# **Probabilistic Graphical Models**

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# **Probabilistic Graphical Models**



- Modeling many real-world problems => a large number of random variables
  - Dependences among variables may be used to reduce the size to encode the model (PCA ?), or
  - They may be the goal by themselves, that is, the idea is to understand the correlations among variables.

# **Modeling the domain**



- Discrete random variables
  - Take 5 random binary variables (A, B, C, D, E)
  - i.i.d. data from a multinomial distribution

A	B	С	D	E
а	~b	~C	~d	~e
а	b	~ <i>C</i>	d	~ <i>e</i>
а	~b	С	d	~e

#### Goals



- (Parameter) Learning: using training data, estimate the joint distribution
  - Which are the values P(A, B, C, D, E),?
  - ... and if there were one hundred binary variables? (Size of model certainly greater than number of atoms on Earth!)
- Inference: Given the distribution *P*(*A*, *B*, *C*, *D*, *E*),
  - Belief updating: compute the probability of an event
    - What is the probability of A=a given E=e?
  - Maximum a posterior: compute the states of variables that maximize their probability.
    - Which state of A maximizes P(A | E=e)? Is it a or  $\sim a$ ?

# The unstructured approach

• To specify the joint distribution, there is an exponential number of values:

$$p(a, b, c, d, e), p(a, b, c, d, \neg e), p(a, b, c, \neg d, e),$$

$$p(a, b, c, \neg d, \neg e), p(a, b, \neg c, d, e), p(a, b, \neg c, d, \neg e),$$

$$p(a, b, \neg c, \neg d, e), p(a, b, \neg c, \neg d, \neg e), \dots$$

• We can compute the probability of events by:

$$p(a) = \sum_{B,C,D,E} p(a, B, C, D, E)$$

$$p(a|d,\neg e) = \frac{p(a,d,\neg e)}{p(d,\neg e)} = \frac{\sum_{B,C} p(a,B,C,d,\neg e)}{\sum_{A,B,C} p(A,B,C,d,\neg e)}$$

There are exponentially many terms in the summations...



# The naïve Bayesian approach

p(a, b) = p(a) p(b)

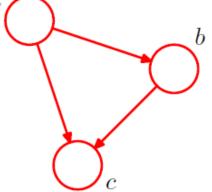
#### • Application: Email spanning

# **Bayesian Networks**



- An arbitrary joint distribution p(a, b, c) over three variables a, b, and c
  - the product rule of probability:

p(a, b, c) = p(c | a, b) p(a, b)= p(c | a, b) p(b / a) p(a)



• General case:  $p(x_1, x_2, \dots, x_K)$ 

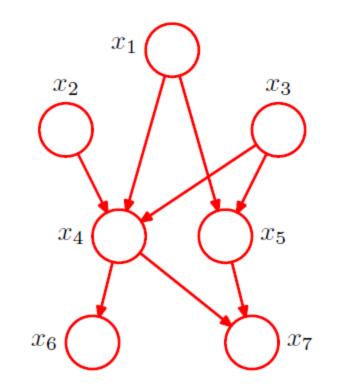
 $p(x_1, \ldots, x_K) = p(x_K | x_1, \ldots, x_{K-1}) \ldots p(x_2 | x_1) p(x_1)$ 

# Not fully connected graph



• Joint distribution:  $p(x_1, x_2, \dots, x_7)$ 

 $p(x_1)p(x_2)p(x_3)p(x_4|x_1, x_2, x_3)p(x_5|x_1, x_3)p(x_6|x_4)p(x_7|x_4, x_5)$ 







• For a graph with *K* nodes, the joint distribution is given by:

$$p(\mathbf{x}) = \prod_{k=1}^{K} p(x_k | \mathrm{pa}_k)$$

• where  $pa_k$  denotes the set of parents of  $x_k$ , and  $x = \{x_1, \ldots, x_K\}$ 



# Definitions

 A set of variables associated with nodes of a Directed Acyclic Graph (DAG).

D

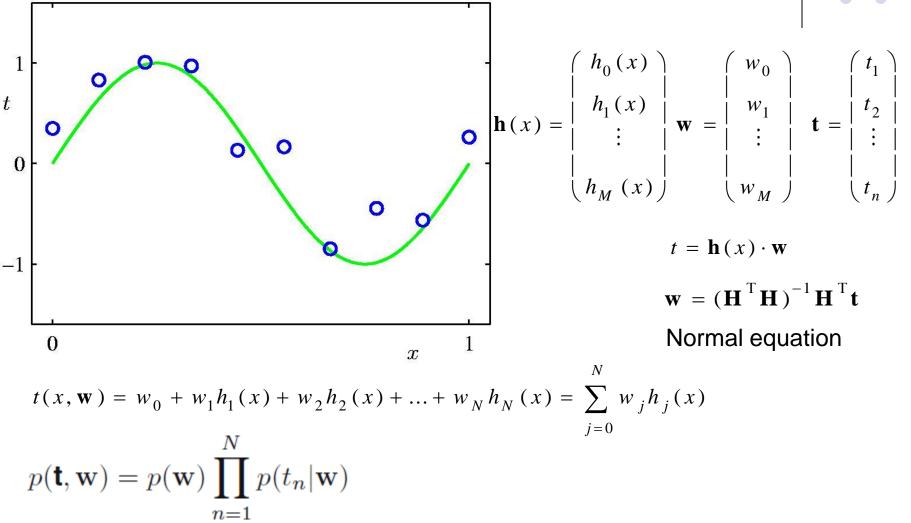
- Markov condition (w.r.t. the DAG): each variable is independent of its non-descendants given its parents.
- For each variable (node), local probability distributions:
  - P(A), P(B|A=a), P(B|A=a), P(C|A=a), P(C|A=~a), P(D|b, c), P(D|~b, c), P(D|~b,c);
     P(D|~b,~c), P(E|c), P(E|~c),

С

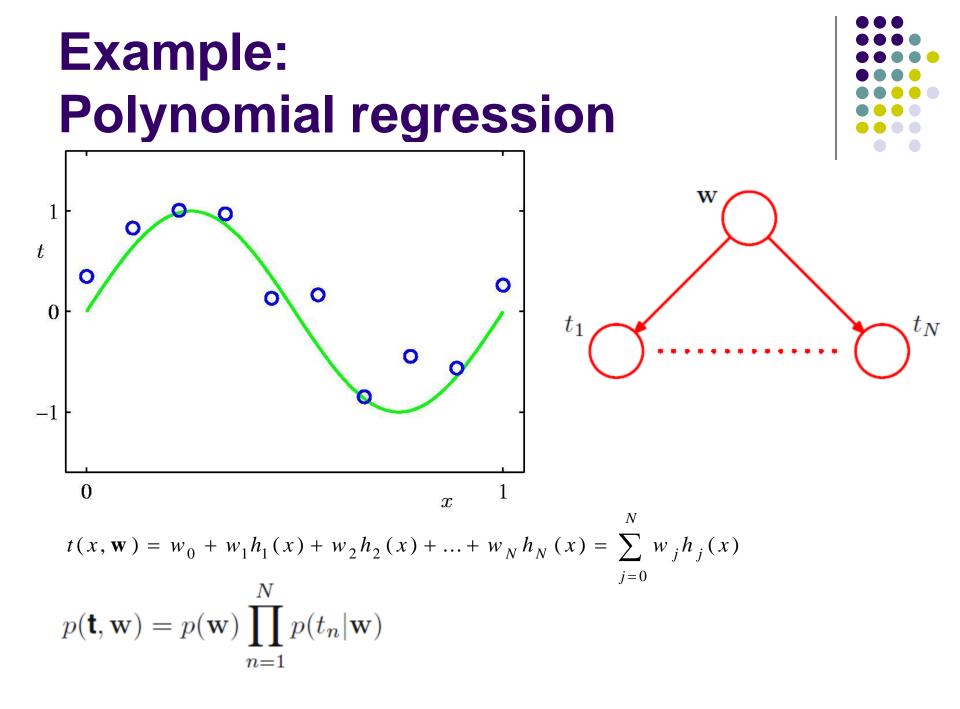
Ε

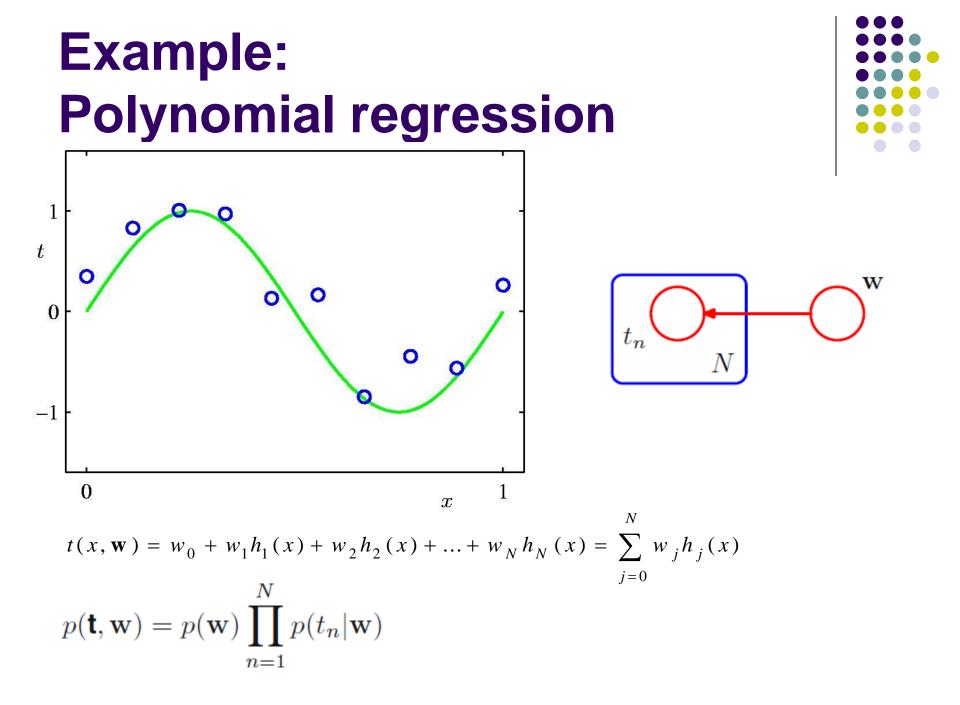
• All these values are precise.

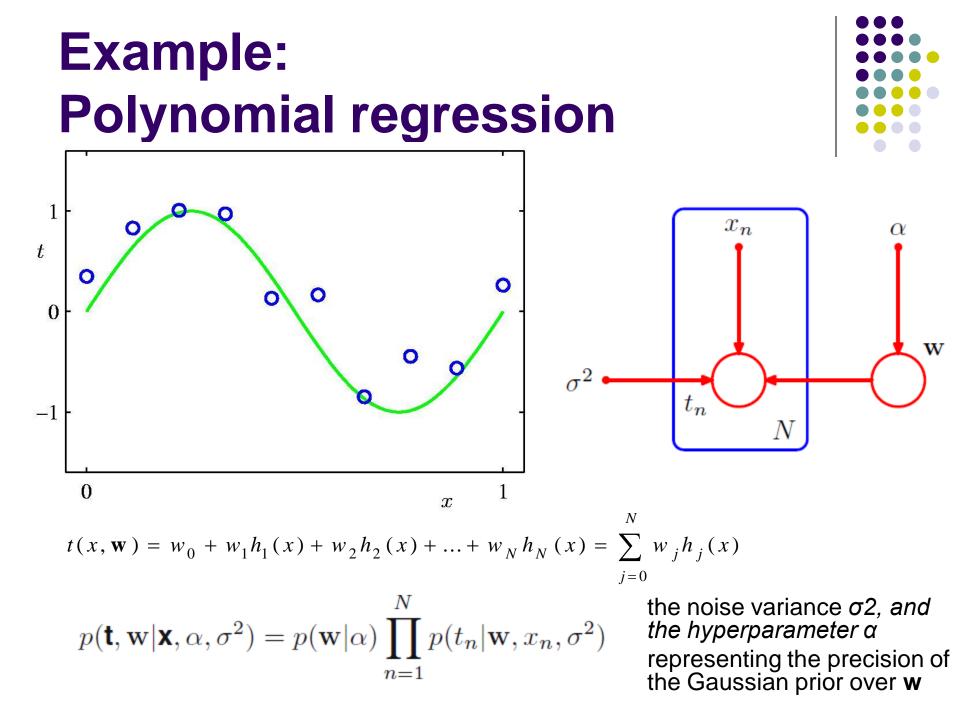
# Regression revisit: Polynomial Curve Fitting













- Consider an arbitrary DAG over *D* variables in which node *i* represents a single continuous random variable x<sub>i</sub> having a *Gaussian distribution*
- The mean of this distribution is taken to be a linear combination of the states of its parent nodes pa<sub>i</sub> of node *i*

$$p(x_i | \text{pa}_i) = \mathcal{N}\left(x_i \left| \sum_{j \in \text{pa}_i} w_{ij} x_j + b_i, v_i \right. \right)$$

$$p(x_i | pa_i) = \mathcal{N}\left(x_i \left| \sum_{j \in pa_i} w_{ij} x_j + b_i, v_i \right. \right)$$

$$\ln p(\mathbf{x}) = \sum_{i=1}^{D} \ln p(x_i | \text{pa}_i)$$
$$= -\sum_{i=1}^{D} \frac{1}{2v_i} \left( x_i - \sum_{j \in \text{pa}_i} w_{ij} x_j - b_i \right)^2 + \text{const}$$

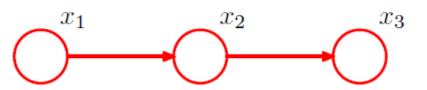


$$p(x_i | pa_i) = \mathcal{N} \left( x_i \left| \sum_{j \in pa_i} w_{ij} x_j + b_i, v_i \right. \right) \right.$$
$$x_i = \sum_{j \in pa_i} w_{ij} x_j + b_i + \sqrt{v_i} \epsilon_i \qquad \mathbb{E}[x_i] = \sum_{j \in pa_i} w_{ij} \mathbb{E}[x_j] + b_i$$

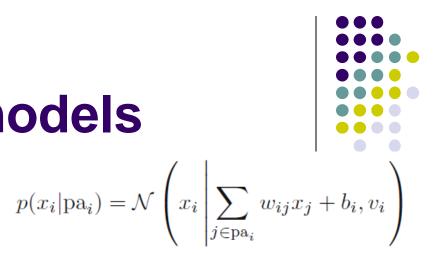
$$\operatorname{cov}[x_i, x_j] = \mathbb{E}\left[ (x_i - \mathbb{E}[x_i])(x_j - \mathbb{E}[x_j]) \right]$$
$$= \mathbb{E}\left[ (x_i - \mathbb{E}[x_i]) \left\{ \sum_{k \in \operatorname{pa}_j} w_{jk}(x_k - \mathbb{E}[x_k]) + \sqrt{v_j} \epsilon_j \right\} \right]$$
$$= \sum_{k \in \operatorname{pa}_j} w_{jk} \operatorname{cov}[x_i, x_k] + I_{ij} v_j$$



- Case 1: no links in the graph
  - The joint distribution:
    - 2D parameters and represents
    - D independent univariate Gaussian distributions.
- Case 2: fully connected graph
  - D(D-1)/2+D independent parameters
- Case 3:



$$\boldsymbol{\mu} = (b_1, b_2 + w_{21}b_1, b_3 + w_{32}b_2 + w_{32}w_{21}b_1)^{\mathrm{T}} \boldsymbol{\Sigma} = \begin{pmatrix} v_1 & w_{21}v_1 & w_{32}w_{21}v_1 \\ w_{21}v_1 & v_2 + w_{21}^2v_1 & w_{32}(v_2 + w_{21}^2v_1) \\ w_{32}w_{21}v_1 & w_{32}(v_2 + w_{21}^2v_1) & v_3 + w_{32}^2(v_2 + w_{21}^2v_1) \end{pmatrix}$$



# **Conditional independence**

- Three random variables: a, b and c
  - a is conditionally independent of b given c iff
  - $P(a \mid b, c) = P(a \mid c)$

$$a \perp b \mid c$$

- This can be re-written in following way
  - P(a, b | c) = P(a | b, c) P(b | c)

 $= P(a \mid c) P(b \mid c)$ 

The joint distribution of a and b factorizes into the product of the marginal distribution of a and ~b.



# Simple example (1)

- Joint distribution:
  - P(a, b, c) = P(a | c) P(b | c) P(c)
- Condition on c:
  - P(a, b | c) = P(a, b, c) / P(c) = P(a | c) P(b | c)• =>  $a \perp b | c$

# Simple example (2)

- Joint distribution:
  - P(a, b, c) = P(a) P(c | a) P(b | c)
- $a \perp b \mid c$

• Factorization:

$$P(a,b) = \sum_{c} P(a,b,c) = P(a) \sum_{c} P(c \mid a) P(b \mid c)$$
$$= P(a) P(b \mid a)$$

• Condition on c:

# Simple example (3)

- Joint distribution:
  - P(a, b, c) = P(a) P(b) P(c | a, b)

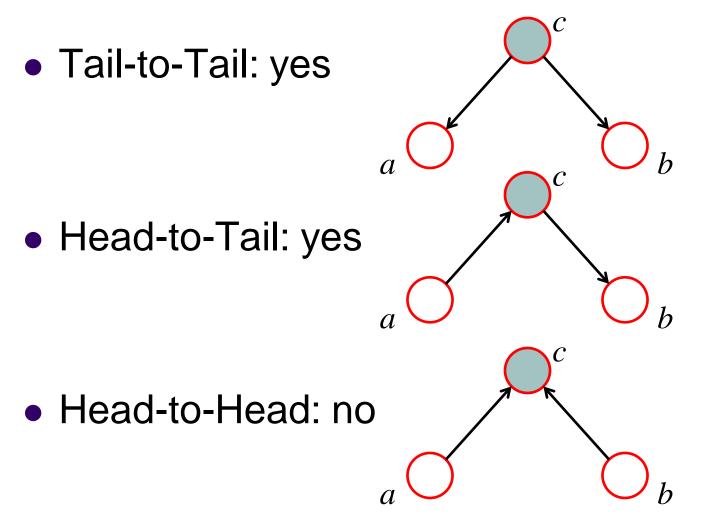
• Factorization:

$$P(a,b) = \sum_{c} P(a,b,c) = P(a)P(b)\sum_{c} P(c \mid a,b)$$
$$= P(a)P(b)$$

• Condition on c:

$$P(a,b \mid c) = \frac{P(a,b,c)}{P(c)} = \frac{P(a)P(b)P(c \mid a,b)}{P(c)}$$
$$\neq P(a \mid c)P(b \mid c)$$

# **Conditional independence**







# Markov condition

 We say that node y is a descendant of node x if there is a path from x to y in which each step of the path follows the directions of the arrows.

В

С

D

E

• If each variable is independent of its non-descendants given its parents, then:

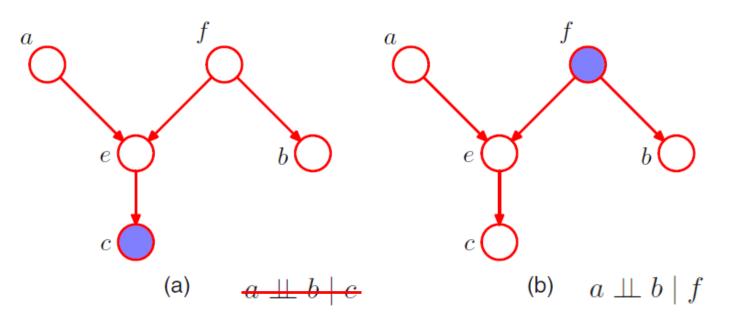
 $B \perp \perp (C, E) | A,$   $D \perp \perp (A, E) | (B, C),$  $E \perp \perp (A, B, D) | C.$ 

# **D-separation**



- All possible paths from any node in A to any node in B. Any such path is said to be *blocked* if it includes a node such that either
  - the arrows on the path meet either head-to-tail or tail-to-tail at the node, and the node is in the set *C*, or
  - the arrows meet head-to-head at the node, and neither the node, nor any of its descendants, is in the set *C*
- If all paths are blocked, then A is said to be *d-separated* from B by C.





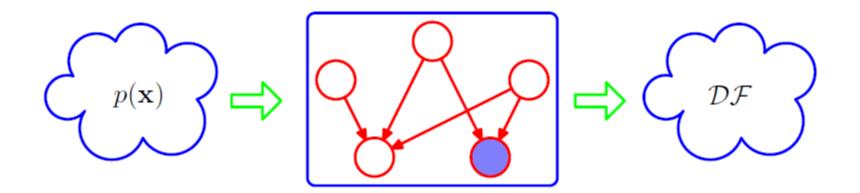
- In graph (a), the path from a to b is not blocked by node c
- In graph (b), the path from a to b is blocked by node f and e



# **D-separation**



- A particular directed graph represents a specific decomposition of a joint probability distribution into a product of conditional probabilities
- A directed graph is a filter



# Markov blanket



 $x_i$ 

Joint distribution p(x<sub>1</sub>, ..., x<sub>D</sub>) represented by a directed graph having D nodes

• 
$$p(\mathbf{x}_i | \mathbf{x}_{\{j \neq i\}}) = \frac{p(\mathbf{x}_1, \dots, \mathbf{x}_D)}{\int p(\mathbf{x}_1, \dots, \mathbf{x}_D) \, \mathrm{d}\mathbf{x}_i}$$
  
$$= \frac{\prod_k p(\mathbf{x}_k | \mathrm{pa}_k)}{\int \prod_k p(\mathbf{x}_k | \mathrm{pa}_k) \, \mathrm{d}\mathbf{x}_i}$$

• The set of nodes comprising the parents, the children and the co-parents is called the Markov blanket

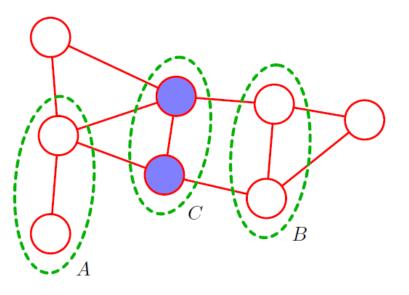
# **Markov Random Fields**



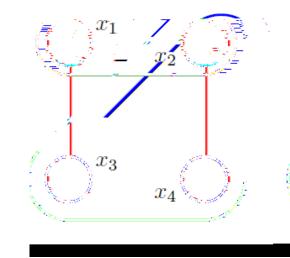
- Also known as a Markov network or an undirected graphical model
- Conditional independence properties:

Conditional dependence exists if there exists a path that connects any node in *A to any* node in *B*.

If there are no such paths, then the conditional independence property must hold.



# Clique

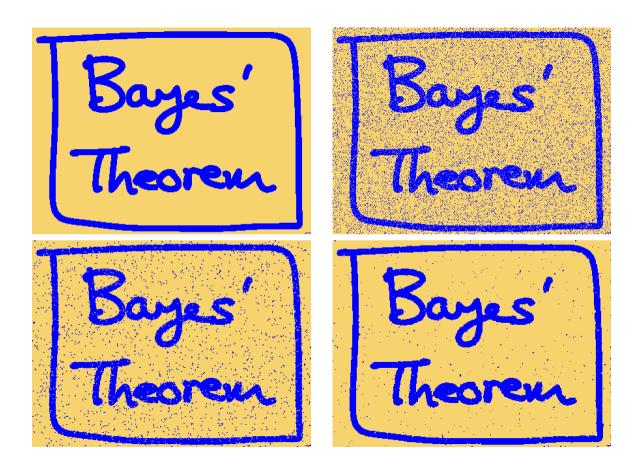


- A subset of the nodes in a graph such that there exists a link between all pairs of nodes in the subset
  - In other words, the set of nodes in a clique is fully connected

- Maximal clique ...
- A four-node undirected graph showing a clique (outlined in green) and a maximal clique (outlined in blue)

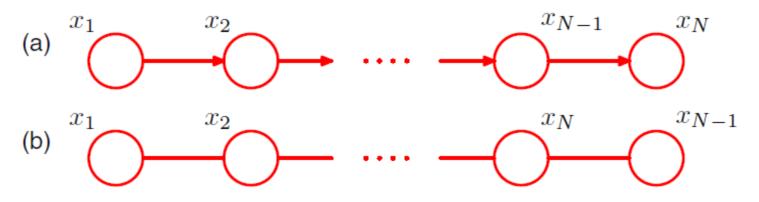


# Image de-noising





# **Relation to directed graphs**



- Joint distribution:
  - Directed:

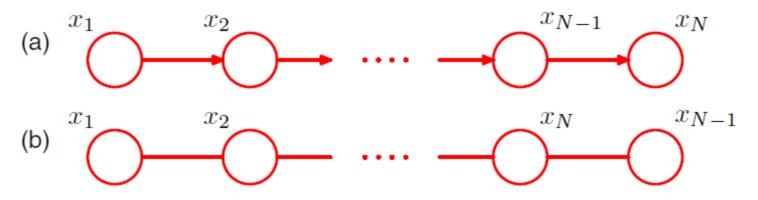
$$p(\mathbf{x}) = p(x_1)p(x_2|x_1)p(x_3|x_2)\cdots p(x_N|x_{N-1})$$

• Undirected:

$$p(\mathbf{x}) = \frac{1}{Z} \psi_{1,2}(x_1, x_2) \psi_{2,3}(x_2, x_3) \cdots \psi_{N-1,N}(x_{N-1}, x_N)$$



# **Relation to directed graphs**



$$\psi_{1,2}(x_1, x_2) = p(x_1)p(x_2|x_1)$$
  

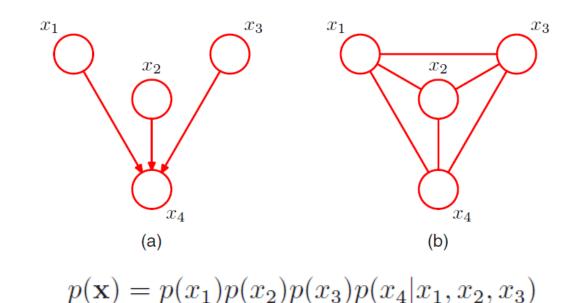
$$\psi_{2,3}(x_2, x_3) = p(x_3|x_2)$$
  

$$\vdots$$
  

$$\psi_{N-1,N}(x_{N-1}, x_N) = p(x_N|x_{N-1})$$



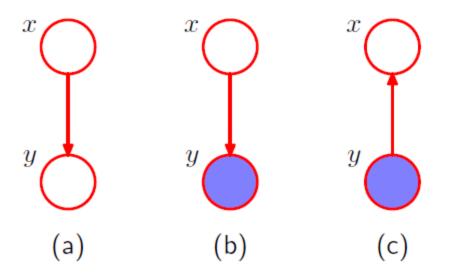
# **Relation to directed graphs**



 this process of 'marrying the parents' has become known as *moralization*, and the resulting undirected graph, after dropping the arrows, is called the *moral graph*.



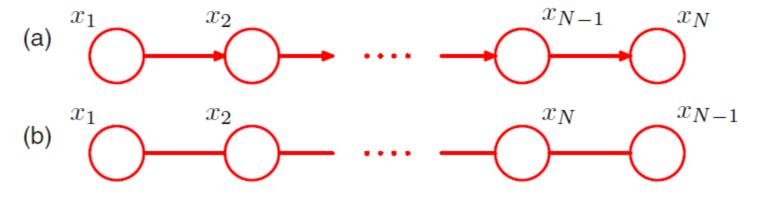
# **Inference in Graphical Models**



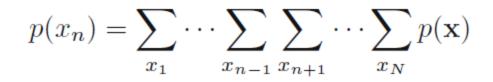
p(x,y) = p(x)p(y|x)

$$p(x|y) = \frac{p(y|x)p(x)}{p(y)}$$

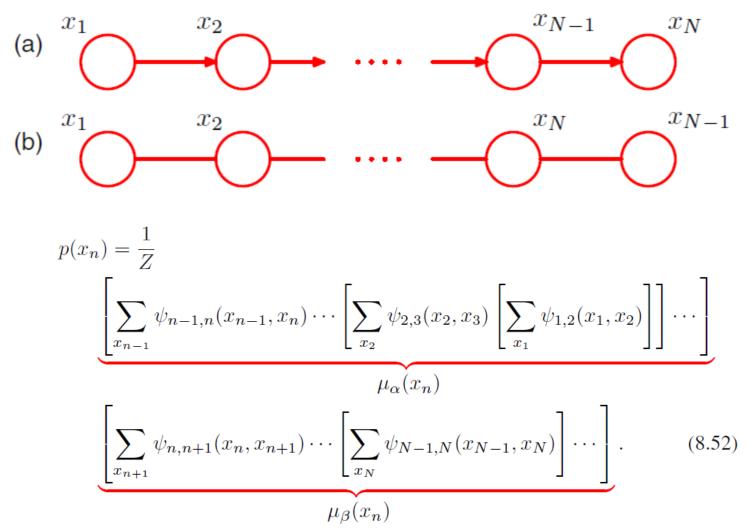




$$p(\mathbf{x}) = \frac{1}{Z} \psi_{1,2}(x_1, x_2) \psi_{2,3}(x_2, x_3) \cdots \psi_{N-1,N}(x_{N-1}, x_N)$$









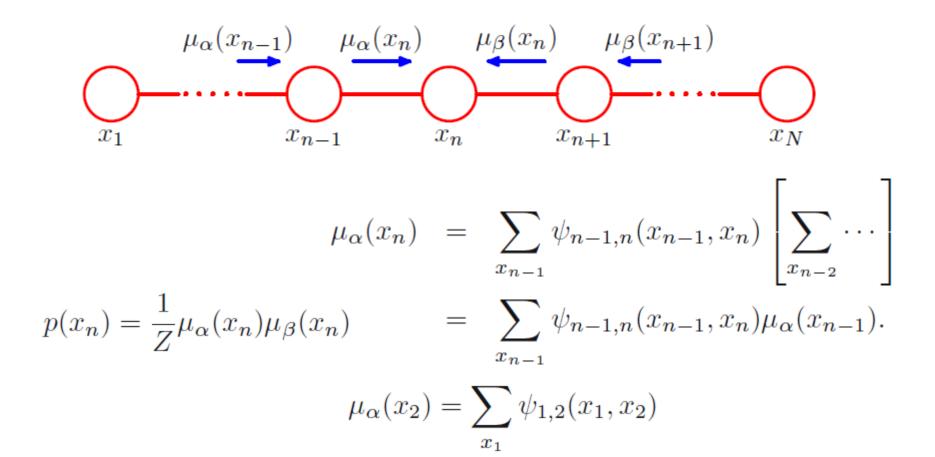
Passing of local messages around on the graph

 $p(x_n) = \frac{1}{Z}$   $\left[\sum_{x_{n-1}} \psi_{n-1,n}(x_{n-1}, x_n) \cdots \left[\sum_{x_2} \psi_{2,3}(x_2, x_3) \left[\sum_{x_1} \psi_{1,2}(x_1, x_2)\right]\right] \cdots\right]$   $\mu_{\alpha}(x_n)$   $\left[\sum_{x_{n+1}} \psi_{n,n+1}(x_n, x_{n+1}) \cdots \left[\sum_{x_N} \psi_{N-1,N}(x_{N-1}, x_N)\right] \cdots\right].$  (8.52)  $\mu_{\beta}(x_n)$ 

$$p(x_n) = \frac{1}{Z} \mu_{\alpha}(x_n) \mu_{\beta}(x_n)$$

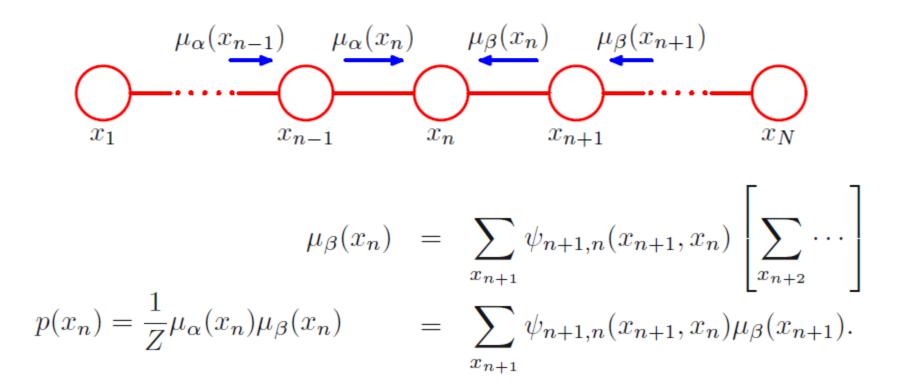


Passing of local messages around on the graph



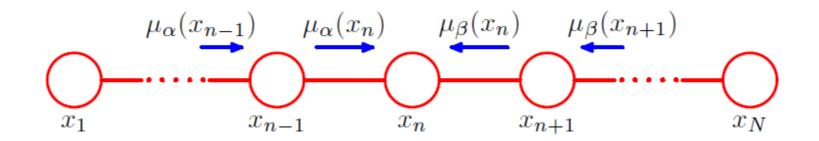


Passing of local messages around on the graph

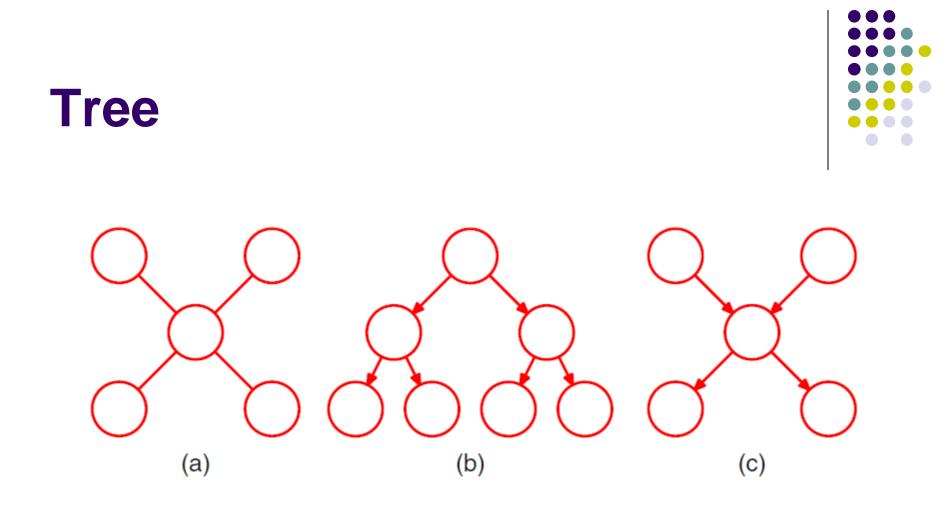




Passing of local messages around on the graph



$$p(x_{n-1}, x_n) = \frac{1}{Z} \mu_{\alpha}(x_{n-1}) \psi_{n-1,n}(x_{n-1}, x_n) \mu_{\beta}(x_n)$$





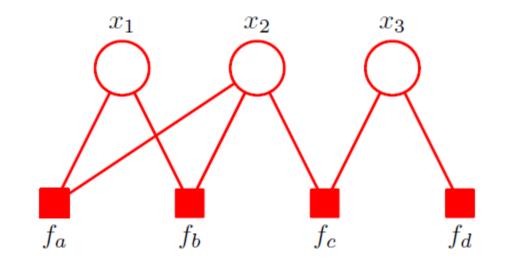
 the joint distribution over a set of variables in the form of a product of factors

$$p(\mathbf{x}) = \prod_{s} f_s(\mathbf{x}_s)$$

• where  $\mathbf{x}_s$  denotes a subset of the variables



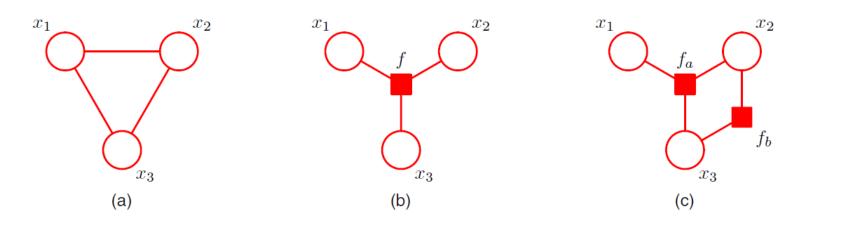
 $p(\mathbf{x}) = f_a(x_1, x_2) f_b(x_1, x_2) f_c(x_2, x_3) f_d(x_3)$ 





- an undirected graph => a factor graph
  - create variable nodes corresponding to the nodes in the original undirected graph
  - create additional factor nodes corresponding to the maximal cliques x<sub>s</sub>
  - Multiple choices of fg.



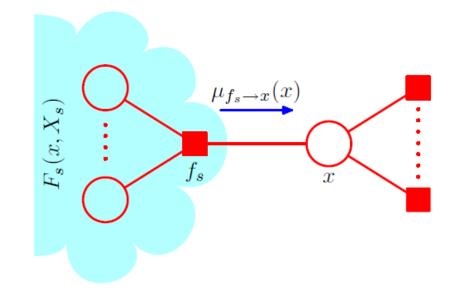


- (a) An undirected graph with a single clique potential  $\psi(x1, x2, x3)$ .
- (b) A factor graph with factor  $f(x1, x2, x3) = \psi(x1, x2, x3)$ representing the same distribution as the undirected graph.
- (c) A different factor graph representing the same distribution, whose factors satisfy  $fa(x1, x2, x3)fb(x1, x2) = \psi(x1, x2, x3)$ .



 The problem of finding the marginal p(x) for particular variable node x

 $p(x) = \sum_{\mathbf{x} \setminus x} p(\mathbf{x})$   $p(\mathbf{x}) = \prod_{s \in ne(x)} F_s(x, X_s)$ 

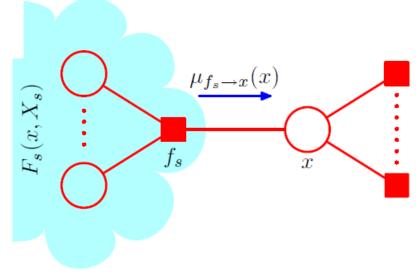




 The problem of finding the marginal p(x) for particular variable node x

$$p(x) = \prod_{s \in ne(x)} \left[ \sum_{X_s} F_s(x, X_s) \right]$$
$$= \prod_{s \in ne(x)} \mu_{f_s \to x}(x).$$

$$\mu_{f_s \to x}(x) \equiv \sum_{X_s} F_s(x, X_s)$$





 The problem of finding the marginal p(x) for particular variable node x

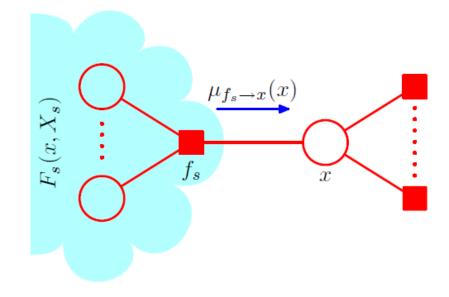
$$\mu_{f_s \to x}(x) \equiv \sum_{X_s} F_s(x, X_s)$$

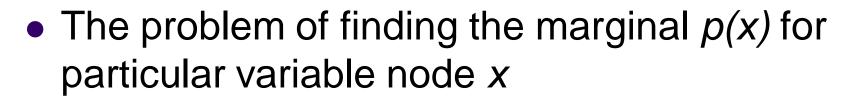


 The problem of finding the marginal p(x) for particular variable node x

$$F_{s}(x, X_{s}) = f_{s}(x, x_{1}, \dots, x_{M})G_{1}(x_{1}, X_{s1}) \dots G_{M}(x_{M}, X_{sM})$$

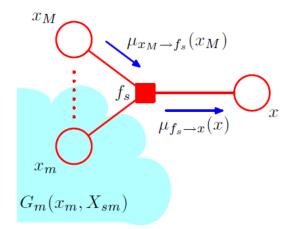
$$\mu_{f_s \to x}(x) \equiv \sum_{X_s} F_s(x, X_s)$$





$$\mu_{f_s \to x}(x) = \sum_{x_1} \dots \sum_{x_M} f_s(x, x_1, \dots, x_M) \prod_{m \in \operatorname{ne}(f_s) \setminus x} \left[ \sum_{X_{xm}} G_m(x_m, X_{sm}) \right]$$
$$= \sum_{x_1} \dots \sum_{x_M} f_s(x, x_1, \dots, x_M) \prod_{m \in \operatorname{ne}(f_s) \setminus x} \mu_{x_m \to f_s}(x_m) \quad (8.66)$$

$$\mu_{x_m \to f_s}(x_m) \equiv \sum_{X_{sm}} G_m(x_m, X_{sm})$$



Г



п

# **Junction tree algorithm**



- deal with graphs having loops
- Algorithm:
  - directed graph => undirected graph (moralization)
  - The graph is triangulated
  - join tree
  - Junction tree
  - a two-stage message passing algorithm, essentially equivalent to the sum-product algorithm

## **Graph inference example**



 Computer-Generated Residential Building Layouts [SIG ASIA 2010]





## MARKOV CHAINS AND HMM



## **Example: Video Textures**

## Problem statement



video clip

## video texture

SIGGRAPH 2000. Schoedl et. al.

## The approach



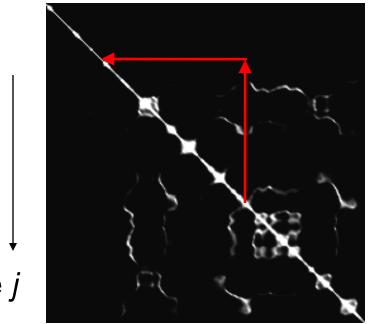
How do we find good transitions?



## **Finding good transitions**



## Compute L<sub>2</sub> distance $D_{i, j}$ between all frames <u>frame</u>



frame j

Similar frames make good transitions

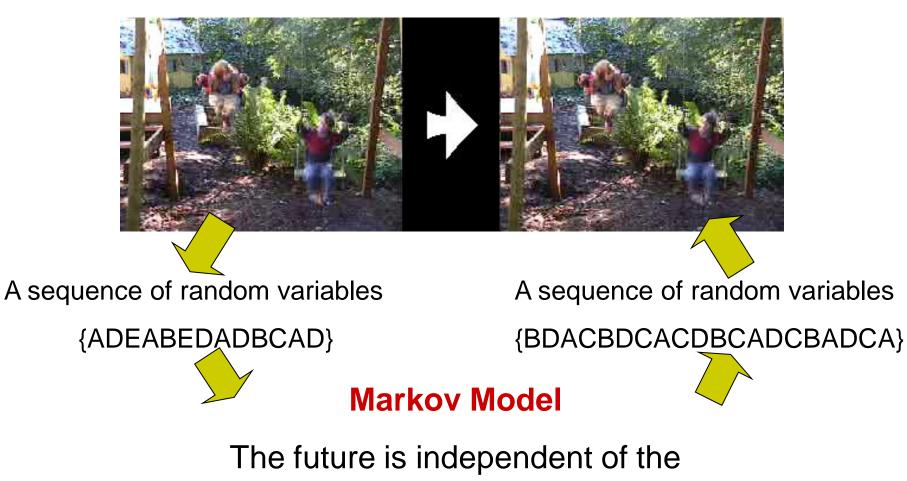
## **Demo: Fish Tank**





## Mathematic model of Video Texture





past and given by the present.

## **Markov Property**

- Formal definition  $x_1 \longrightarrow x_2 \longrightarrow x_3 \longrightarrow x_4$ 
  - Let  $X = \{X_n\}_{n=0...N}$  be a sequence of random variables taking values  $s_k \in N$  if and only if  $P(X_m = s_m / X_0 = s_0, ..., X_{m-1} = s_{m-1}) = P(X_m = s_m / X_{m-1} = s_{m-1})$

then the X fulfills Markov property

#### Informal definition

• The future is independent of the past given the present.

# **History of MC**

- Markov chain theory developed around 1900.
- Hidden Markov Models developed in late 1960's.
- Used extensively in speech recognition in 1960-70.
- Introduced to computer science in 1989.



Andrei Andreyevich Markov

#### **Applications**

- Bioinformatics.
- Signal Processing
- Data analysis and Pattern recognition



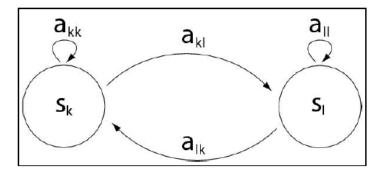
## Markov Chain

- A Markov chain is specified by
  - state space  $S = \{ s_1, s_2, ..., s_n \}$
  - initial distribution  $a_0$
  - transition matrix A

Where 
$$A(n)_{ij} = a_{ij} = P(q_t = s_j / q_{t-1} = s_i)$$



- Vertices represent states
- Edges represent transitions with positive probability





## **Probability Axioms**

 Marginal Probability – sum the joint probability

$$P(x = a_i) \equiv \sum_{y \in A_y} P(x = a_i, y)$$

Conditional Probability

$$P(x = a_i | y = b_j) \equiv \frac{P(x = a_i, y = b_j)}{P(y = b_j)} \text{ if } P(y = b_j) \neq 0.$$



## **Calculating with Markov chains**



- Probability of an observation sequence:
  - Let X={x<sub>t</sub>}<sup>L</sup><sub>t=0</sub> be an observation sequence from the Markov chain {S, a<sub>0</sub>, A}

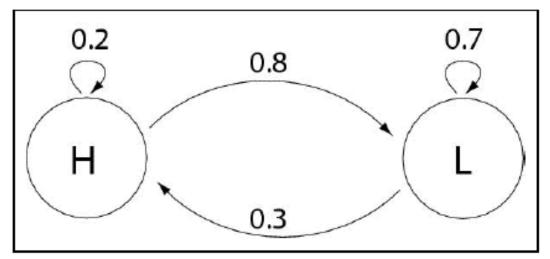
$$P(x) = P(x_{L}, ..., x_{1}, x_{0})$$
  
=  $P(x_{L} | x_{L-1}, ..., x_{0})P(x_{L-1} | x_{L-2}, ..., x_{0}) \cdots P(x_{0})$   
=  $P(x_{L} | x_{L-1})P(x_{L-1} | x_{L-2}) \cdots P(x_{0})$   
=  $\mathbf{b}_{x_{0}} \prod_{i=1}^{L} a_{x_{i-1}x_{i}}$ 



Assume we are modeling a time series of high and low pressures during the Danish autumn.

Let 
$$S = \{H, L\}, \mathbf{b} = \pi = \begin{bmatrix} \frac{3}{11}, \frac{8}{11} \end{bmatrix}, \text{ and } A = \begin{bmatrix} 0.2 & 0.8 \\ 0.3 & 0.7 \end{bmatrix}.$$

#### Graphical representation of A

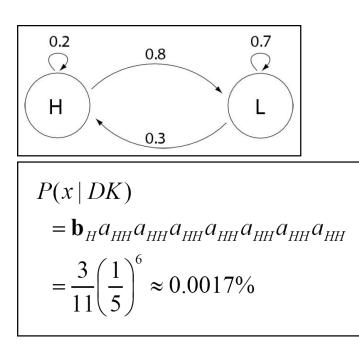


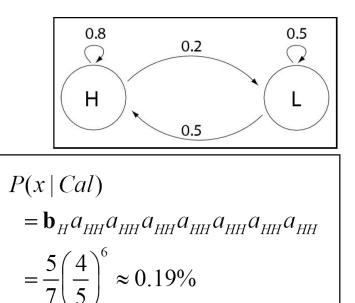


**Comparing likelihoods** 

We want to know the likelihood of one week of high pressure in Denmark (DK) versus California (Cal).

х=ННННННН





# Motivation of Hidden Markov Models



#### Hidden states

- The state of the entity we want to model is often not observable:
  - The state is then said to be hidden.

#### Observables

• Sometimes we can instead observe the state of entities influenced by the hidden state.

#### • A system can be modeled by an HMM if:

- The sequence of hidden states is Markov
- The sequence of observations are independent (or Markov) given the hidden

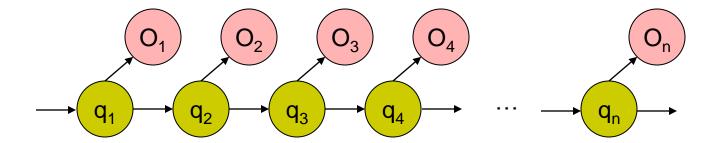
## **Hidden Markov Model**

- Definition  $M = \{S, V, A, B, \pi\}$ 
  - Set of states
  - **Observation symbols**  $V = \{ v_1, v_2, \dots, v_M \}$
  - Transition probabilities
    - A between any two states  $a_{ij} = P(q_t = s_j | q_{t-1} = s_j)$

 $S = \{ S_1, S_2, \dots, S_N \}$ 

- Emission probabilities
  - **B** within each state  $b_j(O_t) = P(O_t = v_j | q_t = s_j)$
- Start probabilities  $\pi = \{a_0\}$

Use  $\lambda = (A, B, \pi)$  to indicate the parameter set of the model.

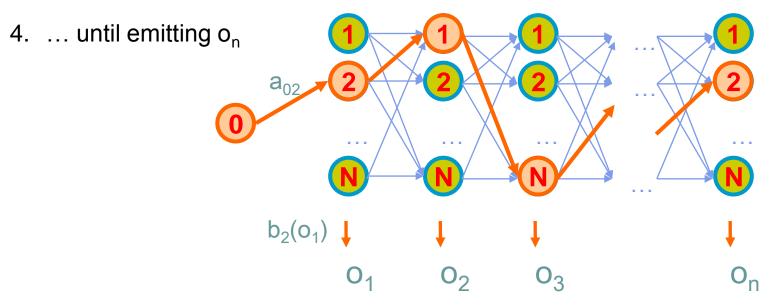




# Generating a sequence by the model

Given a HMM, we can generate a sequence of length n as follows:

- 1. Start at state  $q_1$  according to prob  $a_{0t1}$
- 2. Emit letter  $o_1$  according to prob  $e_{t1}(o_1)$
- 3. Go to state  $q_2$  according to prob  $a_{t1t2}$



## Example

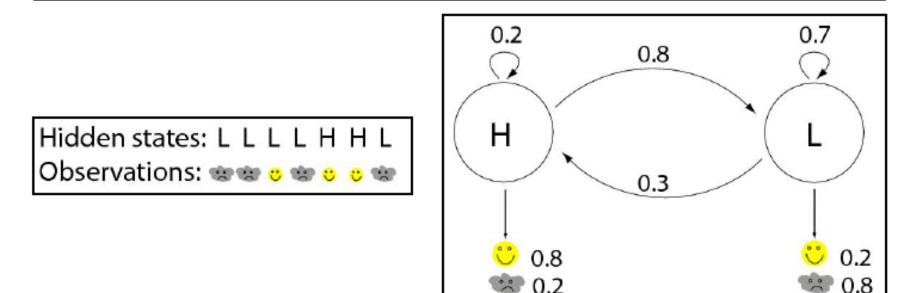


#### Model of high and low pressures

Assume we can not measure high and low pressures.

The state of the weather is influenced by the air pressure.

We make an HMM with hidden states representing high and low pressure and observations representing the weather:



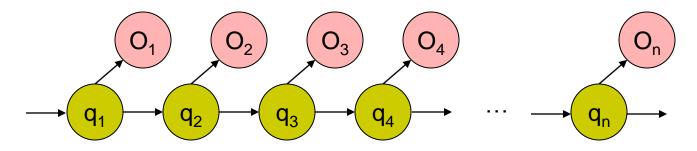
## Calculating with Hidden Markov Model

Consider one such fixed state sequence

$$Q = q_1 q_2 \cdots q_T$$

The observation sequence O for the Q is

$$P(O | Q, \lambda) = \prod_{t=1}^{T} P(O_t | q_t, \lambda)$$
  
=  $b_{q_1}(O_1) \cdot b_{q_2}(O_2) \cdots b_{q_T}(O_T)$ 





### Calculating with Hidden Markov Model (cont.)

The probability of such a state sequence Q

$$P(Q \mid \lambda) = a_{0q_1} a_{q_1q_2} \cdot a_{q_2q_3} \cdots a_{q_{T-1}q_T}$$

The probability that O and Q occur simultaneously, is simply the product of the above two terms, i.e.,

 $P(O, Q \mid \lambda) = P(O \mid Q, \lambda)P(Q \mid \lambda)$ 

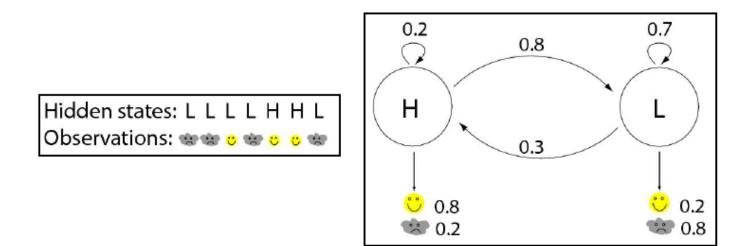
$$P(O, Q \mid \lambda) = a_{0q_1} b_{q_1}(O_1) a_{q_1q_2} b_{q_2}(O_2) a_{q_2q_3} \cdots a_{q_{T-1}q_T} b_{q_T}(O_T)$$



### Example



$$\begin{split} P(x,\pi) &= \left(a_{0L}e_L(R)\right) \left(a_{LL}e_L(R)\right) \left(a_{LL}e_L(S)\right) \left(a_{LL}e_L(R)\right) \left(a_{LH}e_H(S)\right) \left(a_{HH}e_H(S)\right) \left(a_{HL}e_L(R)\right) \\ &= \left(\frac{8}{11}\frac{8}{10}\right) \left(\frac{7}{10}\frac{8}{10}\right) \left(\frac{7}{10}\frac{2}{10}\right) \left(\frac{7}{10}\frac{8}{10}\right) \left(\frac{3}{10}\frac{8}{10}\right) \left(\frac{2}{10}\frac{8}{10}\right) \left(\frac{8}{10}\frac{8}{10}\right) \\ &= 0.0006278 \end{split}$$



# The three main questions on HMMs



#### 1. **Evaluation**

**GIVEN** a HMM  $M=(S, V, A, B, \pi)$ , and a sequence O, **FIND** P[O|M]

#### 2. Decoding

**GIVEN** a HMM  $M=(S, V, A, B, \pi)$ , and a sequence O, **FIND** the sequence Q of states that maximizes  $P(O, Q | \lambda)$ 

### 3. Learning

GIVENa HMM  $M=(S, V, A, B, \pi)$ , with unspecified<br/>transition/emission probabilities and a sequence Q,<br/>parameters  $\theta = (e_i(.), a_{ij})$  that maximize  $P[x|\theta]$ 

# **Evaluation**

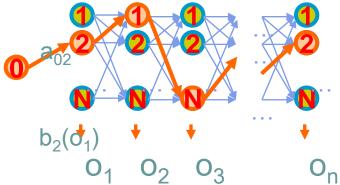


- Find the likelihood a sequence is generated by the model
- A straightforward way (穷举法)
  - The probability of O is obtained by summing all possible state sequences q giving

$$P(O \mid \lambda) = \sum_{all \ Q} P(O \mid Q, \lambda) P(Q \mid \lambda)$$

$$= \sum_{q_1,q_2,\dots,q_T} \pi_{q_1} b_{q_1} (O_1) a_{q_1q_2} b_{q_2} (O_2) a_{q_2q_3} \cdots a_{q_{T-1}q_T} b_{q_T} (O_T)$$

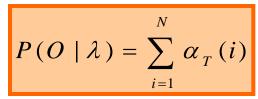
Complexity is *O(N<sup>T</sup>)* Calculations is unfeasible

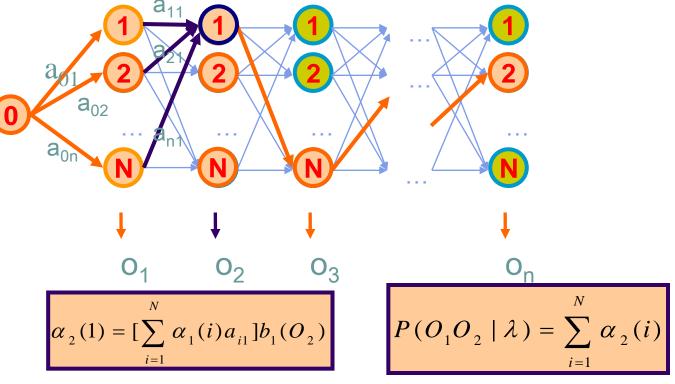




# **The Forward Algorithm**

- A more elaborate algorithm
  - The Forward Algorithm





# **The Forward Algorithm**

The Forward variable

$$\alpha_{t}(i) = P(O_{1}O_{2} \cdots O_{t}, q_{t} = S_{i} | \lambda)$$

We can compute  $\alpha(i)$  for all *N*, *i*, <u>Initialization:</u>  $\alpha_1(i) = a_{0i}b_{0i}(O_1)$  i = 1...N

Ν

**Iteration:** 

$$\alpha_{t+1}(i) = \left[\sum_{i=1}^{n} \alpha_{t}(i)a_{ij}\right]b_{j}(O_{t+1})$$

**Termination:** 

$$P(O \mid \lambda) = \sum_{i=1}^{N} \alpha_{T}(i)$$

 $t = 1 \dots T - 1$   $1 = 1 \dots T - 1$   $2 = 1 \dots T - 1$   $1 = 1 \dots T - 1$ 



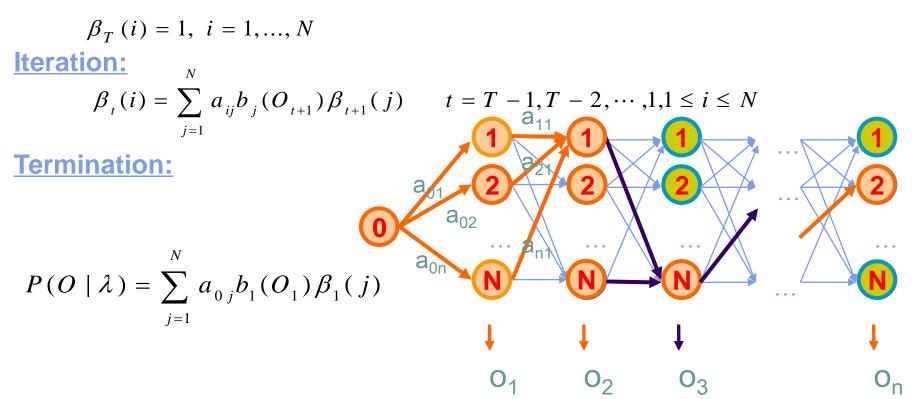


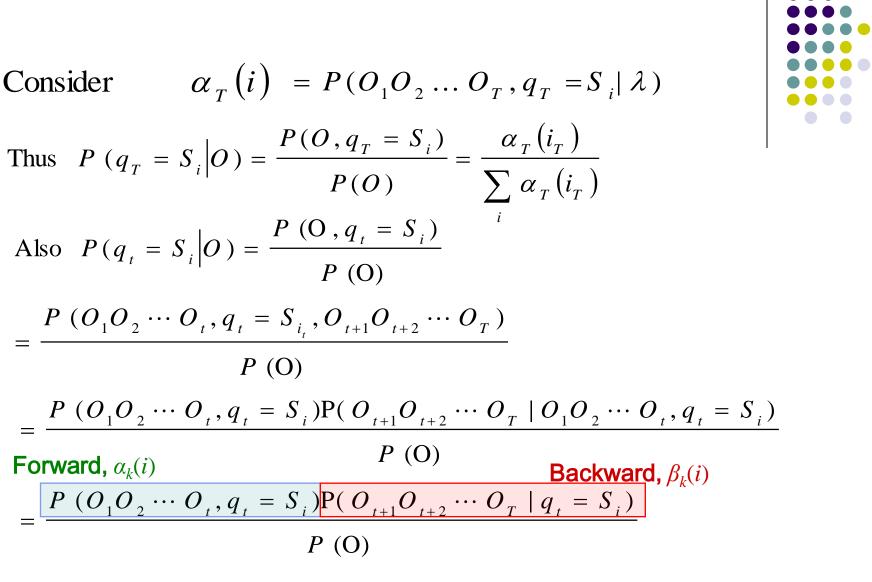
# **The Backward Algorithm**

The backward variable

 $\beta_t(i) = P(O_{t+1}O_{t+2} \cdots O_T | q_t = S_t, \lambda)$ Similar, we can compute backward variable for all *N*, *i*,

**Initialization:** 





 $=\frac{\alpha_{t}(i)\beta_{t}(i)}{\sum_{i}\alpha_{T}(i)}=\gamma(i)$ 

# Decoding



### **GIVEN** a HMM, and a sequence *O*.

Suppose that we know the parameters of the Hidden Markov Model and the observed sequence of observations  $O_1, O_2, \dots, O_T$ .

# **FIND** the sequence Q of states that maximizes $P(Q|O,\lambda)$

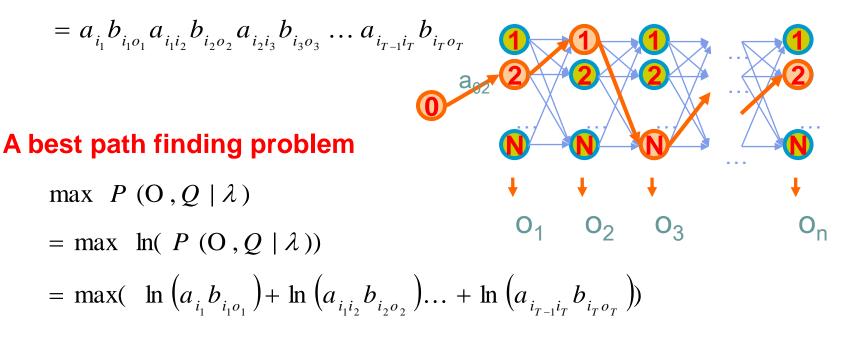
Determining the sequence of States  $q_1, q_2, ..., q_T$ , which is optimal in some meaningful sense. (i.e. best "explain" the observations)

# Decoding



Consider  $P(Q|O,\lambda) = \frac{P(O,Q|\lambda)}{P(O|\lambda)}$ 

To maximize the above probability is equivalent to maximizing  $P(O, Q | \lambda)$ 



# Viterbi Algorithm

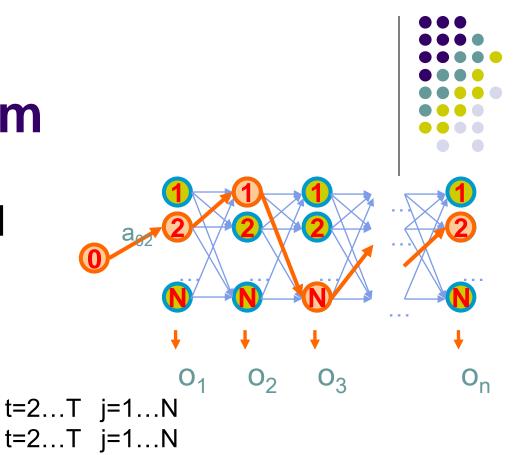
### [Dynamic programming]

Initialization:

$$\begin{split} \delta_{1}(i) &= a_{0i}b_{i}(O_{1}) , \quad i = 1...N \\ \psi_{1}(i) &= 0. \end{split}$$
 **Recursion:**  $\delta_{t}(j) &= \max_{i} \left[ \delta_{t-1}(i) \ a_{ij} \right] b_{j}(O_{t}) \\ \psi_{1}(j) &= \operatorname{argmax}_{i} \left[ \delta_{t-1}(i) \ a_{ij} \right] \end{split}$  **Termination:**  $P^{*} &= \max_{i} \delta_{T}(i) \\ q_{T}^{*} &= \operatorname{argmax}_{i} \left[ \delta_{T}(i) \right] \end{split}$ 

**Traceback:** 

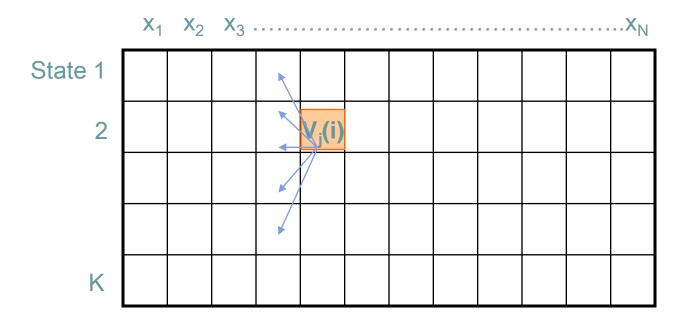
 $q_t^* = \psi_1(q_{t+1}^*)$ 



t=T-1,T-2,...,1.



# **The Viterbi Algorithm**



Similar to "aligning" a set of states to a sequence

 Time:
 O(K<sup>2</sup>N)

 Space:
 O(KN)

# Learning



- Estimation of Parameters of a Hidden Markov Model
  - 1. Both the sequence of observations O and the sequence of states Q is observed

learning  $\lambda = (A, B, \pi)$ 

2. Only the sequence of observations O are observed

learning Q and  $\lambda = (A, B, \pi)$ 



$$L(A, B, \pi) = a_{i_1} b_{i_1 o_1} a_{i_1 i_2} b_{i_2 o_2} a_{i_2 i_3} b_{i_3 o_3} \dots a_{i_{T-1} i_T} b_{i_T o_T}$$



• the log-Likelihood is:

$$l(A, B, \pi) = \ln L(A, B, \pi) = \ln (a_{i_1}) + \ln (b_{i_1 o_1}) + \ln (a_{i_1 i_2}) + \ln (a_{i_2 i_3}) + \ln (b_{i_3 o_3}) \dots + \ln (a_{i_{T-1} i_T}) + \ln (b_{i_T o_T}) = \sum_{i=1}^{M} f_{i0} \ln (a_i) + \sum_{i=1}^{M} \sum_{j=1}^{M} f_{ij} \ln (a_{ij}) + \sum_{i=1}^{M} \sum_{o(i)} \ln (b_{io})$$

where  $f_{i0}$  = the number of times state *i* occurs in the first state  $f_{ij}$  = the number of times state *i* changes to state *j*.  $\beta_{iy} = f(y | \theta_i) \text{ (or } p(y | \theta_i) \text{ in the discrete case})$  $\sum_{o(i)} \bullet =$  the sum of all observations  $o_t$  where  $q_t = S_i$ 



In such case these parameters computed by *Maximum Likelihood Estimation* are:

$$\hat{a}_{i} = \frac{f_{i0}}{1}$$
  $\hat{a}_{ij} = \frac{f_{ij}}{M}$ , and  $\sum_{j=1}^{M} f_{ij}$ 

 $\hat{b}_i$  = the MLE of  $b_i$  computed from the observations  $o_t$  where  $q_t = S_i$ .

Only the sequence of observations O are observed

$$L(A, B, \pi) = \sum_{i_1, i_2 \dots i_T} a_{i_1} b_{i_1 o_1} a_{i_1 i_2} b_{i_2 o_2} a_{i_2 i_3} b_{i_3 o_3} \dots a_{i_{T-1} i_T} b_{i_T o_T}$$

- It is difficult to find the Maximum Likelihood Estimates directly from the Likelihood function.
- The Techniques that are used are
  - 1. The Segmental K-means Algorith
  - 2. The Baum-Welch (E-M) Algorithm



- The E-M algorithm was designed originally to handle "Missing observations".
- In this case the missing observations are the states  $\{q_1, q_2, \dots, q_T\}$ .
- Assuming a model, the states are estimated by finding their expected values under this model. (The E part of the E-M algorithm).



- With these values the model is estimated by Maximum Likelihood Estimation (The M part of the E-M algorithm).
- The process is repeated until the estimated model converges.

#### Initialization:

Pick the best-guess for model parameters (or arbitrary)

#### Iteration:

Forward Backward Calculate  $A_{kl}$ ,  $E_k(b)$ Calculate new model parameters  $a_{kl}$ ,  $e_k(b)$ Calculate new log-likelihood  $P(x | \theta)$ 

#### **GUARANTEED TO BE HIGHER BY EXPECTATION-MAXIMIZATION**

Until  $P(x \mid \theta)$  does not change much



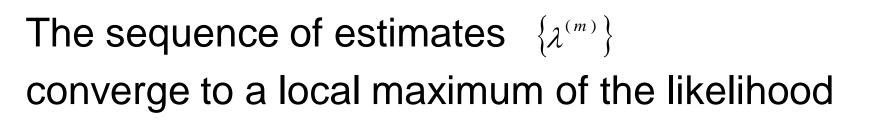


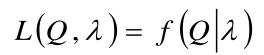
Let  $f(O,Q|\lambda) = L(O,Q,\lambda)$  denote the joint distribution of Q,O. Consider the function:

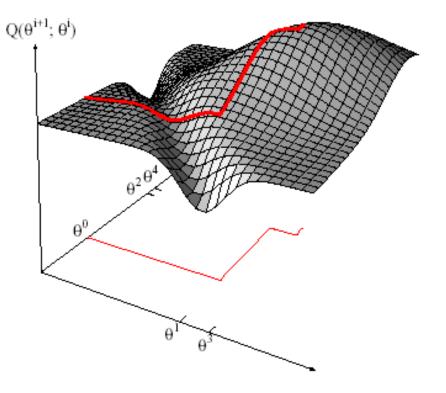
$$Q(\lambda, \lambda') = E_{\mathbf{X}}(\ln L(O, Q, \lambda) | Q, \lambda')$$

Starting with an initial estimate of  $\lambda$  ( $\lambda^{(1)}$ ).

A sequence of estimates  $\{\lambda^{(m)}\}\$  are formed by finding  $\lambda = \lambda^{(m+1)}$  to maximize  $Q(\lambda, \lambda^{(m)})$ with respect to  $\lambda$ .







# **Speech Recognition**



- On-line documents of Java<sup>™</sup> Speech API
  - <a href="http://java.sun.com/products/java-media/speech/">http://java.sun.com/products/java-media/speech/</a>
- On-line documents of Free TTS
  - http://freetts.sourceforge.net/docs/
- On-line documents of Sphinx-II
  - http://www.speech.cs.cmu.edu/sphinx/

# Speech Recognition and Mobile applications

- Siri
- Google speech
- WeChat / Laiwang





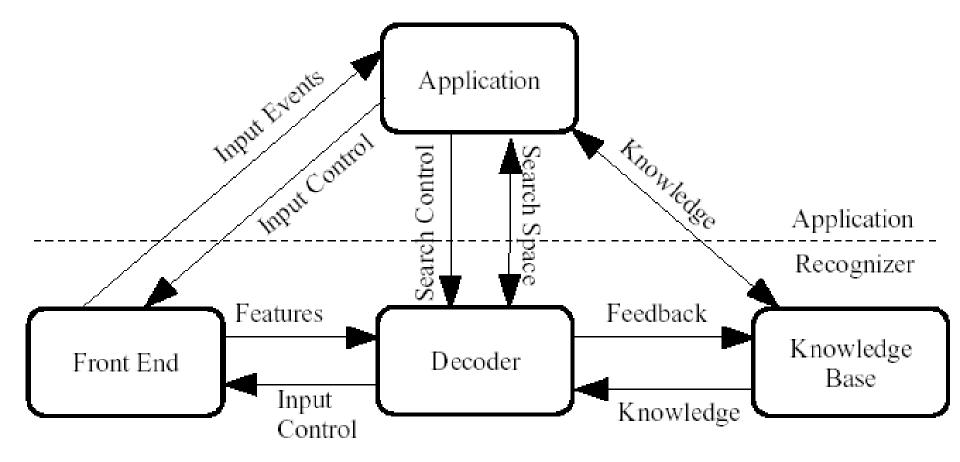
# **Brief History of CMU Sphinx**



- Sphinx-I (1987)
  - The first user independent, high performance ASR of the world.
  - Written in C by Kai-Fu Lee (李開復博士,現任Google副總裁).
- Sphinx-II (1992)
  - Written by Xuedong Huang in C. (黃學東博士,現為Microsoft Speech.NET團 隊領導人)
  - 5-state HMM / N-gram LM.
- Sphinx-III (1996)
  - Built by Eric Thayer and Mosur Ravishankar.
  - Slower than Sphinx-II but the design is more flexible.
- Sphinx-4 (Originally Sphinx 3j)
  - Refactored from Sphinx 3.
  - Fully implemented in Java. (Not finished yet ...)

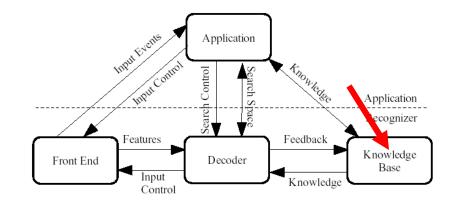


# **Components of CMU Sphinx**



### **Knowledge Base**

- The data that drives the decoder.
- Three sets of data
  - Acoustic Model.
  - Language Model.
  - Lexicon (Dictionary).





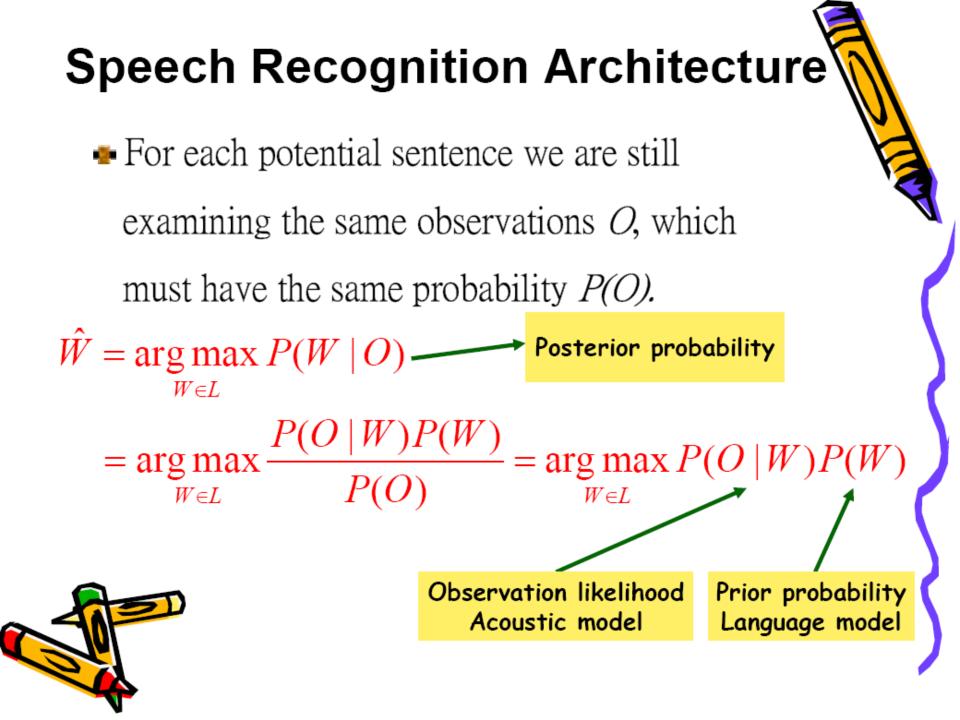
# Speech Recognition Architecture

- Observations :  $O = o_1, o_2, o_3, \dots, o_t$
- Word Sequences :  $W = w_1, w_2, w_3, \dots, w_n$

• Probabilistic implementation can be expressed :  $\hat{W} = \underset{W \in L}{\operatorname{arg\,max}} P(W \mid O)$ 

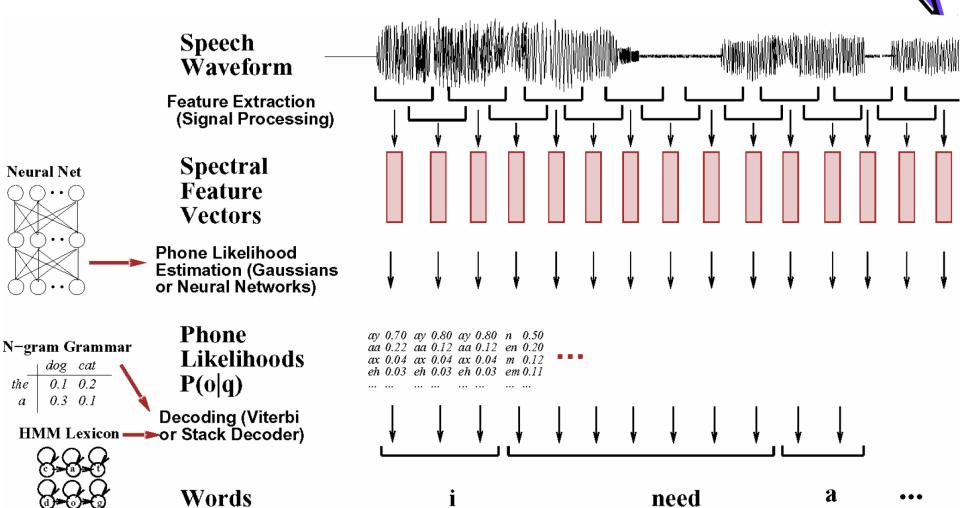
Then we can use Bayes' rule to break it down :

 $\hat{W} = \underset{W \in L}{\operatorname{arg\,max}} P(W \mid O) = \underset{W \in L}{\operatorname{arg\,max}} \frac{P(O \mid W)P(W)}{P(O)}$   $P(O) = \frac{P(WO)}{P(O)} \quad \text{and} \quad P(O \mid W) = \frac{P(WO)}{P(W)}$   $P(W \mid O) \cdot P(O) = P(WO) = P(O \mid W) \cdot P(W)$ 



# Speech Recognition Architecture

Figure 7.2 Schematic architecture for a speech recognition



# **Acoustic Model**



- /model/hmm/6k
- Database of statistical model.
- Each statistical model represents a phoneme.
- Acoustic Models are trained by analyzing large amount of speech data.

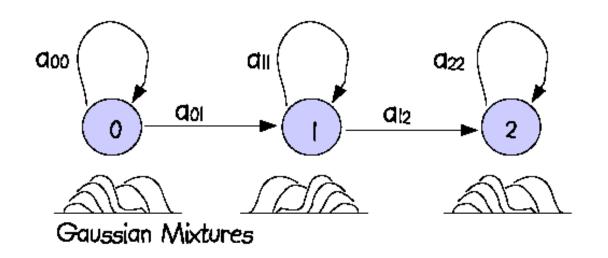
# **HMM in Acoustic Model**



- HMM represent each unit of speech in the Acoustic Model.
- Typical HMM use 3-5 states to model a phoneme.
- Each state of HMM is represented by a set of *Gaussian mixture density functions*.
- Sphinx2 default phone set.

# **Mixture of Gaussians**

- Represent each state in HMM.
- Each set of Gaussian Mixtures are called "senones".
- HMM can share "senones".





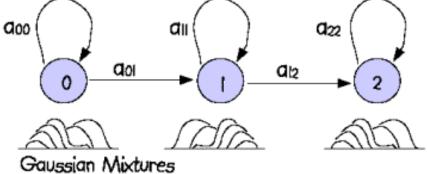


# **Mixture of Gaussians**

$$N(x;\mu,\Sigma) = \frac{1}{(2\pi)^{n/2}} \exp\left[-\frac{1}{2}(x-\mu)^{t}\Sigma^{-1}(x-\mu)\right]$$

$$f(\mathbf{x}) = \sum_{k=1}^{K} c_k N_k(\mathbf{x}; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \qquad \texttt{H} \mathbf{p}$$
$$c_k \ge 0 \quad \texttt{H} \quad \sum_{k=1}^{K} c_k = 1$$

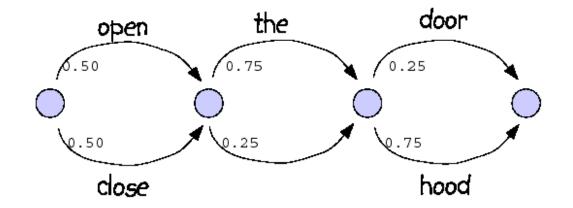
Gaussian mixtures with enough mixture components can approximate any distribution.



# Language Model

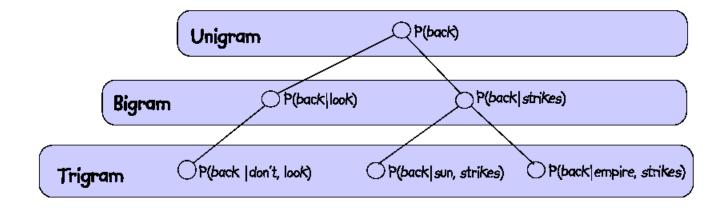


- Describes what is likely to be spoken in a particular context
- Word transitions are defined in terms of transition probabilities
- Helps to constrain the search space



# **N-gram Language Model**

- Probability of word N dependent on word N-1, N-2, ...
- Bigrams and trigrams most commonly used
- Used for large vocabulary applications such as dictation
- Typically trained by very large (millions of words) corpus

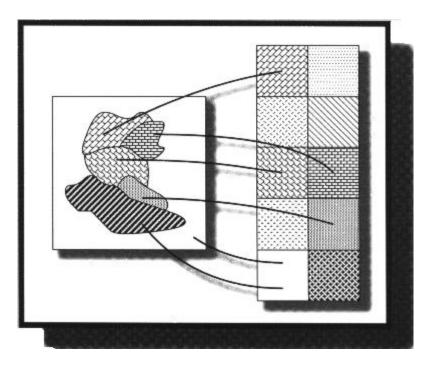




# Markov Random field and CRF



- See webpage
- http://www.nlpr.ia.ac.cn/users/szli/MRF\_Book /MRF\_Book.html



# **Belief Network (Propagation)**



Y. Weiss and W. T. Freeman *Correctness of Belief Propagation in Gaussian Graphical Models of Arbitrary Topology.* in: Advances in Neural Information Processing Systems 12, edited by S. A. Solla, T. K. Leen, and K-R Muller, 2000. <u>MERL-TR99-38.</u> (y<sub>3</sub>) (y<sub>4</sub>) (y<sub>2</sub>)





