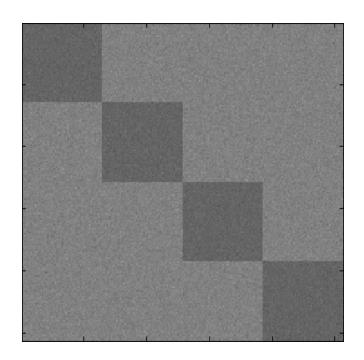
 σ_k : std of k^{th} gaussian component

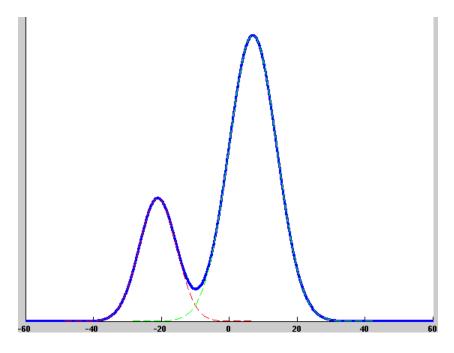
 α_k : mixing probability (weight) of k^{th} gaussian component

$$\Theta = \{\alpha_1, ..., \alpha_k, \mu_1, ..., \mu_k, \sigma_1, ..., \sigma_k\}$$

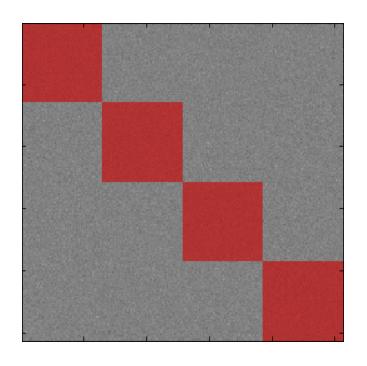
$$f(x|\mathbf{\Theta}) = \sum_{k=1}^{K} \alpha_k \mathcal{N}(x; \mu_k, \sigma_k)$$

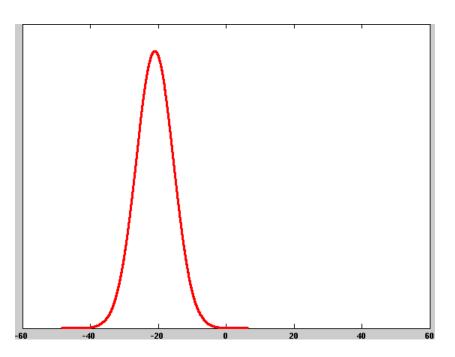
$$K=2,\,\mu=\{-21,7.00\},\,\sigma=\{30,50\},\,\alpha=\{0.25,0.75\}$$





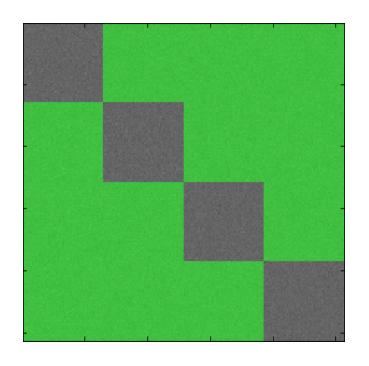
$$K=2,\,\mu=\{-21,7.00\},\,\sigma=\{30,50\},\,\alpha=\{0.25,0.75\}$$

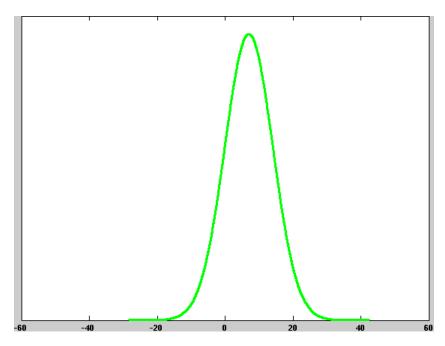




k = 1

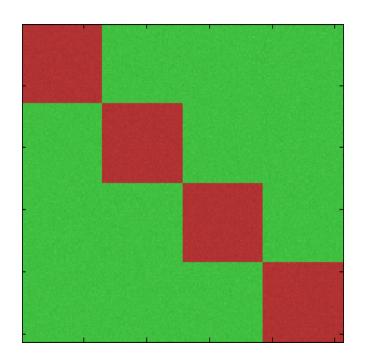
$$K=2,\,\mu=\{-21,7.00\},\,\sigma=\{30,50\},\,\alpha=\{0.25,0.75\}$$

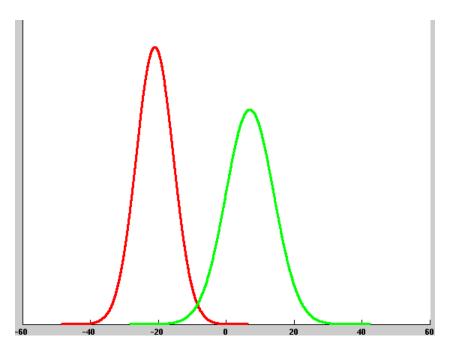




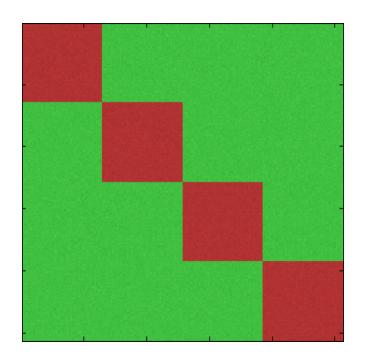
k = 2

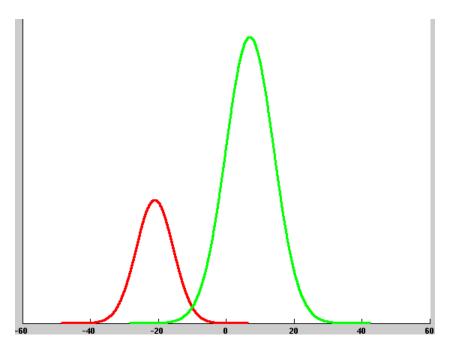
$$K=2,\,\mu=\{-21,7.00\},\,\sigma=\{30,50\},\,\alpha=\{0.25,0.75\}$$



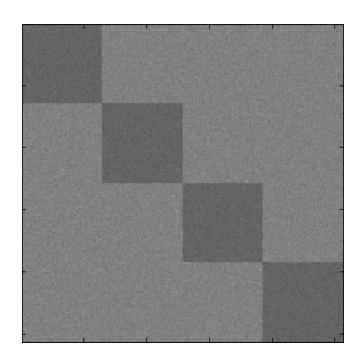


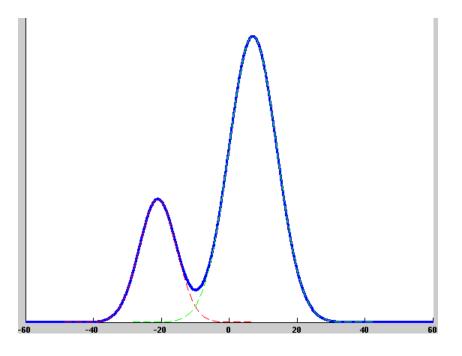
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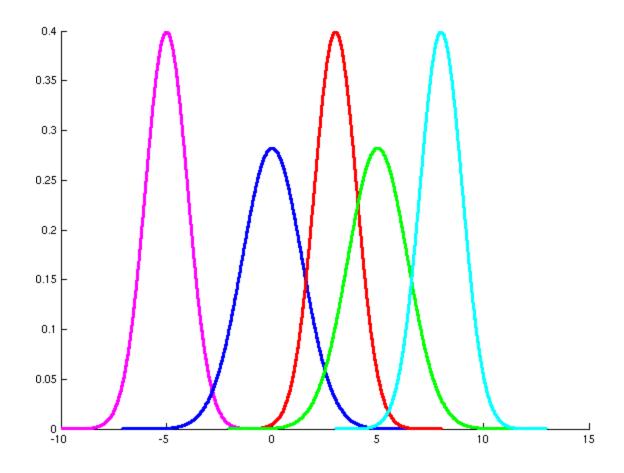


$$K=2,\,\mu=\{-21,7.00\},\,\sigma=\{30,50\},\,\alpha=\{0.25,0.75\}$$

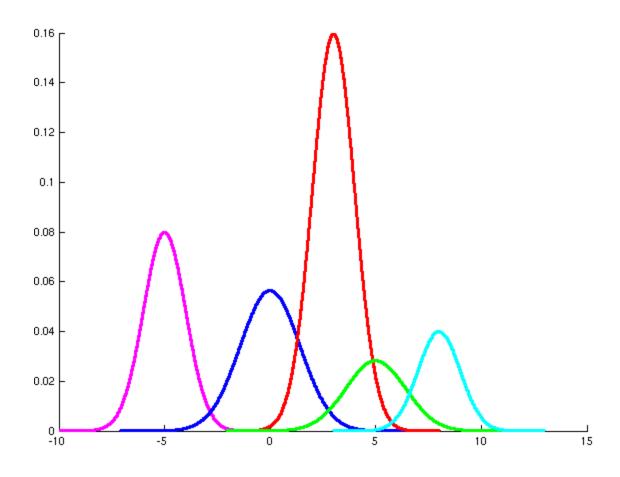




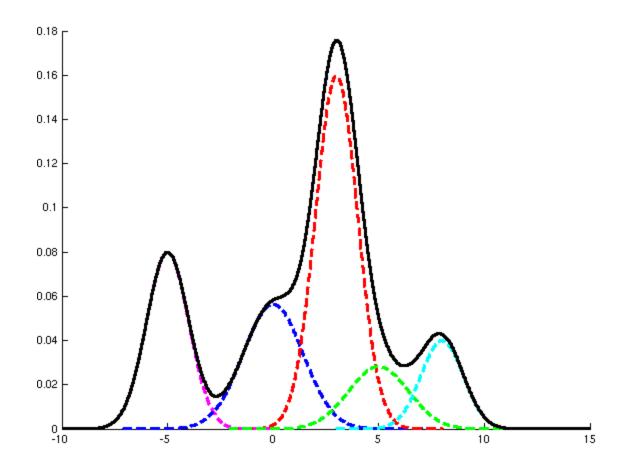
 $K=5,\,\mu=\{-5,0,3,5,8\},\,\sigma=\{1,2,1,2,1\},\,\alpha=\{0.2,0.2,0.4,0.1,0.1\}$



 $K=5,\,\mu=\{-5,0,3,5,8\},\,\sigma=\{1,2,1,2,1\},\,\alpha=\{0.2,0.2,0.4,0.1,0.1\}$

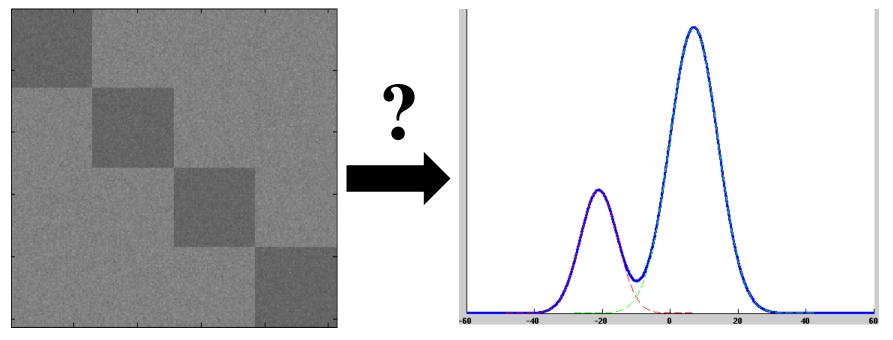


 $K = 5, \ \mu = \{-5, 0, 3, 5, 8\}, \ \sigma = \{1, 2, 1, 2, 1\}, \ \alpha = \{0.2, 0.2, 0.4, 0.1, 0.1\}$



- For most real problems, we don't know how many or which Gaussians the data are sampled from.
- How do we automatically determine the parameters of a GMM from observed samples?

Unsupervised Learning - GMMs



• Given an image we want to determine parameters Θ:

$$\mathbf{\Theta} = \{\alpha_1, ..., \alpha_k, \mu_1, ..., \mu_k, \sigma_1, ..., \sigma_k\}$$

• Assume we know form of model (GMM):

$$p(x|\mathbf{\Theta}) = \sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x; \mu_k, \sigma_k)$$

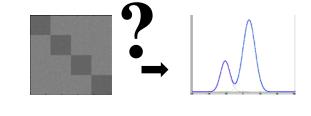
• Assume we know form of model (GMM):

$$p(x|\mathbf{\Theta}) = \sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x; \mu_k, \sigma_k)$$

• Assume observed samples, x_i , are i.i.d.:

$$p(\mathcal{X}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i|\mathbf{\Theta})$$

$$p(\mathcal{X}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i|\mathbf{\Theta})$$



N:# samples, i: sample (pixel) index, $\Theta:$ GMM parameters, K:# components in GMM

$$p(\mathcal{X}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i|\mathbf{\Theta})$$

$$= \prod_{i=1}^{N} \sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k)$$

N: # samples, i: sample (pixel) index, $\Theta:$ GMM parameters, K: # components in GMM

$$p(\mathcal{X}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i|\mathbf{\Theta})$$

$$= \prod_{i=1}^{N} \sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k)$$

• Want to find model parameters, Θ , that maximize likelihood.

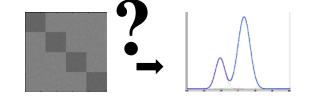
N: # samples, i: sample (pixel) index, $\Theta:$ GMM parameters, K: # components in GMM

$$\mathbf{\Theta}^* = \operatorname*{argmax} p(\mathcal{X}|\mathbf{\Theta})$$



$$\Theta^* = \underset{\Theta}{\operatorname{argmax}} p(\mathcal{X}|\Theta)$$

$$= \underset{\Omega}{\operatorname{argmax}} \log p(\mathcal{X}|\Theta)$$



$$\mathbf{\Theta}^* = \operatorname*{argmax} p(\mathcal{X}|\mathbf{\Theta})$$



$$= \operatorname*{argmax} \log p(\mathcal{X}|\mathbf{\Theta})$$

$$= \underset{\boldsymbol{\Theta}}{\operatorname{argmax}} \log \prod_{i=1}^{N} \sum_{k=1}^{N} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k)$$

$$\mathbf{\Theta}^* = \operatorname*{argmax} p(\mathcal{X}|\mathbf{\Theta})$$



$$= \operatorname*{argmax} \log p(\mathcal{X}|\mathbf{\Theta})$$

$$= \underset{\boldsymbol{\Theta}}{\operatorname{argmax}} \log \prod_{i=1}^{N} \sum_{k=1}^{N} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k)$$

$$= \underset{\boldsymbol{\Theta}}{\operatorname{argmax}} \sum_{i=1}^{N} \log(\sum_{k=1}^{K} \alpha_k \mathcal{N}(x_i; \mu_k, \sigma_k))$$

$$\mathbf{\Theta}^* = \underset{\mathbf{\Theta}}{\operatorname{argmax}} \sum_{i=1}^{N} \log(\sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k))$$

• Difficult to optimize because contains log of summation

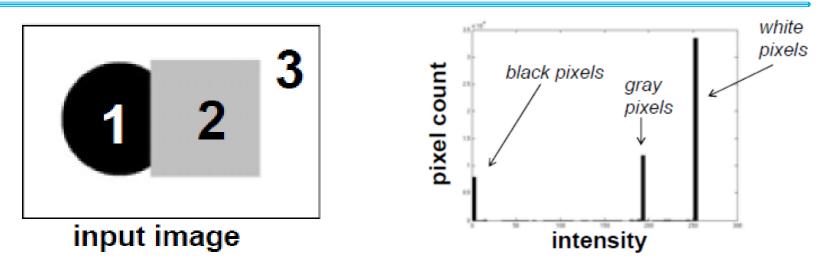
Expectation Maximization (EM)

- Iterative algorithm for finding maximumlikelihood estimate of model parameters from a given data set.
- Expectation: determine which Gaussian component(s) each sample comes from based on current estimate of model parameters
- Maximization: maximize model parameters given current estimates of Gaussian sources of each sample

GMMs for Segmentation

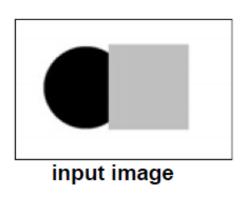
- We have started to look at the use of GMMs / EM for estimating multi-modal probability densities.
- GMMs / EM are also often used for segmentation or clustering tasks
- EM can be thought of as conceptually similar to K-means
- Recall K-means for segmentation....

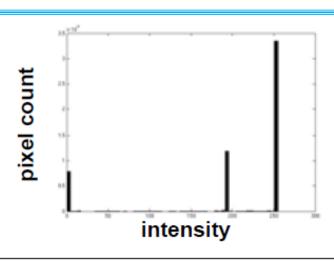
Image Segmentation: toy example

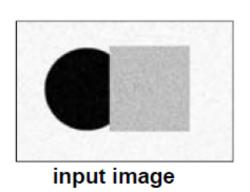


- These intensities define the three groups
- We could label every pixel in the image according to which of these primary intensities it is
 - i.e., *segment* the image based on the intensity feature.
- What if the image isn't quite so simple?

Image Segmentation: toy example







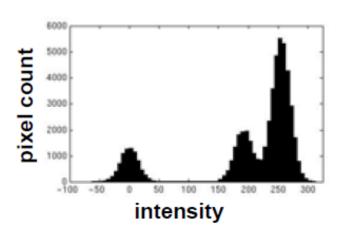
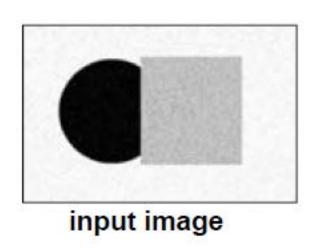
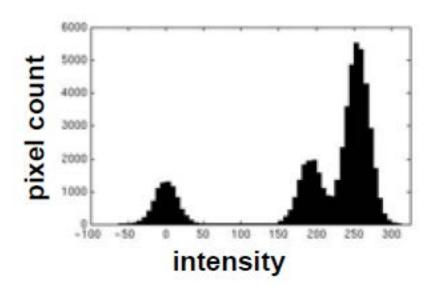


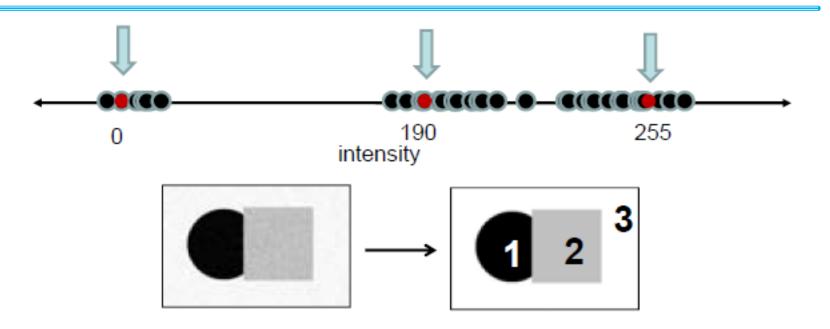
Image Segmentation: toy example





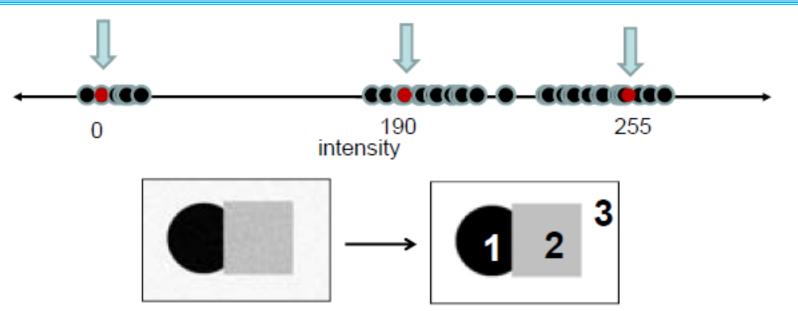
- Now how to determine the three main intensities that define our groups?
- We need to *cluster*

Image Segmentation: Clustering



• Goal: choose three *centers* as the representative intensities, and label every pixel according to which of these centers it is nearest to

Image Segmentation: Clustering

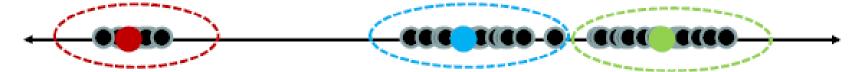


• Best cluster centers are those that minimize SSD between all points and their nearest cluster center c_i

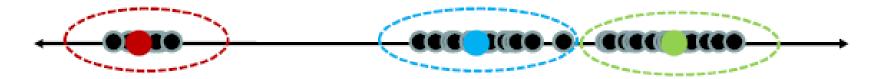
$$\sum_{\text{cluster } i \text{ points } p \text{ in cluster } i} ||p - c_i||^2$$

Image Segmentation: Clustering

- With this objective, it is a "chicken and egg" problem
 - If we knew the cluster centers, we could allocate points to groups by assigning each to its closest center



 If we knew the group memberships, we could get the centers by computing the mean per group



K-means Clustering Algorithm

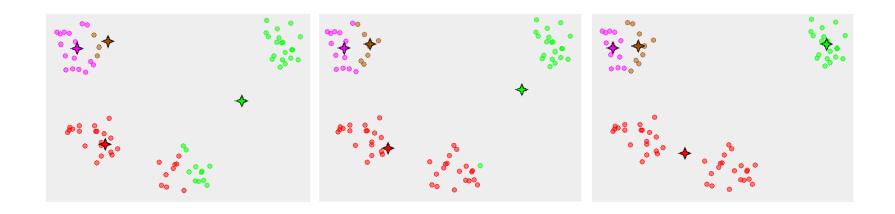
- Randomly initialize the *K* cluster centers, and iterate between the two steps we just saw
 - 1. Randomly initialize the cluster centers c_1, \dots, c_K
 - 2. Given the cluster centers, determine points in each cluster
 - For each point p, find the closest c_i . Put p into cluster i
 - 3. Given points in each cluster, solve for c_i
 - Set c_i to be the mean of points in cluster i
 - 4. If c_i have changed, repeat Step 2

4

Steve Seitz

K-means Clustering

• Converging to a local minimum



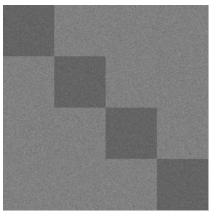
GMMs for Segmentation

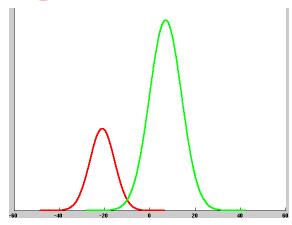
- GMMs / EM are also often used for segmentation or clustering tasks
- We can think of each of the *K* components in our GMM as representing a distinct *class* and each sample as having a membership to one or more of the *K* classes.

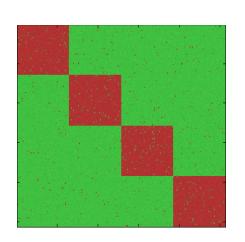
GMMs for Segmentation

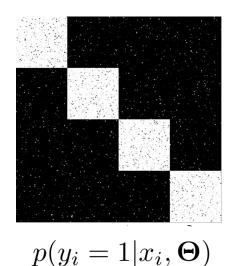
- We can model our image as a GMM consisting of *K* components, each representing a distinct class.
- We can then assign a likelihood of each of the
 K classes for every sample in our image

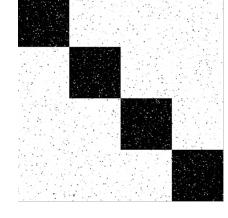
GMMs for Segmentation



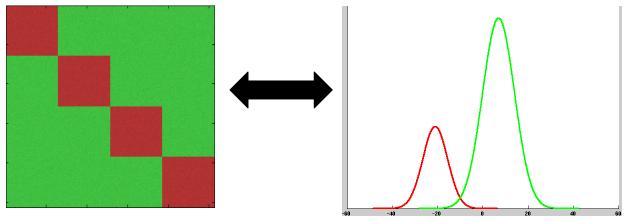




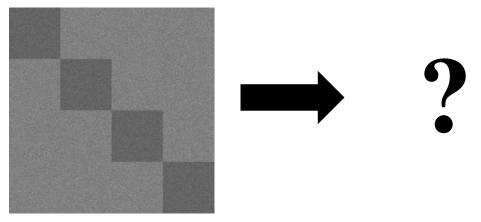




$$p(y_i = 2|x_i, \mathbf{\Theta})$$



- If we know which class (gaussian component) each sample came from, the we can estimate the parameters of our GMM.
- If we know the parameters of the GMM, we can determine the likelihood of each sample belonging to each class.



• In practice we don't know the GMM parameters or the classes of each sample but only have the image.

$$\mathbf{\Theta}^* = \underset{\mathbf{\Theta}}{\operatorname{argmax}} \sum_{i=1}^{N} \log(\sum_{k=1}^{K} \alpha_k \, \mathcal{N}(x_i; \mu_k, \sigma_k))$$

• To simplify our optimization, we introduce a hidden parameter, Y, that acts as a class label (corresponding to one of K gaussian clusters) for each sample, x_i :

$$\mathcal{Y} = \{y_i\}_{i=1}^N, y_i \in \{1, ..., K\}$$

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- For density estimation, y_i acts as a hidden parameter that will allow us to solve for our GMM parameters, Θ .

$$\mathcal{Y} = \{y_i\}_{i=1}^N, y_i \in \{1, ..., K\}$$

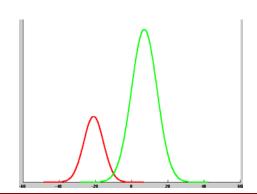
- For segmentation, we can think of each y_i as a class label for sample x_i
- For density estimation, y_i acts as a hidden parameter that will allow us to solve for our GMM parameters, Θ .
- In general, each y_i will be represented as a distribution over all K possible classes.

$$p(x_i, y_i | \mathbf{\Theta}) = p(x_i | y_i, \mathbf{\Theta}) p(y_i | \mathbf{\Theta})$$

$$p(x_i, y_i | \mathbf{\Theta}) = p(x_i | y_i, \mathbf{\Theta}) p(y_i | \mathbf{\Theta})$$

• We can simplify by observing the following:

$$p(y_i|\mathbf{\Theta}) = \alpha_{y_i}$$

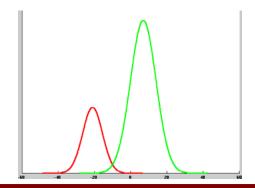


$$p(x_i, y_i | \mathbf{\Theta}) = p(x_i | y_i, \mathbf{\Theta}) p(y_i | \mathbf{\Theta})$$

• We can simplify by observing the following:

$$p(y_i|\mathbf{\Theta}) = \alpha_{y_i}$$

$$p(x_i|y_i, \mathbf{\Theta}) = \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i})$$



$$p(x_i, y_i | \mathbf{\Theta}) = p(x_i | y_i, \mathbf{\Theta}) p(y_i | \mathbf{\Theta})$$

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$$p(x_i, y_i | \mathbf{\Theta}) = \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i}) \alpha_{y_i}$$

$$p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i, y_i|\mathbf{\Theta})$$

$$p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta}) = \prod_{i=1}^{N} p(x_i, y_i|\mathbf{\Theta})$$
$$= \prod_{i=1}^{N} \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i}) \alpha_{y_i}$$

$$\mathbf{\Theta}^* = \operatorname*{argmax} p(\mathcal{X}, \mathcal{Y} | \mathbf{\Theta})$$

$$\mathbf{\Theta}^* = \underset{\mathbf{\Theta}}{\operatorname{argmax}} p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})$$

$$= \underset{\mathbf{\Theta}}{\operatorname{argmax}} \log p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})$$

$$\Theta^* = \underset{\Theta}{\operatorname{argmax}} p(\mathcal{X}, \mathcal{Y}|\Theta)$$

$$= \underset{\Theta}{\operatorname{argmax}} \log p(\mathcal{X}, \mathcal{Y}|\Theta)$$

$$= \underset{i=1}{\operatorname{argmax}} \log \prod_{i=1}^{N} \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i}) \alpha_{y_i}$$

$$\Theta^* = \underset{\Theta}{\operatorname{argmax}} p(\mathcal{X}, \mathcal{Y}|\Theta)$$

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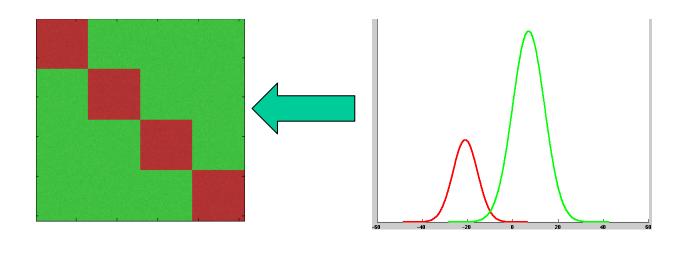
$$= \underset{\Theta}{\operatorname{argmax}} \log \prod_{i=1}^{N} \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i}) \alpha_{y_i}$$

$$= \underset{\Theta}{\operatorname{argmax}} \sum_{i=1}^{N} \log \alpha_{y_i} \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i})$$

$$\mathbf{\Theta}^* = \underset{\mathbf{\Theta}}{\operatorname{argmax}} \sum_{i=1}^{N} \log \alpha_{y_i} \, \mathcal{N}(x_i; \mu_{y_i}, \sigma_{y_i})$$

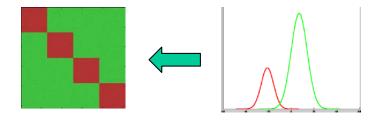
- One problem : we don't actually have y_i ...
- Iterative optimization
 - E-step: estimate y_i based on current estimate of parameters
 - M-step : find parameters, $\boldsymbol{\Theta}$, that maximize expectation of log-likelihood
- Requires initial estimate of parameters, Θ , or of y_i

• Determine density of the hidden parameters, y_{i} , based on current estimate of model parameters

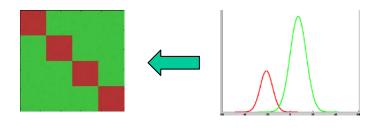


$$p(y_i|x_i, \mathbf{\Theta}^t)$$

$$p(y_i|x_i, \mathbf{\Theta}^t) = \frac{p(x_i|y_i, \mathbf{\Theta}^t)p(y_i|\mathbf{\Theta}^t)}{p(x_i|\mathbf{\Theta}^t)}$$



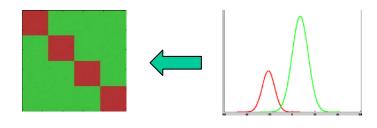
$$p(y_i|x_i, \mathbf{\Theta}^t) = \frac{p(x_i|y_i, \mathbf{\Theta}^t)p(y_i|\mathbf{\Theta}^t)}{p(x_i|\mathbf{\Theta}^t)}$$
$$= \frac{\mathcal{N}(x_i; \mu_{y_i}^t, \sigma_{y_i}^t)\alpha_{y_i}^t}{p(x_i|\mathbf{\Theta}^t)}$$



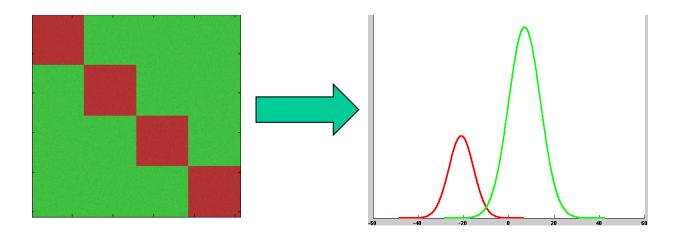
$$p(y_i|x_i, \mathbf{\Theta}^t) = \frac{p(x_i|y_i, \mathbf{\Theta}^t)p(y_i|\mathbf{\Theta}^t)}{p(x_i|\mathbf{\Theta}^t)}$$

$$= \frac{\mathcal{N}(x_i; \mu_{y_i}^t, \sigma_{y_i}^t)\alpha_{y_i}^t}{p(x_i|\mathbf{\Theta}^t)}$$

$$= \frac{\mathcal{N}(x_i; \mu_{y_i}^t, \sigma_{y_i}^t)\alpha_{y_i}^t}{\sum_{k=1}^K \alpha_k^t \mathcal{N}(x_i; \mu_k^t, \sigma_k^t)}$$



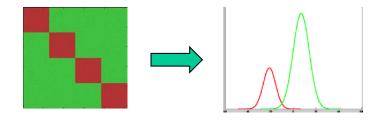
• Find model parameters that maximize expectation of log likelihood based on current estimate of density of hidden parameters



$$E[\log p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})|\mathcal{X}, \mathbf{\Theta}^t] = \sum_{\mathbf{y} \in \mathcal{Y}} \log p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta}) p(\mathbf{y}|\mathcal{X}, \mathbf{\Theta}^t)$$

$$Q(\mathbf{\Theta}, \mathbf{\Theta}^t) = \mathrm{E}[\log p(\mathcal{X}, \mathcal{Y}|\mathbf{\Theta})|\mathcal{X}, \mathbf{\Theta}^t]$$

$$\mathbf{\Theta}^{t+1} = \operatorname*{argmax}_{\mathbf{\Theta}} Q(\mathbf{\Theta}, \mathbf{\Theta}^t)$$



• After a long derivation, we can get analytical equations for our M-step if using GMMs:

$$\alpha_k^{t+1} = \frac{1}{N} \sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)$$

$$\mu_k^{t+1} = \frac{\sum_{i=1}^{N} x_i p(y_i = k | x_i, \mathbf{\Theta}^t)}{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)}$$

$$\sigma_k^{t+1} = \frac{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)(x_i - \mu_k^t)^2}{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)}$$

• The update equations are actually quite intuitive:

$$\alpha_k^{t+1} = \frac{1}{N} \sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)$$

• This represents the total relative weight of samples that are of class k (or gaussian component k), based on estimates of class labels (or hidden parameters), y_i

M-step

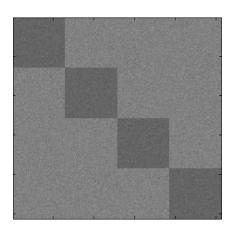
$$\mu_k^{t+1} = \frac{\sum_{i=1}^{N} x_i p(y_i = k | x_i, \mathbf{\Theta}^t)}{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)}$$

This represents the weighted sample mean of samples that are of class k (or gaussian component k), based on estimates of class labels (or hidden parameters), y_i

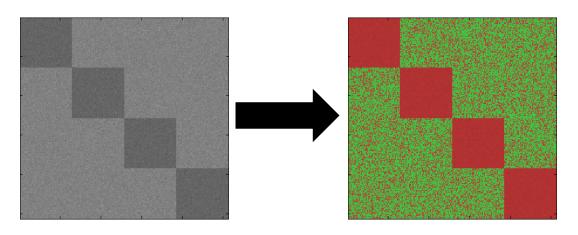
M-step

$$\sigma_k^{t+1} = \frac{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)(x_i - \mu_k^t)^2}{\sum_{i=1}^{N} p(y_i = k | x_i, \mathbf{\Theta}^t)}$$

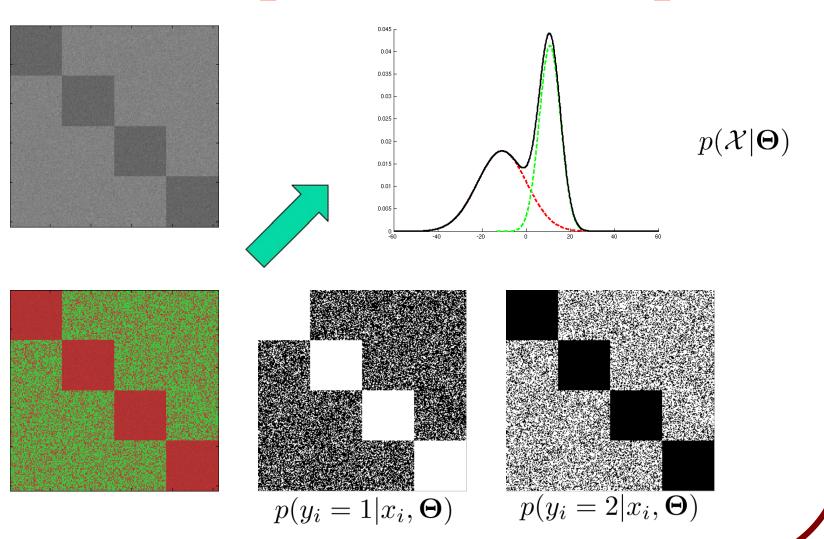
• This represents the weighted sample variance of samples that are of class k (or gaussian component k), based on estimates of class labels (or hidden parameters), y_i

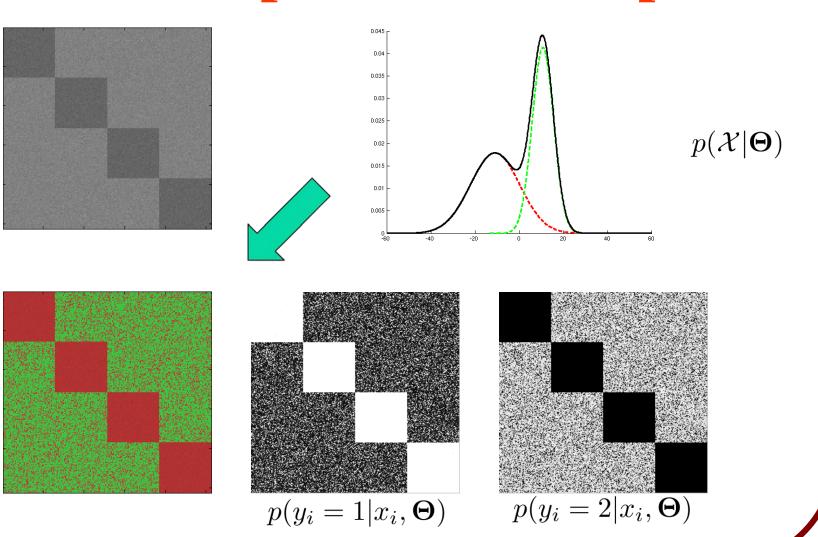


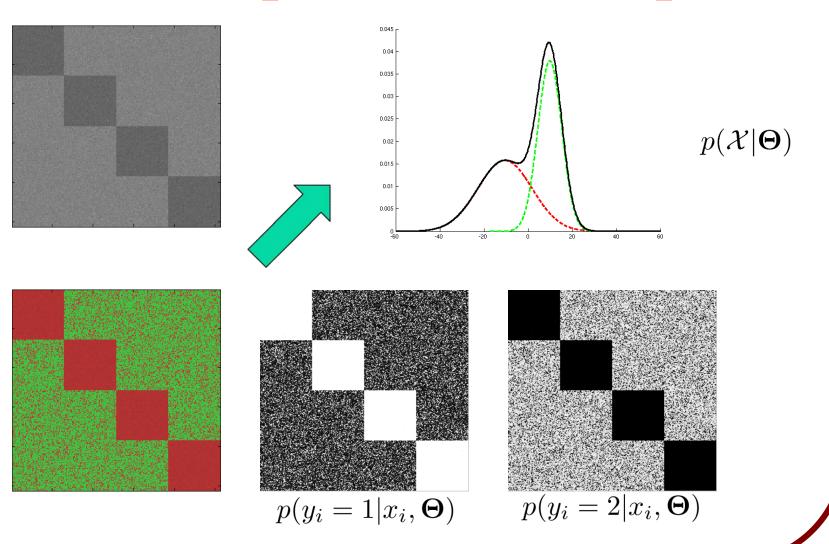
- Assume K = 2
- Initialize y_i such that samples with lowest 50% of intensities are assigned a label of 1, while highest 50% are assigned to label 2

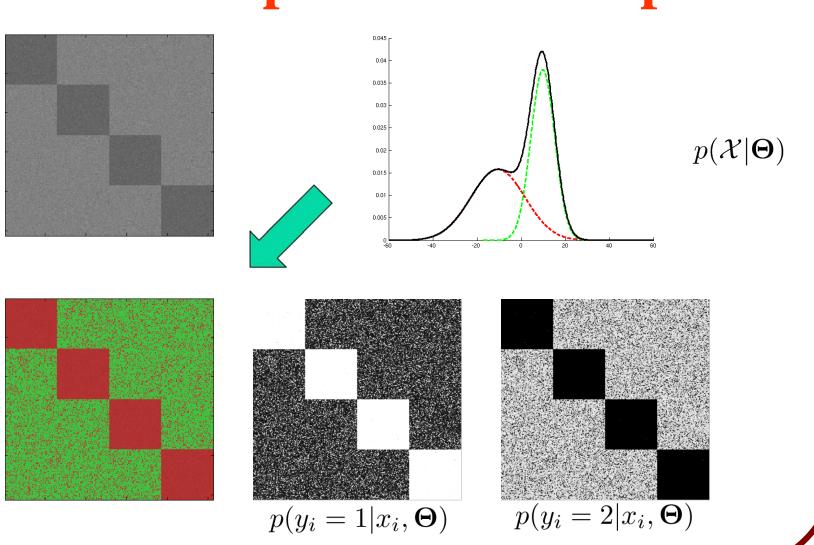


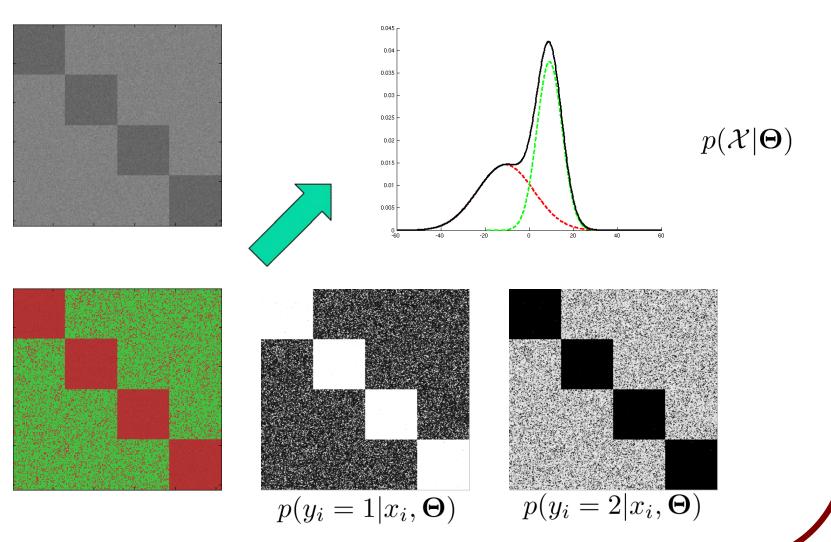
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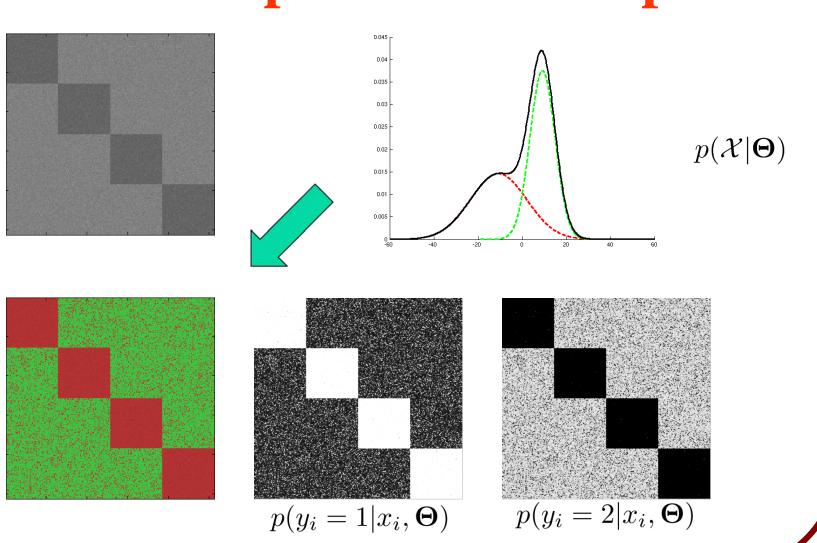


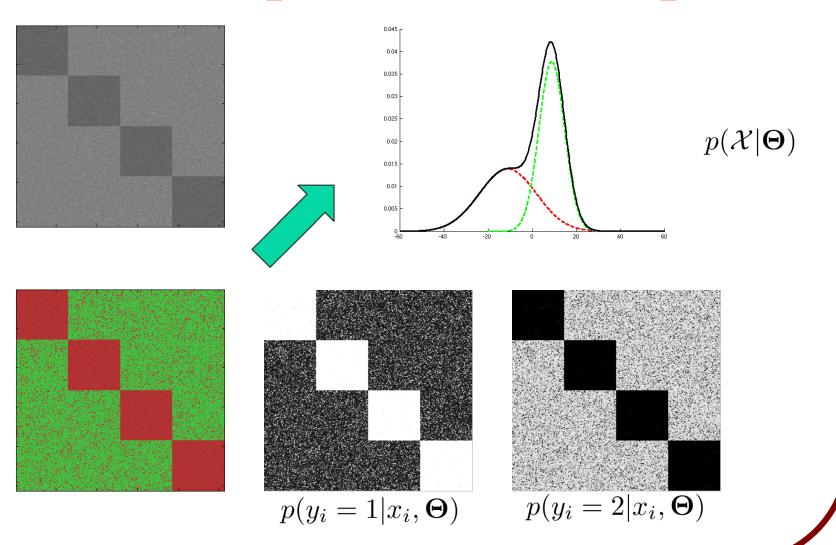


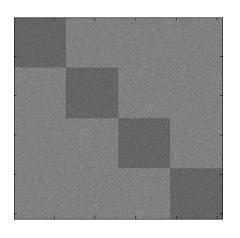


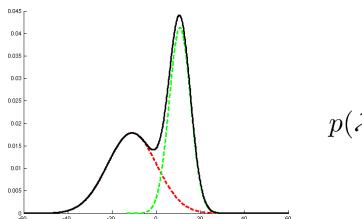




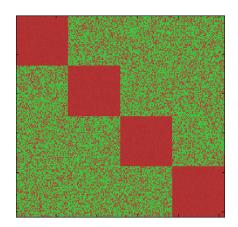


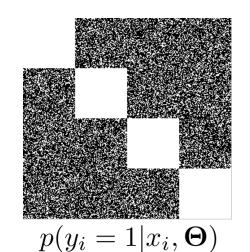


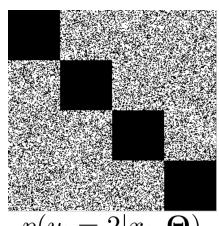


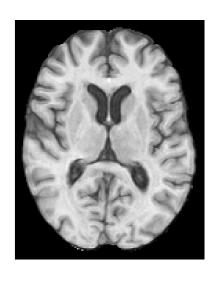


 $p(\mathcal{X}|\mathbf{\Theta})$

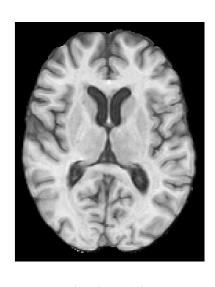




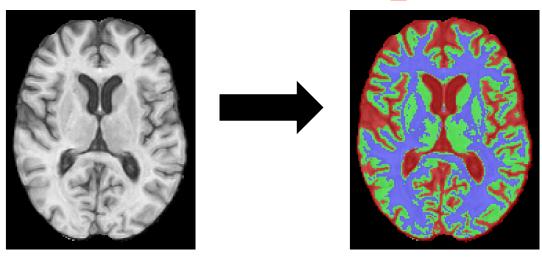




- Want to segment brain MRI into 3 tissue classes:
 - cerebro-spinal fluid (csf)
 - gray matter
 - white matter

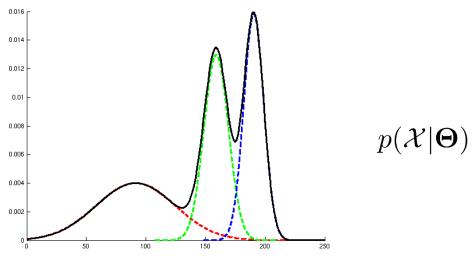


• Initialize y_i by assigning lowest 1/3 of intensities to csf, middle 1/3 to gray matter, and highest 1/3 to white matter

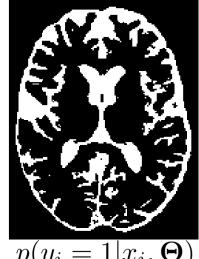


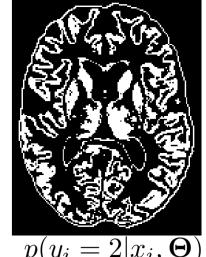
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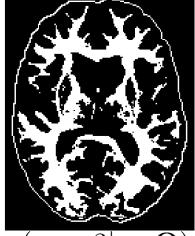






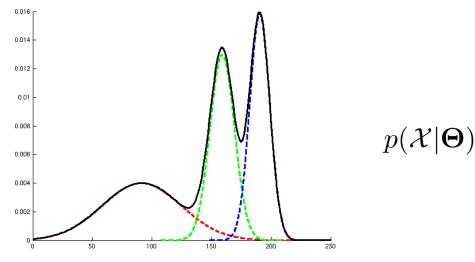


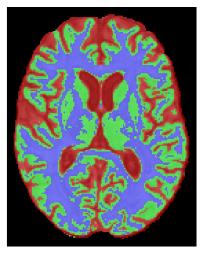


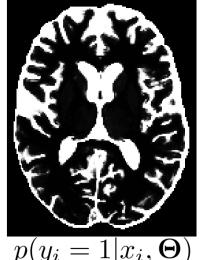


 $p(y_i = 1|x_i, \mathbf{\Theta})$ $p(y_i = 2|x_i, \mathbf{\Theta})$ $p(y_i = 3|x_i, \mathbf{\Theta})$

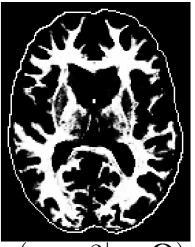




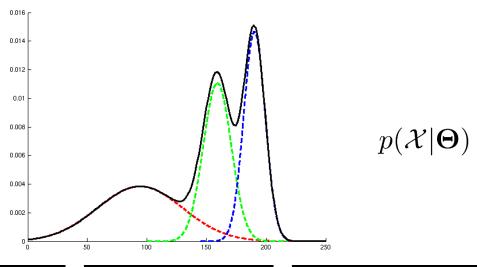


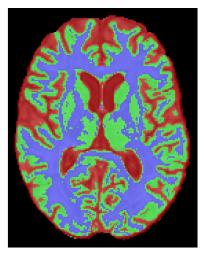










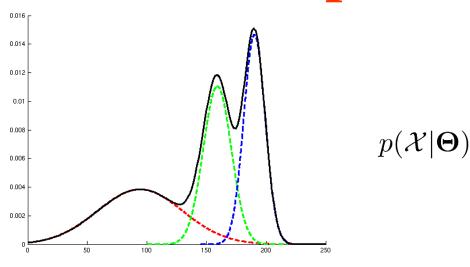










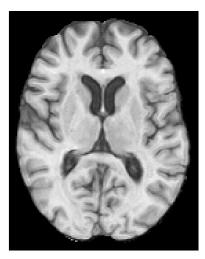


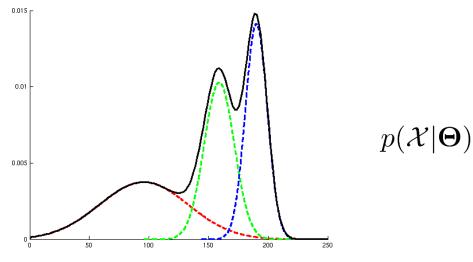


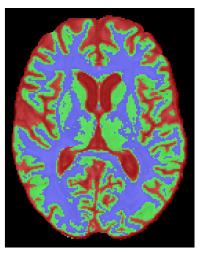


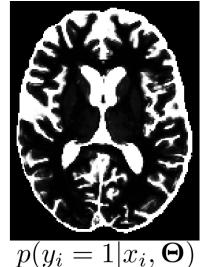


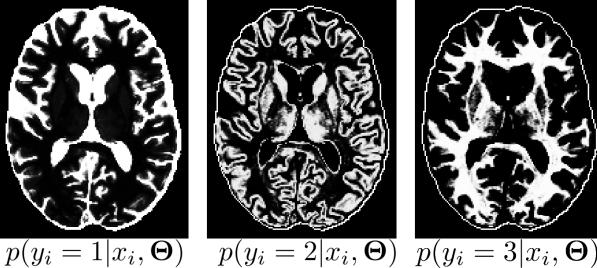






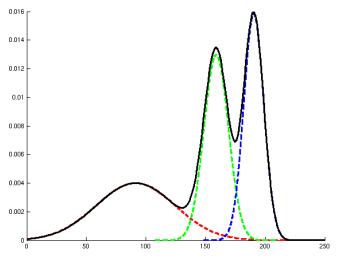






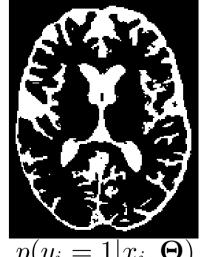


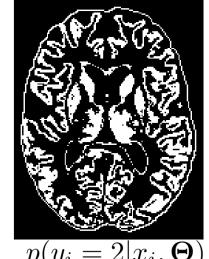


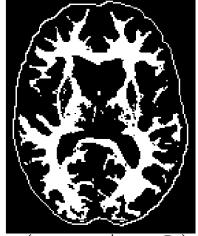


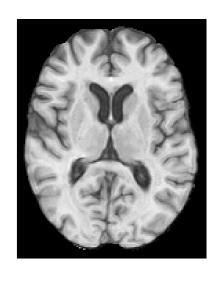
 $p(\mathcal{X}|\mathbf{\Theta})$



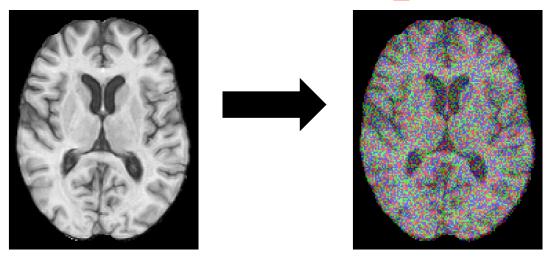








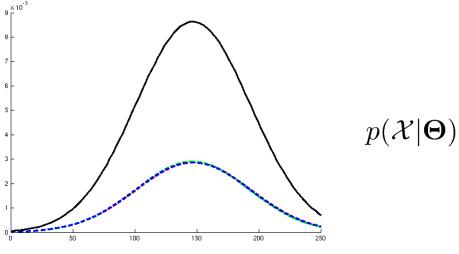
• What if we start with a random initialization?

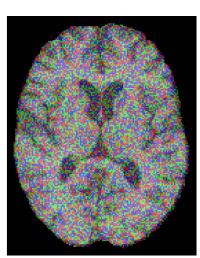


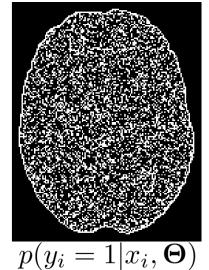
• What if we start with a random initialization?

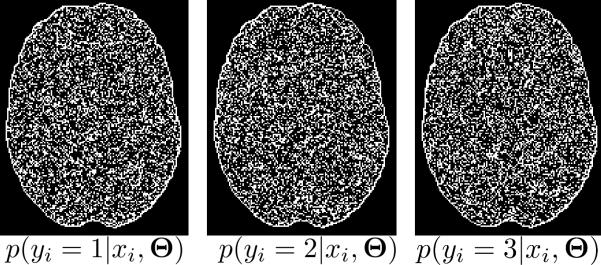


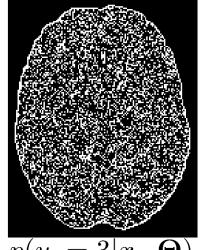


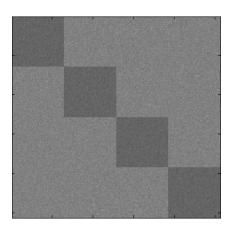




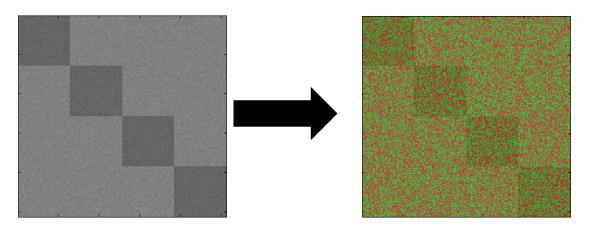




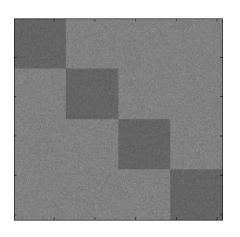


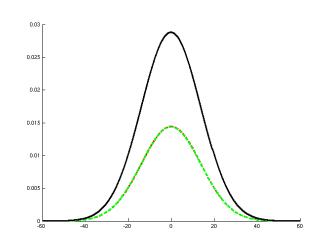


- Assume K = 2
- Random initialization

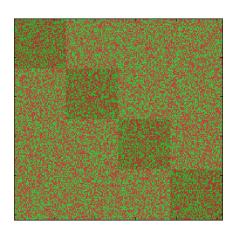


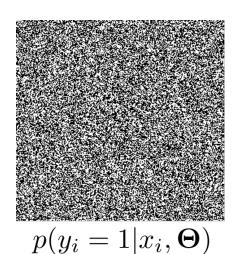
- Assume K = 2
- Random initialization

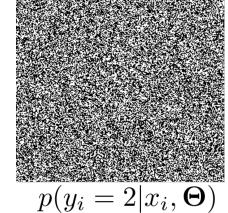


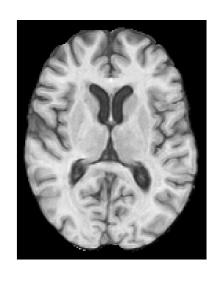


 $p(\mathcal{X}|\mathbf{\Theta})$

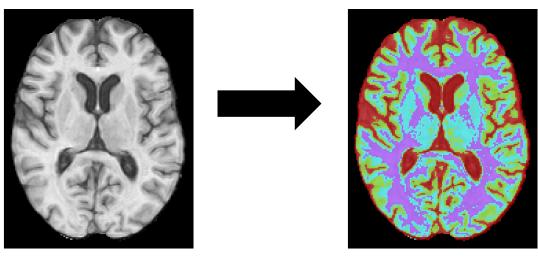




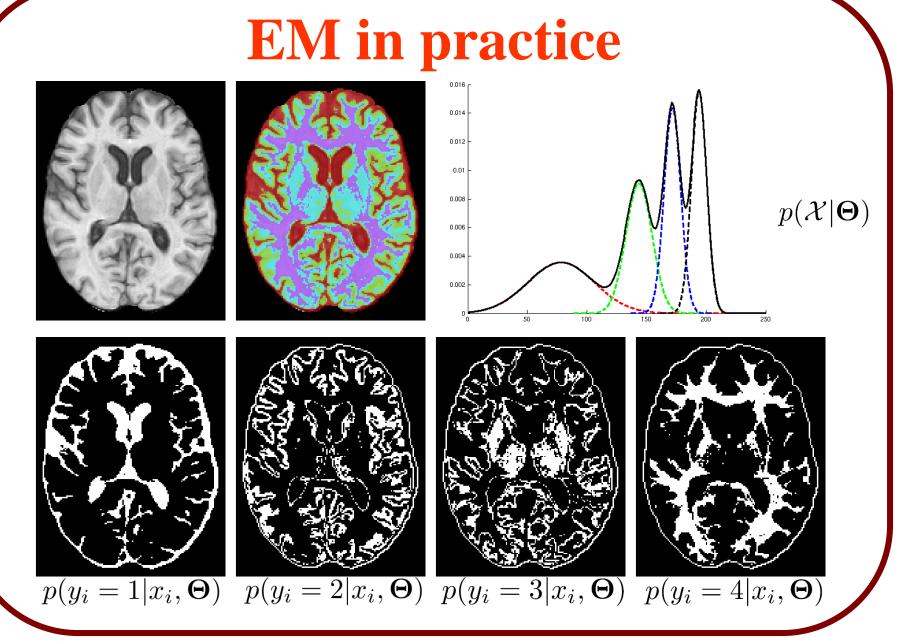


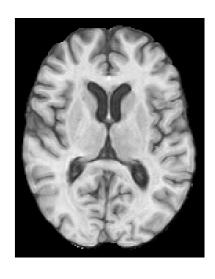


- What if we set K = 4?
- Initialize by setting lowest 25% of intensities to class 1, etc...

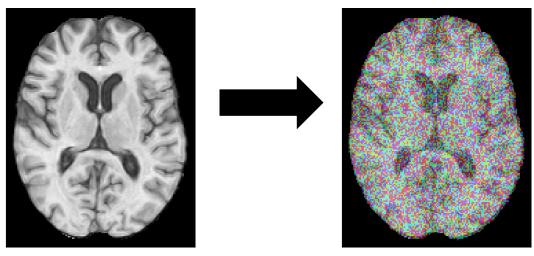


- What if we set K = 4?
- Initialize by setting lowest 25% of intensities to class 1, etc...

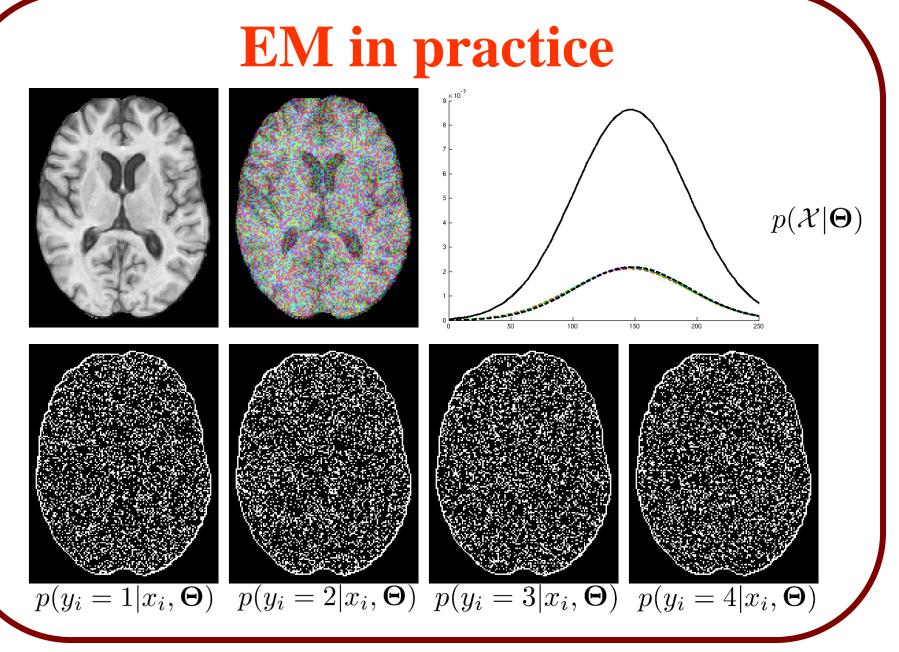




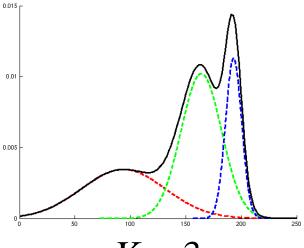
• What if we set K = 4 and initialize randomly?



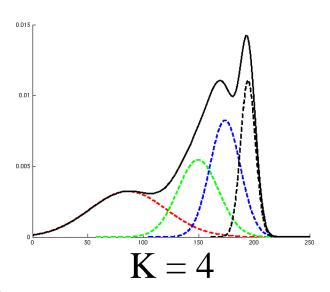
• What if we set K = 4 and initialize randomly?

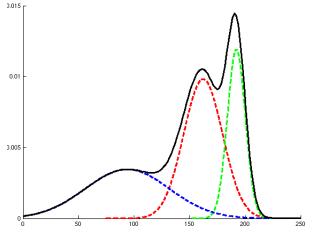


Effect of K and initialization

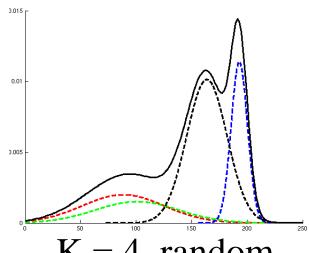


$$K = 3$$





K = 3, random



K = 4, random

Expectation Maximization

- Iterate until convergence guaranteed to converge to at least a local maximum
- More details on the derivation of the update equations (and EM in general) can be found here:

http://www.icsi.berkeley.edu/ftp/global/pub/techreports/1997/tr-97-021.pdf

GMMs / Expectation Maximization

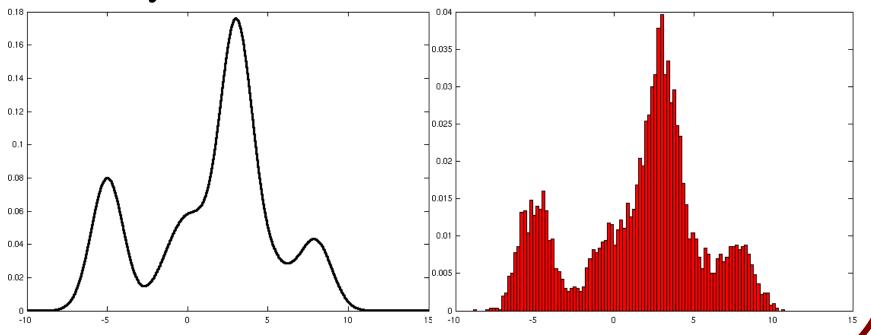
- GMM parameters provide multimodal estimate of a probability density from a set of samples
- Hidden parameters, y_i , can be treated as class labels if used in a segmentation or clustering framework and provide likelihood of class k at each sample, where class k is parametrized by the k^{th} Gaussian component of our GMM

GMMs

- Drawbacks
 - Need to know K or have some way to determine K
 - Final distribution depends on initialization of model parameters
 - Cannot approximate all arbitrary distributions with limited number of Gaussians
 - May take long time to converge
- Advantages
 - Parametric : still have relatively compact representation
 - Can approximate many multimodal distributions reasonably well with relatively small number of Gaussians
 - Relatively easy to implement

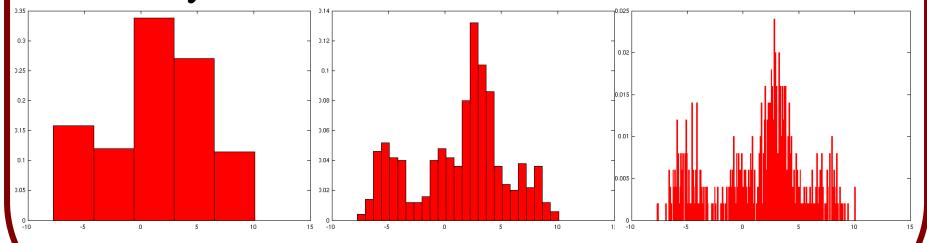
Non-parametric densities

 Non-parametric representations make no assumptions about the form of underlying density



Histogram

- Simple non-parametric representation
- Bins too large: loss of resolution
- Bins too small: zero probability "holes" in density



Kernel Density Estimation

- Method of building non-parametric representations of densities from data samples
- Densities represented as expected value of "smoothed" samples
- Also known as "Parzen Window" method

$$\hat{f}(x) = \frac{1}{N} \sum_{i=1}^{N} K(x - x_i)$$

Kernel Density Estimation

$$\hat{f}(x) = \frac{1}{N} \sum_{i=1}^{N} K(x - x_i)$$

• K(x) is kernel (or window) function

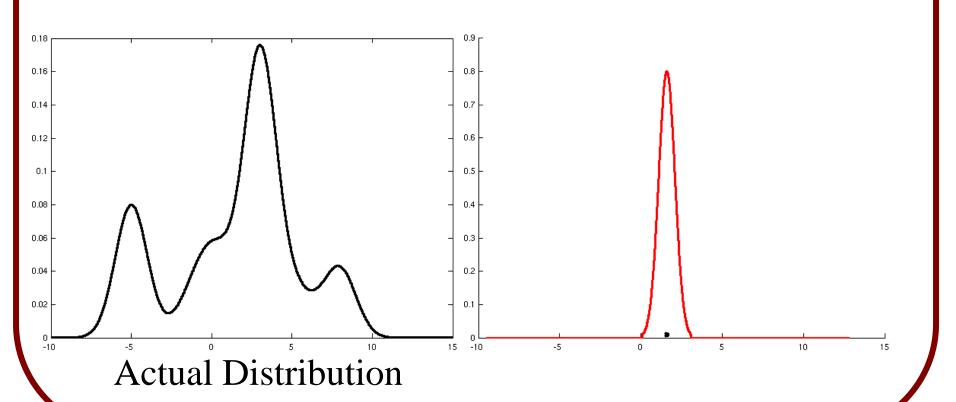
$$\int K(x) = 1$$

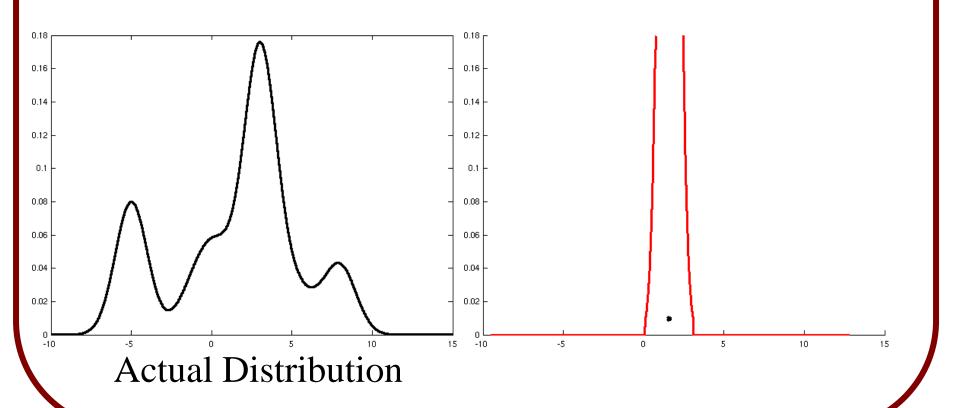
Kernel Density Estimation

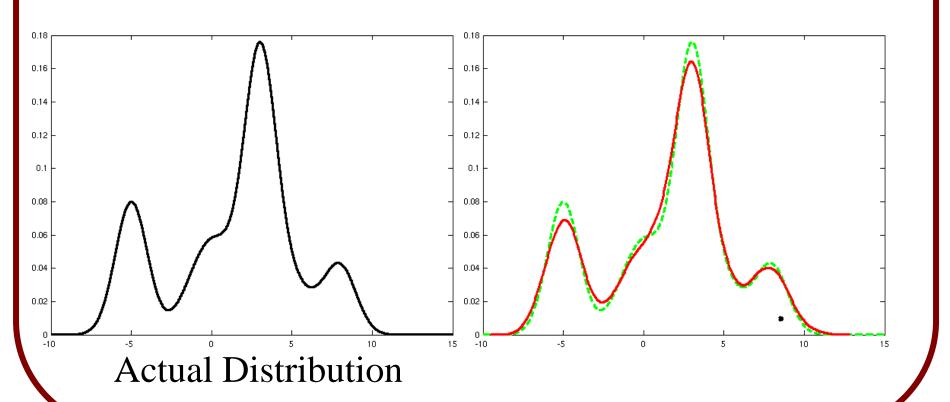
Gaussian is often used as smoothing kernel

$$\hat{f}(x) = \frac{1}{N} \sum_{i=1}^{N} \mathcal{N}(\mu = x_i, \sigma)$$

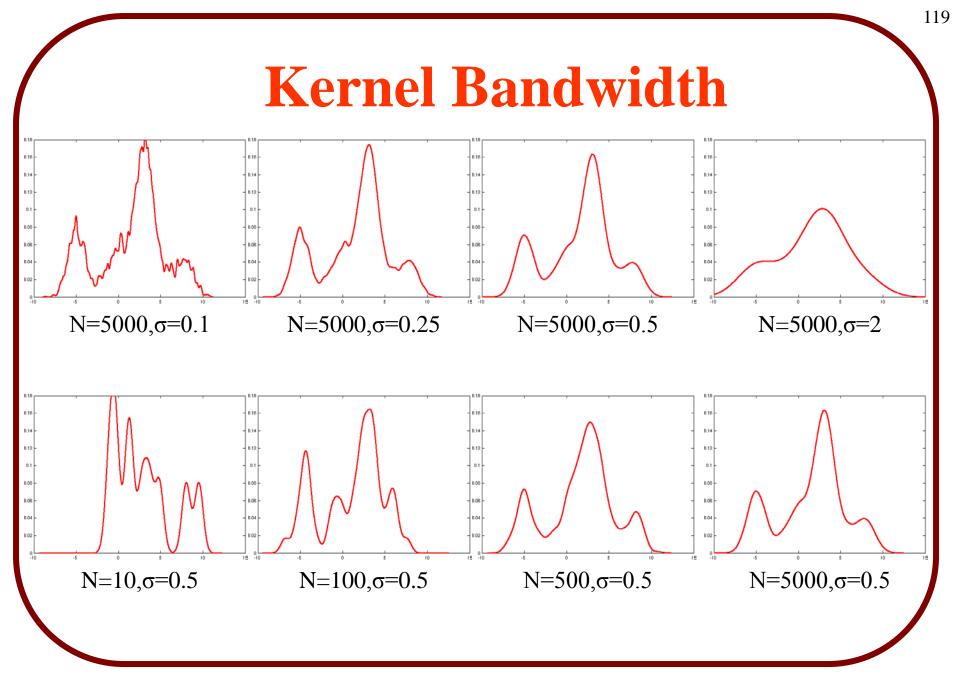
• Density represented as sum of Gaussians, where each Gaussian represents a data sample and has variance σ^2 and mean value equal to the value of the data sample.

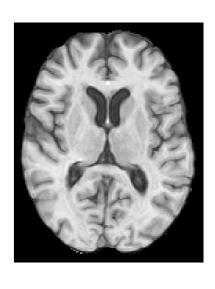


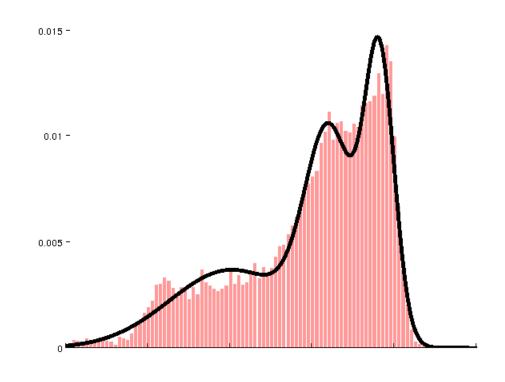




- Require bandwidth parameter (covariance of Gaussian kernel)
 - Determines "smoothness" of representation
 - Similar to bin width selection
 - Choice of kernel bandwidth generally based on #
 data samples and variance of data
 - Several methods exist to select bandwidth based on available sample data

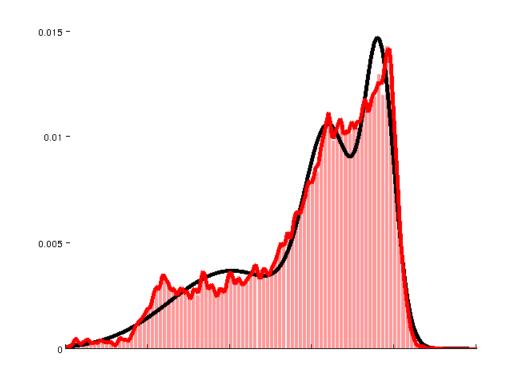




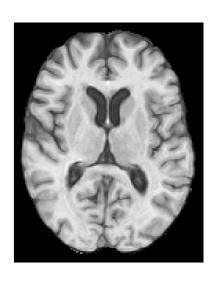


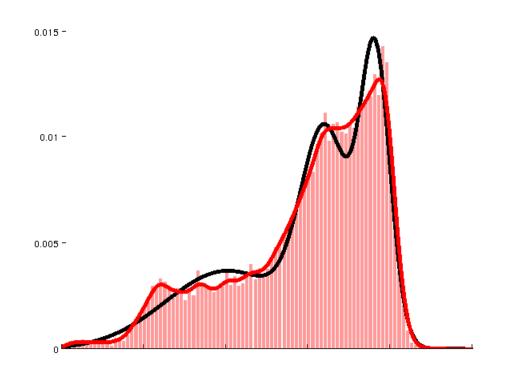
Black = 3 component GMM





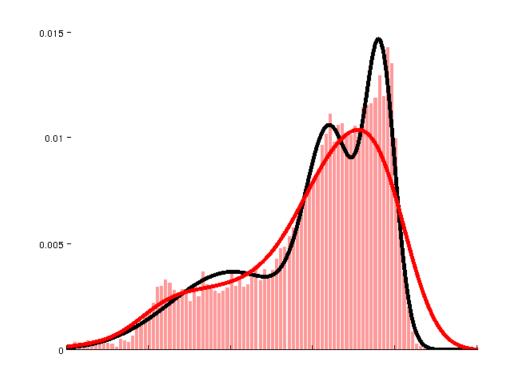
Black = 3 component GMM Red = Parzen Window (BW = 1)





Black = 3 component GMM Red = Parzen Window (BW = 4)





Black = 3 component GMM Red = Parzen Window (BW = 16)

Densities using Parzen Windows

- Advantages
 - Can represent arbitrary densities
 - Does not require initialization
 - Methods exist to select appropriate kernel bandwidth based on available data samples
- Disadvantages
 - Can not represent density in a compact form
 - Need to determine kernel bandwidth
 - Non-trivial implementation details, especially for higherdimensional densities

Summary

• GMMs

- Can approximate multi-modal densities in a relatively compact form.
- Given *K*, EM allows us to generate an estimate of the underlying probability density of our samples.
- For segmentation or clustering tasks, we can model our samples as a GMM and use EM to determine the underlying cluster from which each sample came.

- Can represent arbitrary probability densities without any initialization
- Non-compact representation
- More difficult to implement