### CS 561 Artificial Intelligence Lecture # 19-21

Learning Probabilistic Models

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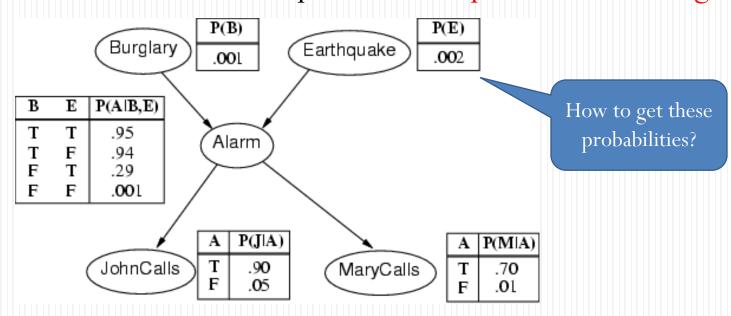
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#### Outline

- Learning Parameters of Bayesian Network
  - Maximum-likelihood approach
  - Bayesian approach
- Learning structure of Bayesian network
- Density estimation with non-parametric models
- Learning with Hidden variables
  - Expectation Maximization algorithm

### Learning Parameters of Bayes Net

- Density estimation: the general task of learning a probability model, given data (defined later) that are assumed to be generated from that model.
- The structure of the Bayesian network is fixed and the task is to learn the conditional probabilities- parameter learning



#### Parameter Learning

- Learning with complete data
  - Maximum-likelihood parameter learning
  - Bayesian parameter learning
- Learning with hidden variables
  - Expectation Maximization Algorithm

#### Parameter learning: Candy example

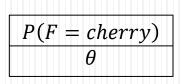
- Candy Example
  - Two flavours: cherry and lime
  - Candy wrapped in the same opaque paper, regardless of flavour
- Data: evidence instantiations of some or all of the random variables describing the domain
  - $D_i$ , i = 1: N represents data and  $D_i$  is a random variable with possible values cherry and lime
- Hypothesis: probabilistic theories of how the domain works
  - H: possible values  $h_1 \dots h_5$
  - $h_1$ : 100% cherry
  - $h_2$ : 75% cherry + 25% lime
  - ....
  - $h_5$ : 100% lime



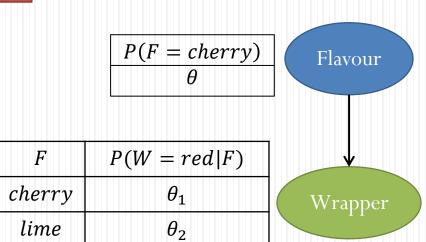
#### Parameter Learning: Candy Example

 We assume that the observations are complete; i.e., each data point contains values for every variable in the probability model being learned.

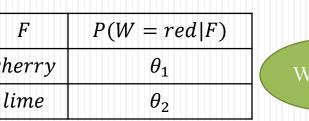




Flavour



P(F = cherry)



## Maximum-likelihood parameter learning – Discrete models

- Given the candy bag, what is the lime-cherry proportion?
  - It (hypothesis) could be anywhere between 0 and 1.
  - ullet heta, proportion of cherry candies and  $h_{ heta}$  is the hypothesis
- Suppose we unwrap *N* candies:
  - c are cherries and l = N c are limes
- Likelihood of this dataset is (assuming i.i.d.):

$$P(\mathbf{d}|h_{\theta}) = \prod_{j=1}^{N} P(d_j|h_{\theta}) = \theta^{c}.(1-\theta)^{l}$$

P(F = cherry)  $\theta$ 



• Maximum-likelihood hypothesis is given by the value of  $\theta$  that maximizes  $P(\mathbf{d}|h_{\theta})$  (easier with  $\log$  likelihood)

## Maximum-likelihood parameter learning – Discrete models

#### Multiple parameters

Red/green wrapper depends probabilistically on flavor:

Likelihood for, e.g., cherry candy in green wrapper:

$$P(F = cherry, W = green | h_{\theta,\theta_1,\theta_2})$$

$$= P(F = cherry | h_{\theta,\theta_1,\theta_2})P(W = green | F = cherry, h_{\theta,\theta_1,\theta_2})$$

$$= \theta \cdot (1 - \theta_1)$$

N candies,  $r_c$  red-wrapped cherry candies, etc.:

$$P(\mathbf{d}|h_{\theta,\theta_1,\theta_2}) = \theta^c(1-\theta)^{\ell} \cdot \theta_1^{r_c}(1-\theta_1)^{g_c} \cdot \theta_2^{r_{\ell}}(1-\theta_2)^{g_{\ell}}$$

$$L = [c \log \theta + \ell \log(1 - \theta)]$$
  
+  $[r_c \log \theta_1 + g_c \log(1 - \theta_1)]$   
+  $[r_\ell \log \theta_2 + g_\ell \log(1 - \theta_2)]$ 

(from text book website)

Flavor

Wrapper

P(W=red)

 $\theta_2$ 

## Maximum-likelihood parameter learning – Discrete models

#### Multiple parameters contd.

Derivatives of L contain only the relevant parameter:

$$\frac{\partial L}{\partial \theta} = \frac{c}{\theta} - \frac{\ell}{1 - \theta} = 0 \qquad \Rightarrow \quad \theta = \frac{c}{c + \ell}$$

$$\frac{\partial L}{\partial \theta_1} = \frac{r_c}{\theta_1} - \frac{g_c}{1 - \theta_1} = 0 \qquad \Rightarrow \quad \theta_1 = \frac{r_c}{r_c + g_c}$$

$$\frac{\partial L}{\partial \theta_2} = \frac{r_\ell}{\theta_2} - \frac{g_\ell}{1 - \theta_2} = 0 \qquad \Rightarrow \quad \theta_2 = \frac{r_\ell}{r_\ell + g_\ell}$$

With complete data, parameters can be learned separately

• Learning the parameters of a Gaussian density function on a single variable:

$$P(x) = \frac{1}{\sqrt{2\pi\sigma}} e^{\frac{-(x-\mu)^2}{2\sigma^2}}$$

Parameters for this model:  $\mu$  (mean) and  $\sigma$  (standard deviation)

• for observed values:  $x_1, ..., x_N$ , the log likelihood can be given as

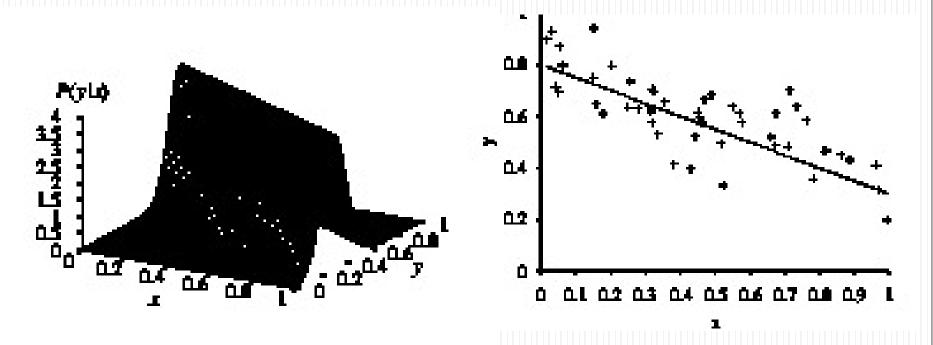
$$L = \sum_{j=1}^{N} \log \frac{1}{\sqrt{2\pi\sigma}} e^{\frac{-(x_j - \mu)^2}{2\sigma^2}} = N(-\log \sqrt{2\pi} - \log \sigma) - \sum_{j=1}^{N} \frac{(x_j - \mu)^2}{2\sigma^2}$$

Taking the derivative and setting to zero,

$$\frac{\partial L}{\partial \mu} = -\frac{1}{\sigma^2} \sum_{j=1}^{N} (x_j - \mu) = 0 \implies \mu = \frac{\sum_j x_j}{N} \frac{\partial L}{\partial \sigma} = -\frac{N}{\sigma} + \frac{1}{\sigma^3} \sum_{j=1}^{N} (x_j - \mu)^2$$

$$\mu^2 = 0 \implies \sigma = \sqrt{\frac{\sum_j (x_j - \mu)^2}{N}}$$

• Maximum-likelihood value of the mean is the sample average and the maximum-likelihood value of the standard deviation is the square root of the sample variance.



- Linear Gaussian Model with one continuous parent X and a continuous childY.
- Y has Gaussian distribution, mean depends linearly on X and standard deviation is fixed.

• To learn the conditional distribution P(Y|X), maximize the conditional likelihood

$$P(y|x) = \frac{1}{\sqrt{2\pi\sigma}} e^{\frac{-(y-(\theta_1x+\theta_2))^2}{2\sigma^2}}$$

Parameters for this model:  $heta_1$  and  $heta_2$  and  $\sigma$ 

- For observations,  $(x_1, y_1), (x_2, y_2), ... (x_N, y_N)$ 
  - maximizing P(y|x) w.r.t.  $\theta_1$  and  $\theta_2$  is same as minimizing the numerator  $(y (\theta_1 x + \theta_2))^2$  i.e. minimizing the sum of squared errors

$$E = \sum_{j=1}^{N} (y_j - (\theta_1 x_j + \theta_2))^2$$

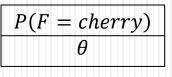
minimizing the sum of squared errors gives the ML solution for a linear fit assuming Gaussian noise of fixed variance

# Maximum-likelihood parameter learning

- ML parameter learning approach
  - Write down an expression for the likelihood of the data as a function of the parameter(s).
  - Write down the derivative of the log likelihood with respect to each parameter.
  - Find the parameter values such that the derivatives are zero.
- Maximum-likelihood learning has deficiencies with small datasets.

### Bayesian Parameter Learning

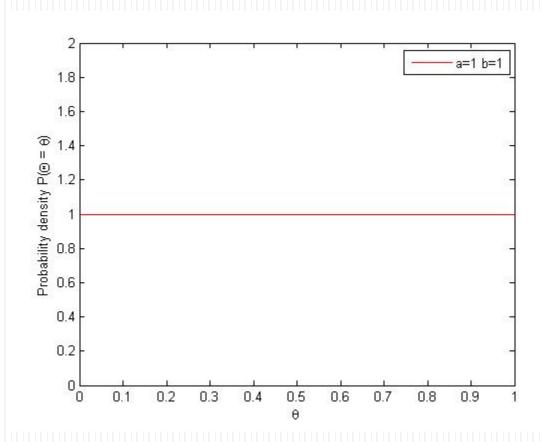
- Bayesian approach to parameter learning
  - define a prior probability distribution over the possible hypothesis (hypothesis prior)
  - as data arrives, update the posterior probability distribution
- Bayesian view:  $\theta$  is the (unknown) value of a random variable  $\Theta$  that defines the hypothesis space: the hypothesis prior is the prior distribution  $P(\Theta)$
- $P(\Theta = \theta)$  is the prior probability that the bag has a fraction  $\theta$  of cherry candies
- $\theta$  can be any value between 0 and 1, then  $P(\theta)$  must be a continuous distribution that is non-zero only between 0 and 1 and that integrates to 1.

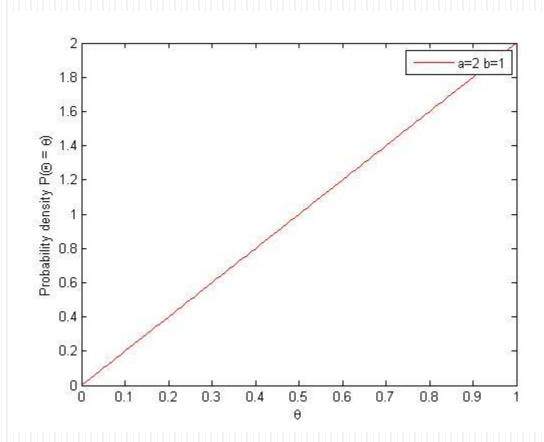


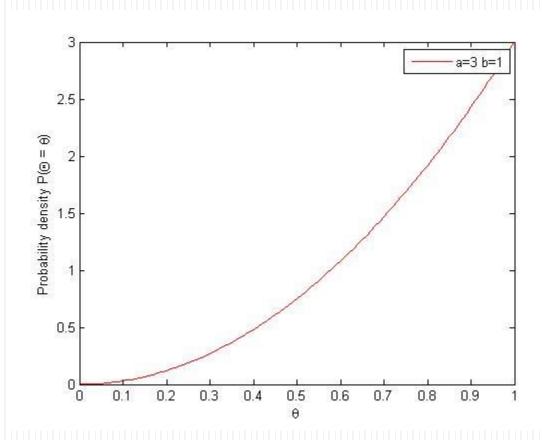


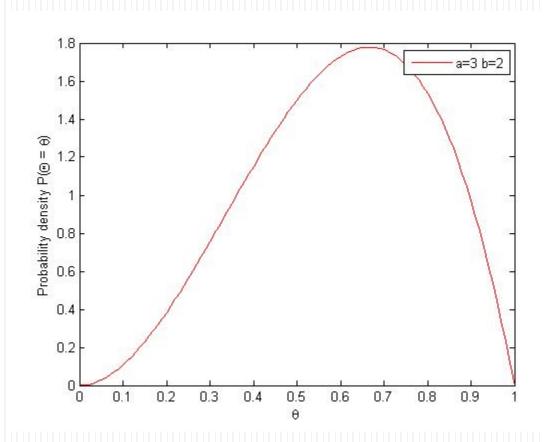
### Bayesian Parameter Learning

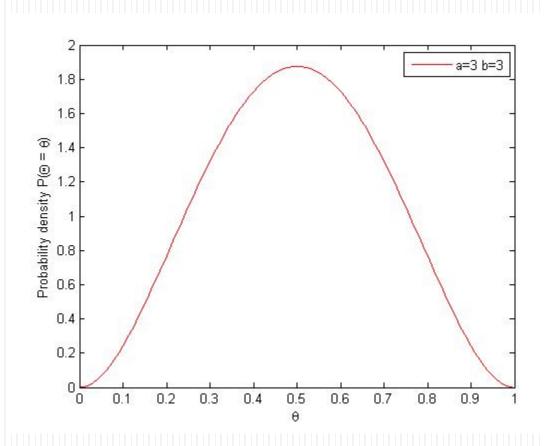
- $P(\theta) = \text{beta distribution}$ : candidate prior probability
  - beta $[a,b](\theta) = \alpha \theta^{a-1} (1-\theta)^{b-1}$  for  $\theta$  in the range [0,1]
  - Mean of the distribution  $\frac{a}{a+b}$
  - Larger values of a suggest a belief that  $\Theta$  is closer to 1 than to 0
  - Larger values of a + b make the distribution more peaked
  - beta distribution is closed under update
- Suppose we observe cherry candy

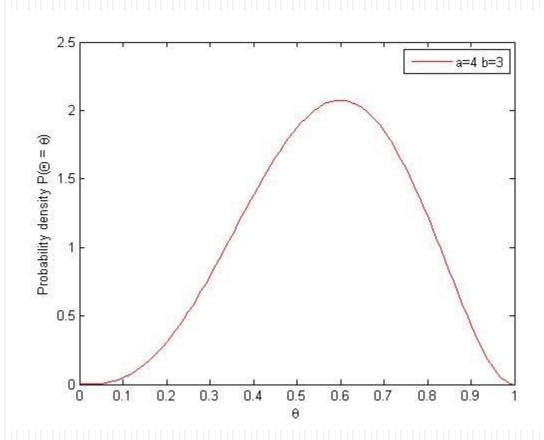


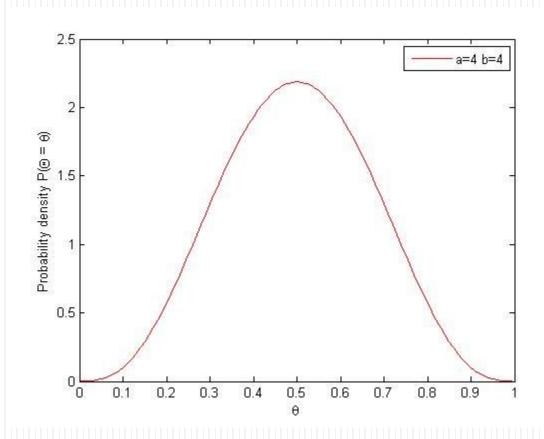


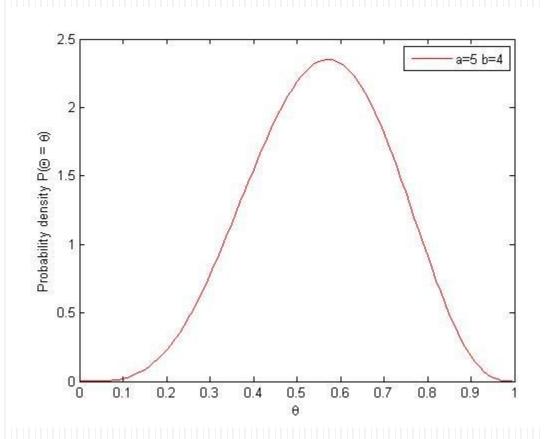


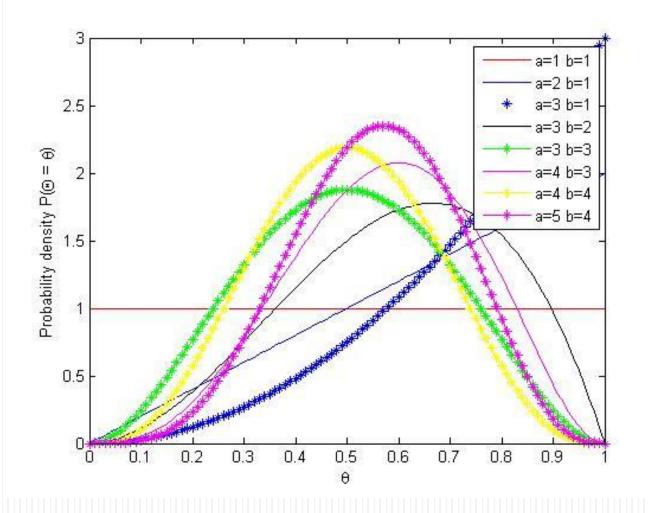












### Density estimation using Nonparametric models

- Density estimation (repeated from slide # 3)
  - general task of learning a probability model, given data that are assumed to be generated from that model

#### Background

• Continuous random variable *X* can take an infinite number of possible values, the probability density function describes the relative likelihood that the random variable takes on a given value.

$$P(a \le X \le b) = \int_{a}^{b} p(x)$$

We are interested in construction of estimate of the density function from the observed data

• For a Discrete random variable, X the probability distribution is given by a list of probabilities associated with each possible outcome, i.e.  $P(x_i) = P(X = x_i)$ , which satisfies the conditions:

$$\sum_{i=1}^{n} P(x_i) = 1 \text{ and } 0 \le P(x_i) \le 1$$

### Parametric approach vs. Nonparametric approach

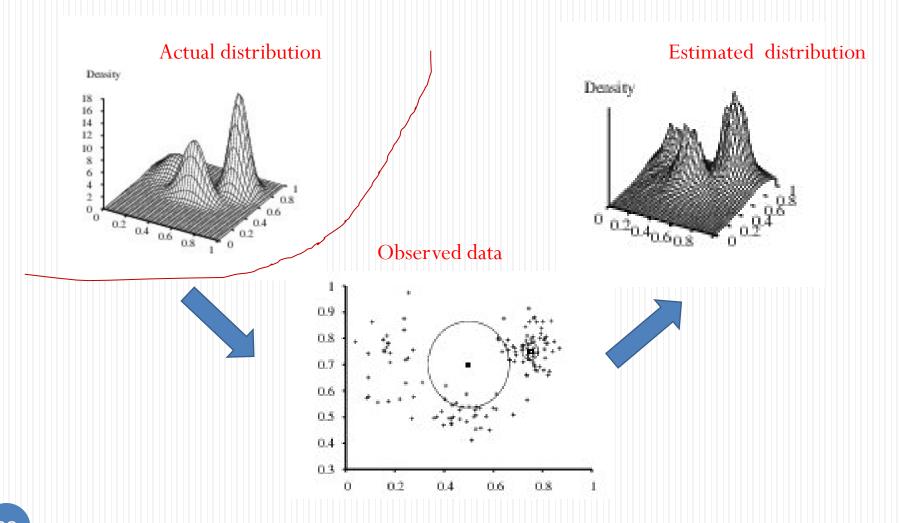
#### • Parametric approach

- assumes that the data are drawn from one of a known parametric family of distributions, for example the normal distribution with mean  $\mu$  and variance  $\sigma^2$
- density p underlying the data could then be estimated by finding estimates of  $\mu$  and  $\sigma^2$  from the data and substituting these estimates into the formula for the normal density.

#### Non-parametric approach

- No assumptions about the distribution of data is made
- ullet Data will speak for themselves in determining the estimate of density p

### Non-parametric density estimation



#### Non-parametric density estimation

- Histogram simplest form of non-parametric density estimation
- Kernel density estimation
- k-nearest neighbour approach

### Non-parametric density estimation

#### Histogram

- sample space is divided into a number of bins and density is approximated at the centre of each bin by the fraction of points in the observed data that fall into corresponding bin.
- Given an origin  $x_0$  and a bin width h, the bins of the histogram are intervals  $[x_0 + mh, x_0 + (m+1)h]$  for positive and negative integer m.
- Histogram is then defined by
  - $\hat{p}(x) = \frac{1}{nh} (\# X_i \text{ in same bin as } x)$
  - Generally,  $\hat{p}(x) = \frac{1}{n} \frac{(\# X_i \text{ in same bin as } x)}{(\text{width of bin containing } x)}$

#### Histogram

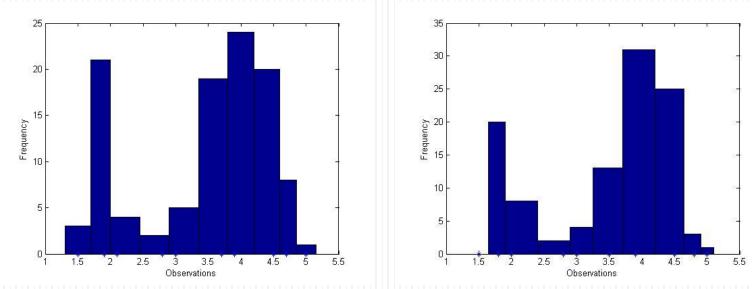


Figure 1. Histograms for the same data with different origins

- histogram requires bin width and origin to be defined
- final shape of the density estimate depends on the origin
- histogram unsuitable for most practical applications, useful for quick visualization of results in one or two dimensions

### Kernel density estimate

- Basic idea: take the weighted local density estimates at each observation  $x_i$  and then aggregate them to yield an overall density.
- Consider a vector  ${\bf x}$  , the probability P that a vector falls in a region  ${\mathcal R}$  can be given as

$$P = \int_{\mathcal{R}} p(\mathbf{x}) dx$$

• Assuming that  $\mathcal{R}$  is very small such that  $p(\mathbf{x})$  does not vary much within it, we can write

$$P = \int_{\mathcal{R}} p(\mathbf{x}) dx \approx p(\mathbf{x}) \int_{\mathcal{R}} dx = p(x)V$$

where V is the "volume" of region  ${\cal R}$ 

#### Kernel density estimate

• Suppose that n samples  $\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n$  are independently drawn according to the probability density function  $p(\mathbf{x})$ , and there are k out of n samples falling within the region  $\mathcal{R}$ , we have

$$P = k/n$$

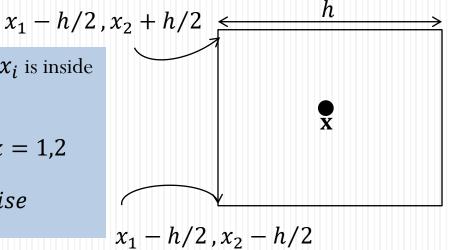
• Thus, the estimate of  $p(\mathbf{x})$  can be given as

$$\hat{p}(\mathbf{x}) = \frac{\kappa}{n} / V$$

• Consider that  $\mathcal{R}$  is a square with length of edge = h

Let us define a function that indicates whether  $\boldsymbol{x_i}$  is inside the square or not.

$$\emptyset\left(\frac{\mathbf{x}_{i} - \mathbf{x}}{h}\right) = \begin{cases} 1 & \frac{|x_{ik} - x_{k}|}{h} \le 1/2, k = 1, 2\\ 0 & otherwise \end{cases}$$



### Kernel density estimate (KDE)

• # samples within the region  $\mathcal{R}$  out of n samples is given by

$$k = \sum_{i=1}^{n} \emptyset\left(\frac{\mathbf{x}_{i} - \mathbf{x}}{h}\right)$$

• The density estimate for 2-D is given by

$$\hat{p}(\mathbf{x}) = \frac{k/n}{V} = \frac{1}{n} \sum_{i=1}^{n} \frac{1}{h^2} \emptyset \left(\frac{\mathbf{x}_i - \mathbf{x}}{h}\right)$$

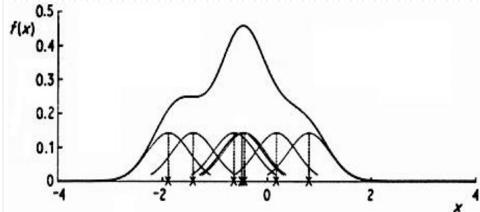
Kernel (or window) function)

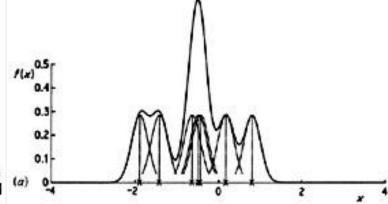
where *h* is the window width, also called the smoothing parameter or bandwidth.

• For d-dimension

$$\hat{p}(\mathbf{x}) = \frac{k/n}{V} = \frac{1}{n} \sum_{i=1}^{n} \frac{1}{h^d} \emptyset\left(\frac{\mathbf{x}_i - \mathbf{x}}{h}\right)$$

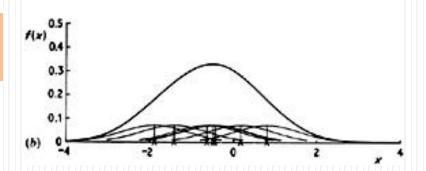
#### Kernel density estimate





1-D Gaussian function is used here for 1-D with various values of bandwidth parameter.

$$\hat{p}(x) = \frac{1}{n} \sum_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma}} exp\left(-\frac{(x_i - x)^2}{2\sigma^2}\right)$$



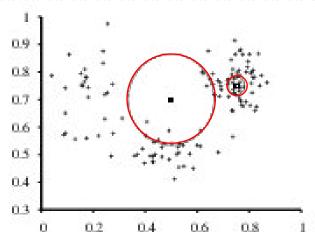
#### k-nearest neighbour

• Recall the generic expression for density estimation

$$\hat{p}(\mathbf{x}) = \frac{k}{n} / V$$

- In KDE, V is fixed and that determines k, the number of samples inside V.
- In k-nearest neighbour approach, k is fixed and we find the V that contains k points inside.

10 nearest neighbours of square points



Bayesian Belief Network Training

### Algorithm 3 PC algorithm for skeleton learning.

```
1: Start with a complete undirected graph G on the set \mathcal V of all vertices.

2: i=0

3: repeat

4: for x \in \mathcal V do

5: for y \in Adj \{x\} do

Determine if there a subset \mathcal S of size i of the neighbours of x (not including y) for which x \perp \!\!\! \perp y \mid \mathcal S. If this set exists remove the x-y link from the graph G and set \mathcal S_{xy} = \mathcal S.

7: end for

8: end for

9: i=i+1.

10: until all nodes have \leq i neighbours.
```

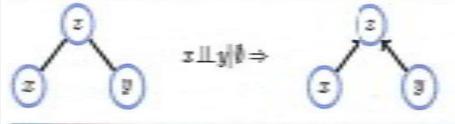
From the Book: Bayesian Reasoning and Machine Learning, David Barber (Chapter 9)

Bayesian Belief Network Training

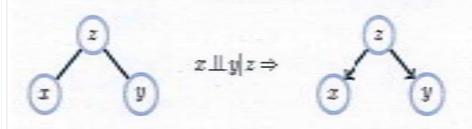
### Algorithm 4 Skeleton orientation algorithm (returns a DAG).

- 1: Unmarried Collider: Examine all undirected links x-z-y. If  $z \notin S_{xy}$  set  $x \to z \leftarrow y$ .
- 2: repeat
- 3:  $x \rightarrow z y \Rightarrow x \rightarrow z \rightarrow y$
- 4: For x-y, if there is a directed path from x to y orient  $x \to y$
- 5: If for x-z-y there is a w such that  $x\to w$ ,  $y\to w$ , z-w then orient  $z\to w$
- 6: until No more edges can be oriented.
- 7: The remaining edges can be arbitrarily oriented provided that the graph remains a DAG and no additional colliders are introduced.

### Example 41 (Skeleton orienting).

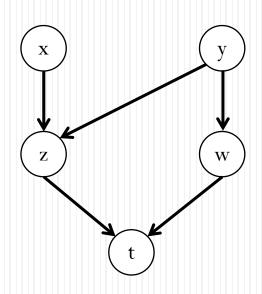


If x is (unconditionally) independent of y, it must be that z is a collider since otherwise marginalising over z would introduce a dependence between x and y.

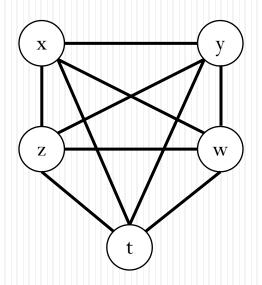


If x is independent of y conditioned on z, z must not be a collider. Any other orientation is appropriate.

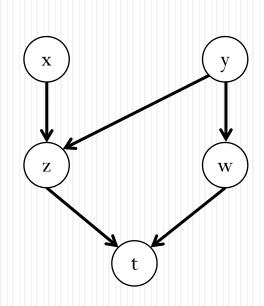
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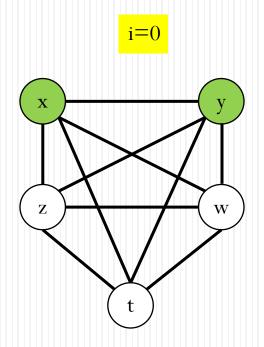
Bayesian Network from which data is assumed to be generated and against which conditional independence tests will be performed.



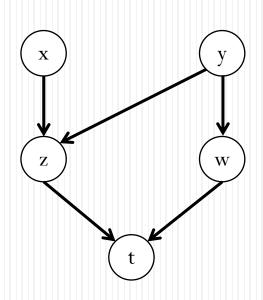
We start with a structure (skeleton) that is fully connected.



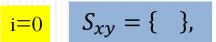
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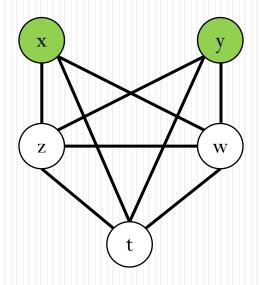


i=0, size of subset S = 0 for which  $x \perp y \mid S$ 

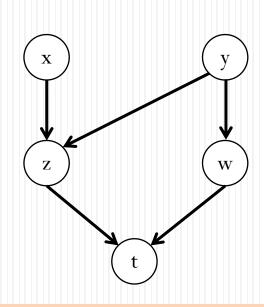


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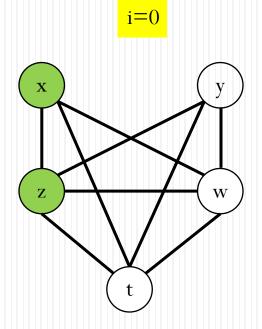




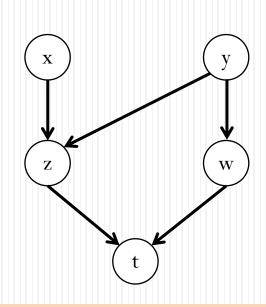
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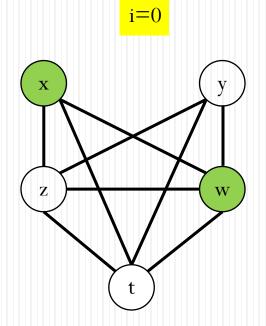
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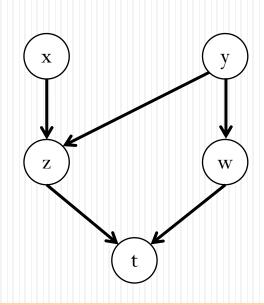
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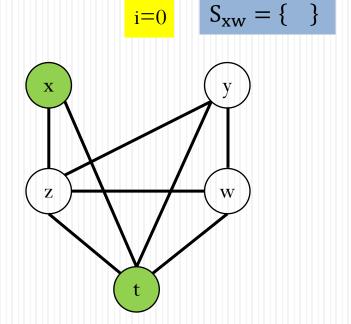
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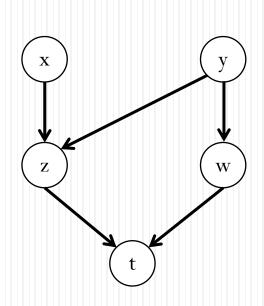
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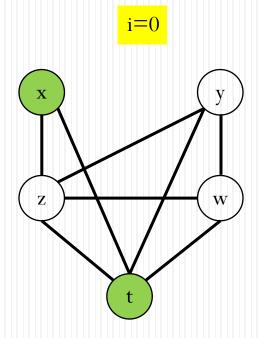
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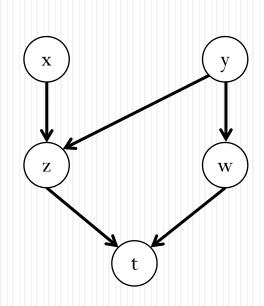
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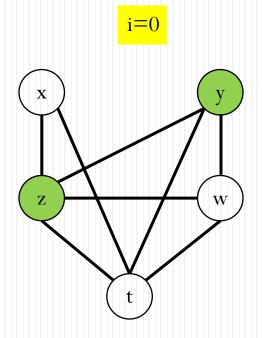
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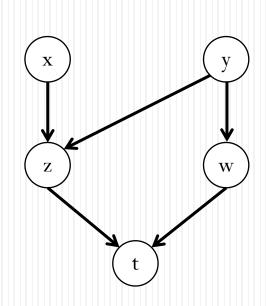
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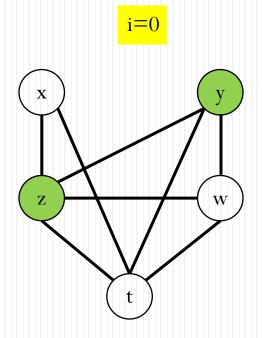
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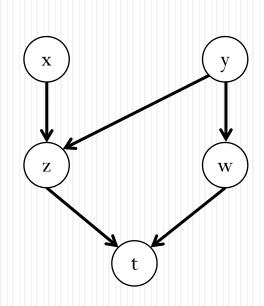
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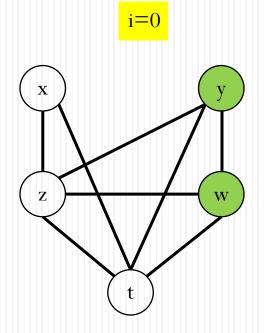
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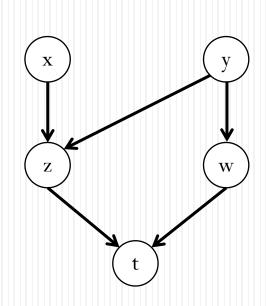
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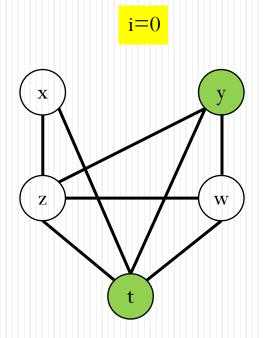
Bayesian Network from which data is assumed to be generated and against which conditional independence tests will be performed.



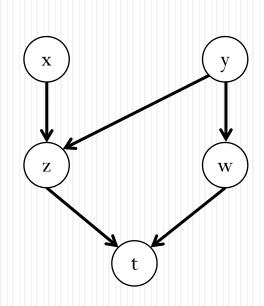
i=0, size of subset S = 0 for which  $x \perp y \mid S$ 



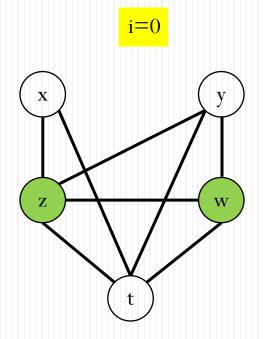
Bayesian Network from which data is assumed to be generated and against which conditional independence tests will be performed.



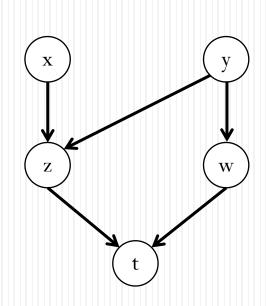
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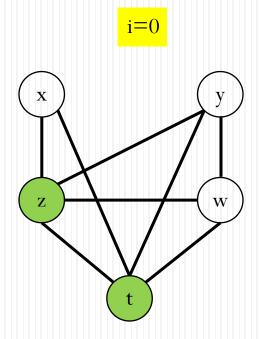
Bayesian Network from which data is assumed to be generated and against which conditional independence tests will be performed.



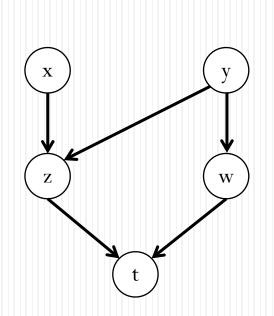
i=0, size of subset S = 0 for which  $x \perp y \mid S$ 

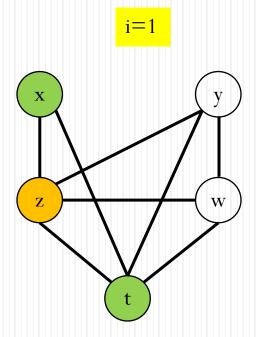


Bayesian Network from which data is assumed to be generated and against which conditional independence tests will be performed.

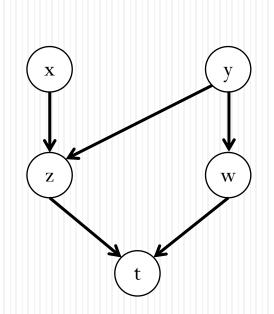


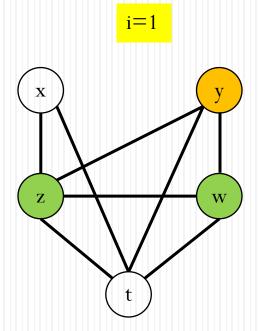
i=0, size of subset S = 0 for which  $x \perp y \mid S$ 



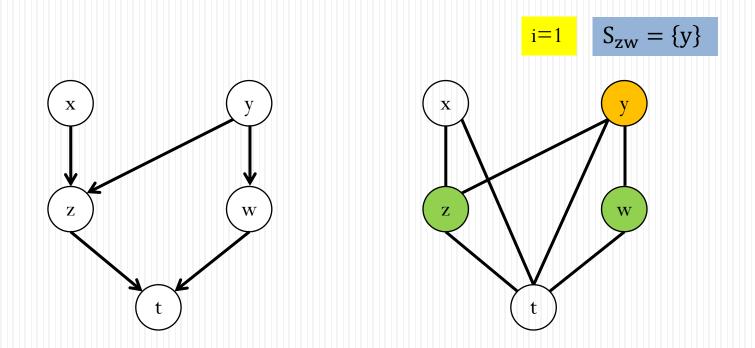


i=1, size of subset S=1 for which  $x\perp y|S$  or  $x\perp y|z$ ,  $z\in S_{xy}$  we test all pairs x-y for independence, conditioned on single neighbour variable z and for such pairs the link is removed.

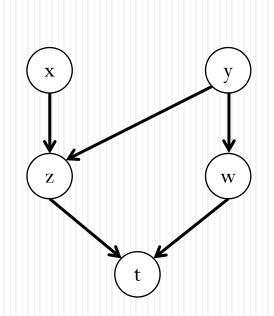


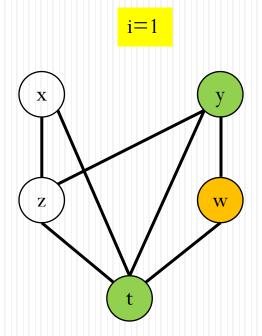


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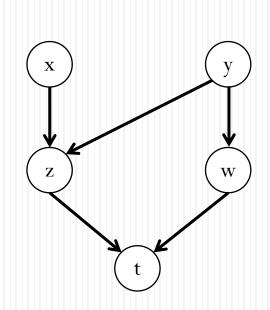


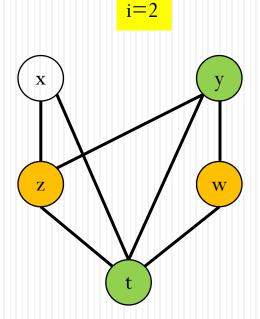
i=0, size of subset S=1 for which  $x\perp y|S$  or  $x\perp y|z$ ,  $z\in S_{xy}$  we test all pairs x-y for independence, conditioned on single neighbour variable z and for such pairs the link is removed.

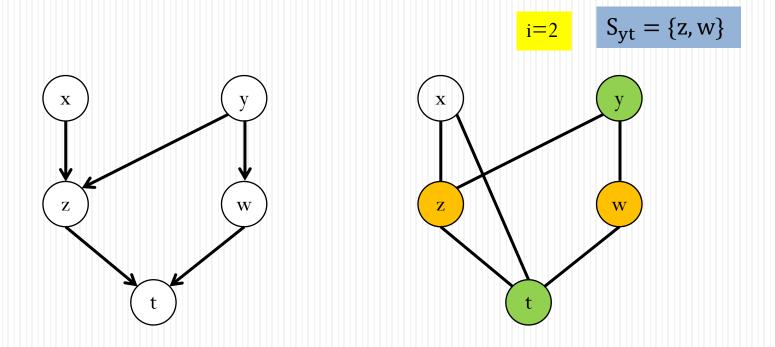


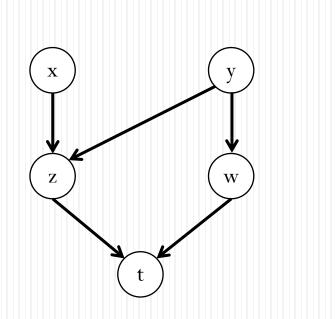


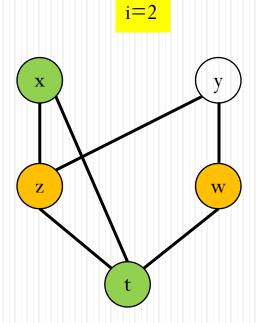
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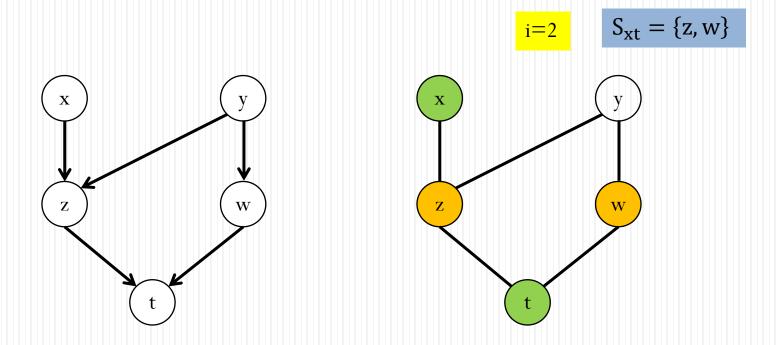


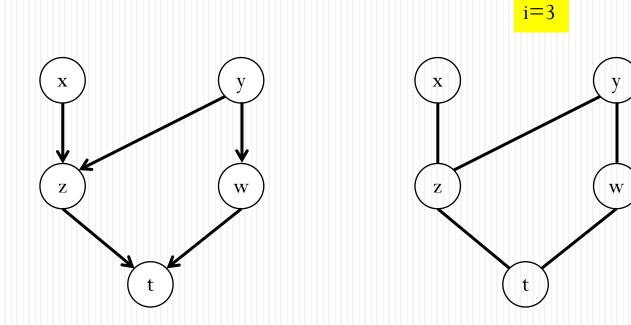








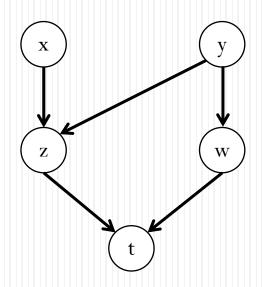


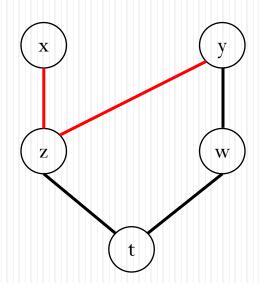


The algorithm terminates after this round as there are no nodes with 3 or more neighbours

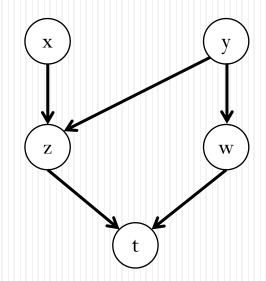
$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

### Determine the orientation

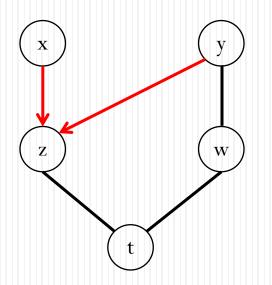




### Determine the orientation



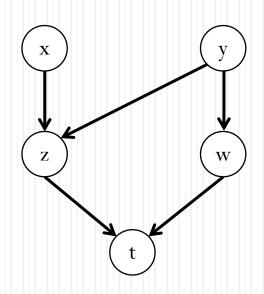
 $z \notin S_{xy}$ 

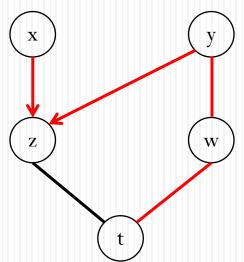


$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

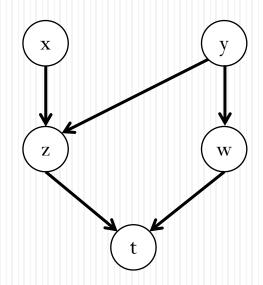
### Determine the orientation



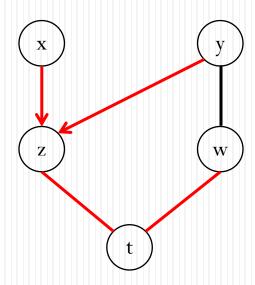




### Determine the orientation

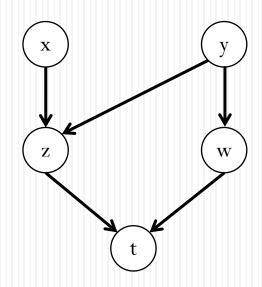


 $t \notin S_{zw}$ 

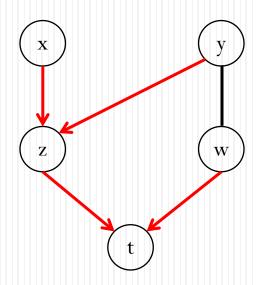


$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

### Determine the orientation

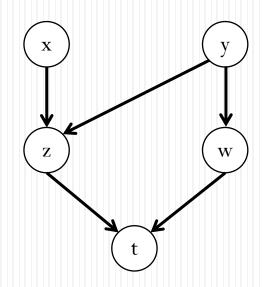




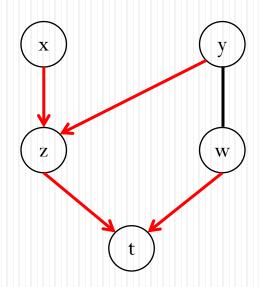


$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

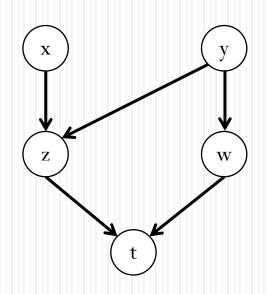
### Determine the orientation

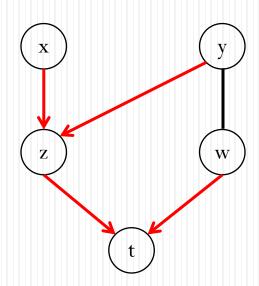






$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

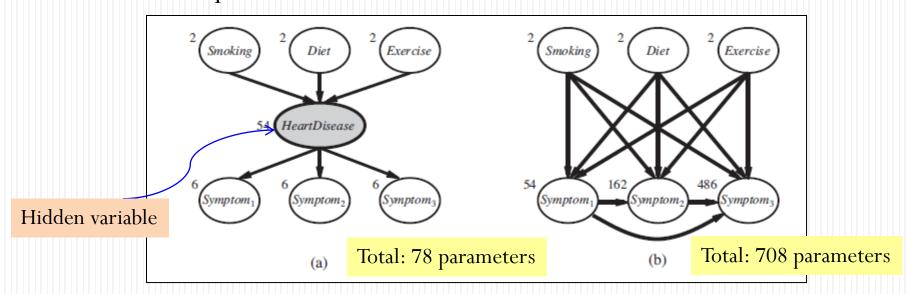




Partial DAG: the remaining edges can be arbitrarily oriented provided that the graph remains a DAG and no additional convergent edges are introduced.

$$S_{xy} = \{ \}, S_{xw} = \{ \}, S_{zw} = \{y\}, S_{xt} = \{z, w\}, S_{yt} = \{z, w\}$$

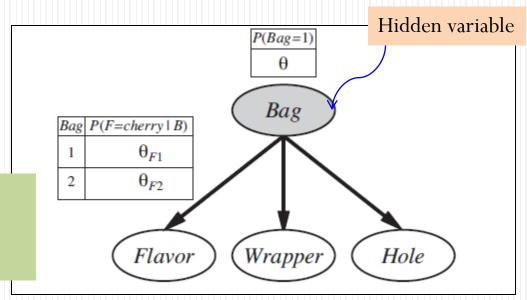
- Hidden variables (or latent variables) not observable in the data that are available for learning.
- Latent variables dramatically reduce the number of parameters in a Bayesian network and, in turn, reduces the amount of data needed to learn the parameters.



Each variable has three possible values and is labeled with the no. of independent parameters in its conditional distribution.

- Example: Consider a situation where two bags of candies are mixed together.
- Candies are describe by three features:
  - *Flavour* (lime and cherry), *Wrapper* (red and green), *Hole* (candy with hole or without hole)
- Proportions of different flavours, wrappers, presence of holes depend on the bag, which is not observed.

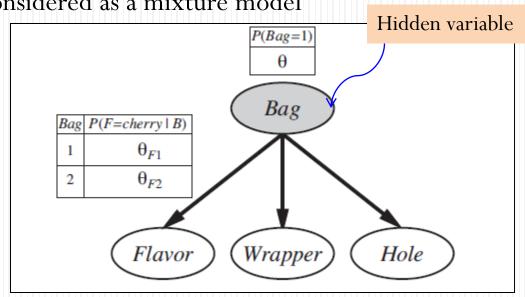
We want to predict for each candy, which was its original bag, from its features.



• Parameters:  $\theta$  is the prior probability that a candy comes from Bag 1  $\theta_{F1}$  and  $\theta_{F2}$  are the probabilities that the flavor is cherry, given that the candy comes from Bag 1 or Bag 2 respectively;  $\theta_{W1}$  and  $\theta_{W2}$  give the probabilities that the wrapper is red; and  $\theta_{H1}$  and  $\theta_{H2}$  give the probabilities that the candy has a hole.

the overall model can be considered as a mixture model

We will use Expectation
 —Maximization (EM)
 algorithm to learn the
 parameters of the model
 where Bag is hidden
 variable.



- Expectation Maximization- applied to unsupervised clustering
- Clustering: revealing multiple categories in collection of objects where category labels are not given.
- In clustering, the data is assumed to be generated from a mixture distribution *P* with *k* components, each of which is a distribution in its own right.
- Let the random variable C denote the component, with values 1, ..., k; then the mixture distribution is given by

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

• For continuous data, the component distributions can be mixture of Gaussians that gives the mixture of Gaussians distributions.

- Parameters of mixture of Gaussians:
  - $w_i = P(C = i)$ , weight of each component
  - ullet  $oldsymbol{\mu}_i$  , mean of each component
  - $\Sigma_i$ , covariance of each component

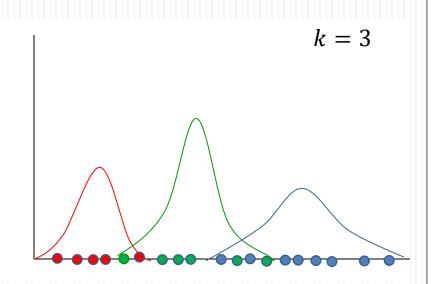
unsupervised clustering problem, then, is to recover a mixture model like the one

### Model reconstructed by Original distribution with 3 Raw data sampled from EM from data in (b) components model in (a) 0.8 0.8 0.8 0.6 0.6 0.6 0.4 0.4 0.4 0.2 0.2 0.20.4 0.6 0.8 0.6 0.6 0.8 (a) (b) (c)

- If it is known that which data sample came from which component distribution then we can estimate the parameters (i.e. the  $\mu$  and  $\sigma$  of the Gaussians).
- If we know the parameters of the components (Gaussians) then we can probabilistically assign each data point to a component.

We know neither the assignments nor the parameters —Problem!

EM algorithm assumes that the parameters are known, it then infers the probability that each data point belongs to each component. After that, it again refits the component to entire data and this is iterated over until convergence.



- EM algorithm
  - Initialize the model parameters (randomly) and iterate the following steps
  - 1. E-step: Compute the probabilities  $p_{ij} = P(C = i \mid \mathbf{x}_j)$ , the probability that datum  $\mathbf{x}_j$  was generated by component i. By Bayes' rule, we have  $p_{ij} = \alpha P(\mathbf{x}_j \mid C = i)P(C = i)$ . The term  $P(\mathbf{x}_j \mid C = i)$  is just the probability at  $\mathbf{x}_j$  of the ith Gaussian, and the term P(C = i) is just the weight parameter for the ith Gaussian. Define  $n_i = \sum_j p_{ij}$ , the effective number of data points currently assigned to component i.
  - M-step: Compute the new mean, covariance, and component weights using the following steps in sequence:

$$\mu_i \leftarrow \sum_j p_{ij} \mathbf{x}_j / n_i$$

$$\Sigma_i \leftarrow \sum_j p_{ij} (\mathbf{x}_j - \mu_i) (\mathbf{x}_j - \mu_i)^\top / n_i$$

$$w_i \leftarrow n_i / N$$

EM algorithm

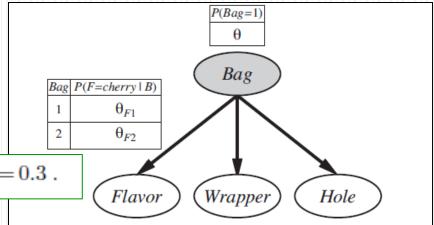
where N is the total number of data points. The E-step, or *expectation* step, can be viewed as computing the expected values  $p_{ij}$  of the hidden **indicator variables**  $Z_{ij}$ , where  $Z_{ij}$  is 1 if datum  $\mathbf{x}_j$  was generated by the *i*th component and 0 otherwise. The M-step, or *maximization* step, finds the new values of the parameters that maximize the log likelihood of the data, given the expected values of the hidden indicator variables.

• EM increases the log-likelihood of data at every iteration (this can be proved in general).

### Coming back to initial problem!

- Parameters:  $\theta$  is the prior probability that a candy comes from Bag 1;  $\theta_{F1}$  and  $\theta_{F2}$  are the probabilities that the flavor is cherry, given that the candy comes from Bag 1 or Bag 2 respectively;  $\theta_{W1}$  and  $\theta_{W2}$  give the probabilities that the wrapper is red; and  $\theta_{H1}$  and  $\theta_{H2}$  give the probabilities that the candy has a hole.
  - 1000 samples are generated from the following (true) model:

$$\theta = 0.5, \ \theta_{F1} = \theta_{W1} = \theta_{H1} = 0.8, \ \theta_{F2} = \theta_{W2} = \theta_{H2} = 0.3$$
.



	W = red		W = green	
	H = 1	H = 0	H = 1	H = 0
F = cherry	273	93	104	90
F = lime	79	100	94	167

• For EM algorithm initialize the parameters (arbitrarily):

$$\theta^{(0)} = 0.6, \ \ \theta_{F1}^{(0)} = \theta_{W1}^{(0)} = \theta_{H1}^{(0)} = 0.6, \ \ \theta_{F2}^{(0)} = \theta_{W2}^{(0)} = \theta_{H2}^{(0)} = 0.4 \ .$$

- Learning of Parameter  $\theta$ 
  - E-step:

find the expected count of number of candies in Bag1 : Sum of the probabilities that each of the N data points comes from bag 1

$$\hat{\mathbf{N}}(\mathrm{Bag} = 1) = \sum_{j=1}^{N} P(Bag = 1|flavor_{j}, wrapper_{j}, \ hole_{j})$$

$$\hat{\mathbf{N}}(\mathrm{Bag} = 1) = \sum_{j=1}^{N} P(Bag = 1|flavor_{j}, wrapper_{j}, whole_{j})$$

$$= \sum_{j=1}^{N} \frac{P(flavor_{j}, wrapper_{j}, hole_{j}|Bag = 1)P(Bag = 1)}{P(flavor_{j}, wrapper_{j}, hole_{j})}$$

	W = red		W = green	
	H = 1	H = 0	H = 1	H = 0
F = cherry	273	93	104	90
F = lime	79	100	94	167

$$=\sum_{j=1}^{N}\frac{P(flavor_{j}|Bag=1)P(wrapper_{j}|Bag=1)P(hole_{j}|Bag=1)P(Bag=1)}{\sum_{i}P(flavor_{j}|Bag=i)P(wrapper_{j}|Bag=i)P(hole_{j}|Bag=i)P(Bag=i)}$$

This summation can be broken down for 8 candy groups

For example, sum over 273 cheery candies with red wrapper and holes is:

$$= 273 \frac{\theta_{F1}^{(0)} \theta_{W1}^{(0)} \theta_{H1}^{(0)} \theta^{(0)}}{\theta_{F1}^{(0)} \theta_{W1}^{(0)} \theta_{H1}^{(0)} \theta^{(0)} + \theta_{F2}^{(0)} \theta_{W2}^{(0)} \theta_{H2}^{(0)} (1 - \theta^{(0)})} =$$

$$273 \frac{0.6^4}{0.6^4 + 0.4^4} = 273 \frac{0.1296}{0.1552} = 227.97$$

• summing over other 7 groups in the table

$$\hat{N}(Bag = 1) = 612.4$$

- M-step
  - ullet Compute new  $oldsymbol{ heta}$  using the expected count of candies that came from Bag 1.

$$\theta(1) = \frac{\hat{N}(Bag = 1)}{N} = 0.6124$$

- Similarly, we can learn other 5 parameters.
- Learning parameter  $heta_{F1}$

E-step: compute the expected count of cherry candies from Bag 1

$$\hat{N}(Bag = 1 \land Flavor = cherry) = \sum_{j:Flavor_j = cherry} P(Bag = 1 \mid Flavor_j = cherry , wrapper_j, hole_j)$$

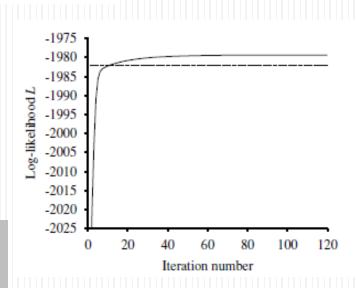
M-step: refine  $\theta_{F1}$  by computing the corresponding expected frequencies

$$\theta_{{\scriptscriptstyle F1}}^{(1)} = \frac{\hat{N}(Bag=1 \land Flavor=cherry)}{\hat{N}(Bag=1)}$$

After a complete cycle through all the parameters, we get

$$\theta^{(1)} = 0.6124;$$
 $\theta_{F1}^{(1)} = 0.6684;$ 
 $\theta_{W1}^{(1)} = 0.6483;$ 
 $\theta_{H1}^{(1)} = 0.658;$ 
 $\theta_{F2}^{(1)} = 0.3887;$ 
 $\theta_{W2}^{(1)} = 0.3817;$ 
 $\theta_{H2}^{(1)} = 0.3827;$ 

• EM increases the log likelihood of the data at every iteration



### What did we discussed in L19-21?

- How to learn parameters (CPTs) of a Bayesian Network when complete observations are given?
  - Maximum-likelihood approach
  - Bayesian approach
- How to learn the structure of Bayesian network?
- Some non-parametric approaches of Density estimation
  - Histograms
  - KDE
  - k-NN
- How to learn the parameters of Bayesian Network with Hidden variables?
  - Expectation Maximization algorithm