

# Machine Learning

10-701/15-781, Spring 2010

## Hidden Markov Model (II)

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Lecture 11, February 24, 2010



Reading: Chap. 13 CB

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## Three Main Questions on HMMs



### 1. Evaluation

GIVEN an HMM  $M$ , and a sequence  $x$ ,  
FIND Prob ( $x | M$ )  
ALGO. Forward

### 2. Decoding

GIVEN an HMM  $M$ , and a sequence  $x$ ,  
FIND the sequence  $y$  of states that maximizes, e.g.,  $P(y | x, M)$ ,  
or the most probable subsequence of states  
ALGO. Viterbi, Forward-backward

### 3. Learning

GIVEN an HMM  $M$ , with unspecified transition/emission probs.,  
and a sequence  $x$ ,  
FIND parameters  $\theta = (\pi_i, a_{ij}, \eta_{ik})$  that maximize  $P(x | \theta)$   
ALGO. Baum-Welch (EM)

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## Example:

$x = 1, 2, 1, 5, 6, 2, 1, 6, 2, 4$



$$\begin{aligned} P(1|F) &= 1/6 & P(1|L) &= 1/10 \\ P(2|F) &= 1/6 & P(2|L) &= 1/10 \\ P(3|F) &= 1/6 & P(3|L) &= 1/10 \\ P(4|F) &= 1/6 & P(4|L) &= 1/10 \\ P(5|F) &= 1/6 & P(5|L) &= 1/10 \\ P(6|F) &= 1/6 & P(6|L) &= 1/2 \end{aligned}$$

$$\begin{aligned} \alpha_t^k &= P(x_t | y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k} \\ \beta_t^k &= \sum_i a_{k,i} P(x_{t+1} | y_{t+1}^i = 1) \beta_{t+1}^i \end{aligned}$$

$$P(y_t^k = 1 | x) = \frac{P(y_t^k = 1, x)}{P(x)} = \frac{\alpha_t^k \beta_t^k}{P(x)}$$

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$x = 1, 2, 1, 5, 6, 2, 1, 6, 2, 4$



### Alpha (actual)

0.0833	0.0500
0.0136	0.0052
0.0022	0.0006
0.0004	0.0001
0.0001	0.0000
0.0000	0.0000
0.0000	0.0000
0.0000	0.0000
0.0000	0.0000
0.0000	0.0000

### Beta (actual)

0.0000	0.0000
0.0000	0.0000
0.0045	0.0055
0.0264	0.0112
0.1633	0.1033
1.0000	1.0000

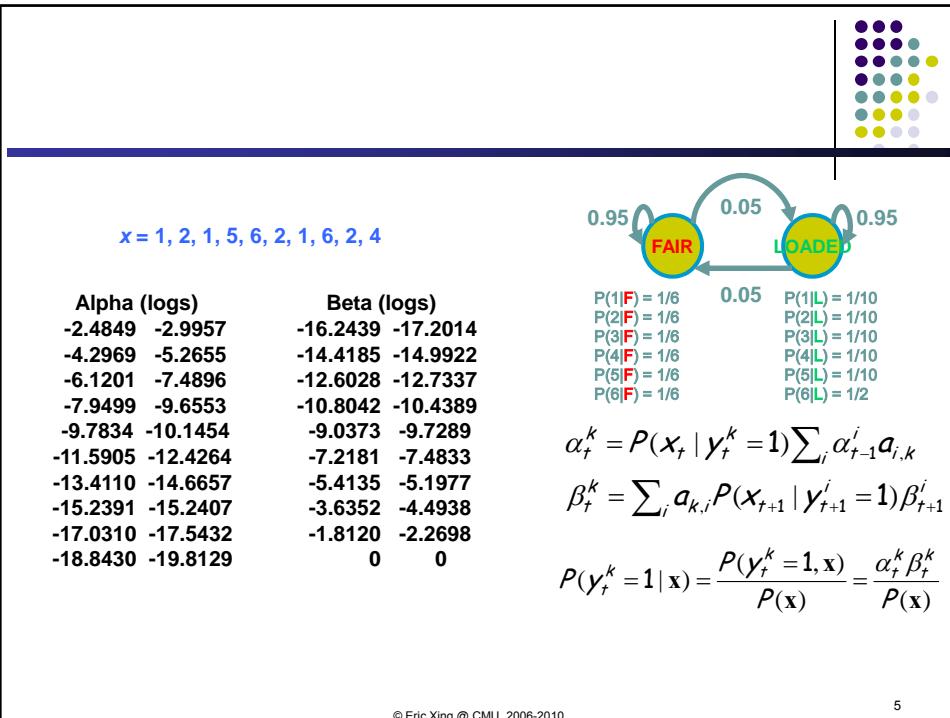
$$\begin{aligned} P(1|F) &= 1/6 & P(1|L) &= 1/10 \\ P(2|F) &= 1/6 & P(2|L) &= 1/10 \\ P(3|F) &= 1/6 & P(3|L) &= 1/10 \\ P(4|F) &= 1/6 & P(4|L) &= 1/10 \\ P(5|F) &= 1/6 & P(5|L) &= 1/10 \\ P(6|F) &= 1/6 & P(6|L) &= 1/2 \end{aligned}$$

$$\begin{aligned} \alpha_t^k &= P(x_t | y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k} \\ \beta_t^k &= \sum_i a_{k,i} P(x_{t+1} | y_{t+1}^i = 1) \beta_{t+1}^i \end{aligned}$$

$$P(y_t^k = 1 | x) = \frac{P(y_t^k = 1, x)}{P(x)} = \frac{\alpha_t^k \beta_t^k}{P(x)}$$

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## What is the probability of a hidden state prediction?

- A single state:

$$P(y_t|\mathbf{X})$$

- What about a hidden state sequence ?

$$P(y_1, \dots, y_T|\mathbf{X})$$

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## Posterior decoding

- We can now calculate

$$P(y_t^k = 1 | \mathbf{x}) = \frac{P(y_t^k = 1, \mathbf{x})}{P(\mathbf{x})} = \frac{\alpha_t^k \beta_t^k}{P(\mathbf{x})}$$

- Then, we can ask

- What is the most likely state at position  $t$  of sequence  $\mathbf{x}$ :

$$k_t^* = \arg \max_k P(y_t^k = 1 | \mathbf{x})$$

- Note that this is an MPA of a **single hidden state**, what if we want to a MPA of a whole hidden state sequence?

- Posterior Decoding:  $\{y_t^{k_t^*} = 1 : t = 1 \dots T\}$

- This is different from MPA of a **whole sequence states**

- This can be understood as **bit error rate** vs. **word error rate**

of hidden

$x$	$y$	$P(x, y)$
0	0	0.35
0	1	0.05
1	0	0.3
1	1	0.3

Example:  
MPA of  $X$ ?  
MPA of  $(X, Y)$ ?

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## Viterbi decoding

- GIVEN  $\mathbf{x} = x_1, \dots, x_T$ , we want to find  $\mathbf{y} = y_1, \dots, y_T$ , such that  $P(\mathbf{y}|\mathbf{x})$  is maximized:

$$y^* = \operatorname{argmax}_y P(\mathbf{y}|\mathbf{x}) = \operatorname{argmax}_{\pi} P(\mathbf{y}, \mathbf{x})$$

- Let

$$V_t^k = \max_{\{y_1, \dots, y_{t-1}\}} P(x_1, \dots, x_{t-1}, y_1, \dots, y_{t-1}, x_t, y_t^k = 1)$$

= Probability of most likely **sequence of states** ending at state  $y_t = k$

- The recursion:

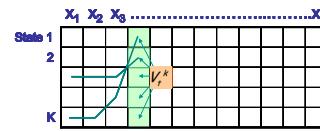
$$V_t^k = p(x_t | y_t^k = 1) \max_i a_{i,k} V_{t-1}^i$$

- Underflows are a significant problem

$$p(x_1, \dots, x_t, y_1, \dots, y_t) = \pi_{y_1} a_{y_1, y_2} \cdots a_{y_{t-1}, y_t} b_{y_t, x_t}$$

- These numbers become extremely small – underflow

- Solution: Take the logs of all values:  $V_t^k = \log p(x_t | y_t^k = 1) + \max_i (\log(a_{i,k}) + V_{t-1}^i)$



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## The Viterbi Algorithm – derivation



- Define the viterbi probability:

$$\begin{aligned}
 V_{t+1}^k &= \max_{\{y_1, \dots, y_t\}} P(x_1, \dots, x_t, y_1, \dots, y_t, x_{t+1}, y_{t+1}^k = 1) \\
 &= \max_{\{y_1, \dots, y_t\}} P(x_{t+1}, y_{t+1}^k = 1 | x_1, \dots, x_t, y_1, \dots, y_t) P(x_1, \dots, x_t, y_1, \dots, y_t) \\
 &= \max_{\{y_1, \dots, y_t\}} P(x_{t+1}, y_{t+1}^k = 1 | y_t) P(x_1, \dots, x_{t-1}, y_1, \dots, y_{t-1}, x_t, y_t) \\
 &= \max_i P(x_{t+1}, y_{t+1}^k = 1 | y_t^i = 1) \max_{\{y_1, \dots, y_{t-1}\}} P(x_1, \dots, x_{t-1}, y_1, \dots, y_{t-1}, x_t, y_t^i = 1) \\
 &= \max_i P(x_{t+1} | y_{t+1}^k = 1) a_{i,k} V_t^i \\
 &= P(x_{t+1} | y_{t+1}^k = 1) \max_i a_{i,k} V_t^i
 \end{aligned}$$

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## The Viterbi Algorithm



- Input:  $x = x_1, \dots, x_T$

### Initialization:

$$V_1^k = P(x_1 | y_1^k = 1) \pi_k$$

### Iteration:

$$\begin{aligned}
 V_t^k &= P(x_t, | y_t^k = 1) \max_i a_{i,k} V_{t-1}^i \\
 \text{Ptr}(k, t) &= \arg \max_i a_{i,k} V_{t-1}^i
 \end{aligned}$$

### Termination:

$$P(x, y^*) = \max_k V_T^k$$

### TraceBack:

$$\begin{aligned}
 y_T^* &= \arg \max_k V_T^k \\
 y_{t-1}^* &= \text{Ptr}(y_t^*, t)
 \end{aligned}$$

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## Viterbi Vs. MPA (individual)

$V_t^k$ (log)	$ptr(k, t)$	Seq	Viterbi	MPA	$p(y_t^k = 1   x)$
-2.4849 -2.9957	N/A	1	1	1	0.8128 0.1872
-4.3280 -5.3496	1 2	2	1	1	0.8238 0.1762
-6.1710 -7.7035	1 2	1	1	1	0.8176 0.1824
-8.0141 -10.0574	1 2	5	1	1	0.7925 0.2075
-9.8571 -10.8018	1 2	6	1	1	0.7415 0.2585
-11.7002 -13.1557	1 2	2	1	1	0.7505 0.2495
-13.5432 -15.5096	1 2	1	1	1	0.7386 0.2614
-15.3863 -16.2540	1 2	6	1	1	0.7027 0.2973
-17.2293 -18.6079	1 2	2	1	1	0.7251 0.2749
-19.0724 -20.9618	1 2	4	1	1	0.7251 0.2749

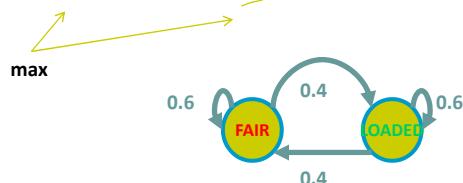
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## Another Example

X = 6, 2, 3, 5, 6, 2, 6, 3, 6, 6

$V_t^k$ (log)	$ptr(k, t)$	Seq	Viterbi	MPA	$p(y_t^k = 1   x)$
-2.4849 -1.3863	N/A	6	2	2	0.2733 0.7267
-4.0943 -4.1997	2 2	2	1	1	0.6040 0.3960
-6.3969 -7.0131	1 2	3	1	1	0.6538 0.3462
-8.6995 -9.6158	1 1	5	1	1	0.6062 0.3938
-11.0021 -10.3090	1 1	6	2	2	0.2861 0.7139
-13.0170 -13.1224	2 2	2	2	1	0.5342 0.4658
-15.3196 -14.3263	1 2	6	2	2	0.2734 0.7266
-17.0344 -17.1397	2 2	3	2	1	0.5226 0.4774
-19.3370 -18.3437	1 2	6	2	2	0.2252 0.7748
-21.0518 -19.5477	2 2	6	2	2	0.2159 0.7841



Same transition probabilities

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## Computational Complexity and implementation details



- What is the running time, and space required, for Forward, and Backward?

$$\begin{aligned}\alpha_t^k &= p(x_t | y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k} \\ \beta_t^k &= \sum_i a_{k,i} p(x_{t+1} | y_{t+1}^i = 1) \beta_{t+1}^i \\ V_t^k &= p(x_t | y_t^k = 1) \max_i a_{i,k} V_{t-1}^i\end{aligned}$$

Time:  $O(K^2N)$ ; Space:  $O(KN)$ .

- Useful implementation technique to avoid underflows

- Viterbi: sum of logs
- Forward/Backward: rescaling at each position by multiplying by a constant

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 ALGO. Baum-Welch (EM)

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## Learning HMM: two scenarios

- **Supervised learning:** estimation when the “right answer” is known
  - **Examples:**
    - GIVEN:** a genomic region  $x = x_1 \dots x_{1,000,000}$  where we have good (experimental) annotations of the CpG islands
    - GIVEN:** the casino player allows us to observe him one evening, as he changes dice and produces 10,000 rolls
- **Unsupervised learning:** estimation when the “right answer” is unknown
  - **Examples:**
    - GIVEN:** the porcupine genome; we don't know how frequent are the CpG islands there, neither do we know their composition
    - GIVEN:** 10,000 rolls of the casino player, but we don't see when he changes dice
- **QUESTION:** Update the parameters  $\theta$  of the model to maximize  $P(x|\theta)$  --- Maximal likelihood (ML) estimation

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## MLE

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## Supervised ML estimation

- Given  $x = x_1 \dots x_N$  for which the true state path  $y = y_1 \dots y_N$  is known,

- Define:

$$\begin{aligned} A_{ij} &= \# \text{ times state transition } i \rightarrow j \text{ occurs in } y \\ B_{ik} &= \# \text{ times state } i \text{ in } y \text{ emits } k \text{ in } x \end{aligned}$$

- We can show that the maximum likelihood parameters  $\theta$  are:

$$a_{ij}^{ML} = \frac{\#(i \rightarrow j)}{\#(i \rightarrow \bullet)} = \frac{\sum_n \sum_{t=2}^T Y_{n,t-1}^i Y_{n,t}^j}{\sum_n \sum_{t=2}^T Y_{n,t-1}^i} = \frac{A_{ij}}{\sum_j A_{ij}}$$

$$b_{ik}^{ML} = \frac{\#(i \rightarrow k)}{\#(i \rightarrow \bullet)} = \frac{\sum_n \sum_{t=1}^T Y_{n,t}^i X_{n,t}^k}{\sum_n \sum_{t=1}^T Y_{n,t}^i} = \frac{B_{ik}}{\sum_k B_{ik}}$$

- What if  $y$  is continuous? We can treat  $\{(x_{n,t}, y_{n,t}) : t = 1:T, n = 1:N\}$  as  $N \times T$  observations of, e.g., a Gaussian, and apply learning rules for Gaussian ...

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## Supervised ML estimation, ctd.

- Intuition:**

- When we know the underlying states, the best estimate of  $\theta$  is the average frequency of transitions & emissions that occur in the training data

- Drawback:**

- Given little data, there may be **overfitting**:
  - $P(x|\theta)$  is maximized, but  $\theta$  is unreasonable
  - 0 probabilities – VERY BAD

- Example:**

- Given 10 casino rolls, we observe

$x = 2, 1, 5, 6, 1, 2, 3, 6, 2, 3$   
 $y = F, F, F, F, F, F, F, F, F, F$

- Then:  
 $a_{FF} = 1; a_{FL} = 0$   
 $b_{F1} = b_{F3} = .2;$   
 $b_{F2} = .3; b_{F4} = 0; b_{F5} = b_{F6} = .1$

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## Pseudocounts

- Solution for small training sets:

- Add pseudocounts

$$A_{ij} = \# \text{ times state transition } i \rightarrow j \text{ occurs in } y + R_{ij}$$

$$B_{ik} = \# \text{ times state } i \text{ in } y \text{ emits } k \text{ in } x + S_{ik}$$

- $R_{ij}$ ,  $S_{ik}$  are pseudocounts representing our prior belief
  - Total pseudocounts:  $R_i = \sum_j R_{ij}$ ,  $S_i = \sum_k S_{ik}$ ,
  - --- "strength" of prior belief,
    - --- total number of imaginary instances in the prior

- Larger total pseudocounts  $\Rightarrow$  strong prior belief

- Small total pseudocounts: just to avoid 0 probabilities --- smoothing

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## Unsupervised ML estimation

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## Unsupervised ML estimation

- Given  $x = x_1 \dots x_N$  for which the true state path  $y = y_1 \dots y_N$  is unknown,

- EXPECTATION MAXIMIZATION**

- Starting with our best guess of a model  $M$ , parameters  $\theta$ .

- Estimate  $A_{ij}$ ,  $B_{ik}$  in the training data

- How?  $A_{ij} = \sum_{n,t} \langle y_{n,t-1}^i y_{n,t}^j \rangle$     $B_{ik} = \sum_{n,t} \langle y_{n,t}^i \rangle x_{n,t}^k$ ,
- Update  $\theta$  according to  $A_{ij}$ ,  $B_{ik}$
- Now a "supervised learning" problem

- Repeat 1 & 2, until convergence

This is called the Baum-Welch Algorithm

We can get to a provably more (or equally) likely parameter set  $\theta$  each iteration

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## The Baum Welch algorithm

- The complete log likelihood

$$\ell_c(\theta; x, y) = \log p(x, y) = \log \prod_n \left( p(y_{n,1}) \prod_{t=2}^T p(y_{n,t} | y_{n,t-1}) \prod_{t=1}^T p(x_{n,t} | x_{n,t}) \right)$$

- The expected complete log likelihood

$$\langle \ell_c(\theta; x, y) \rangle = \sum_n \left( \langle y_{n,1}^i \rangle_{p(y_{n,1}|x_n)} \log \pi_i \right) + \sum_n \sum_{t=2}^T \left( \langle y_{n,t-1}^i y_{n,t}^j \rangle_{p(y_{n,t-1}, y_{n,t}|x_n)} \log a_{i,j} \right) + \sum_n \sum_{t=1}^T \left( x_{n,t}^k \langle y_{n,t}^i \rangle_{p(y_{n,t}|x_n)} \log b_{i,k} \right)$$

- EM

- The E step

$$\gamma_{n,t}^i = \langle y_{n,t}^i \rangle = p(y_{n,t}^i = 1 | x_n)$$

$$\xi_{n,t}^{i,j} = \langle y_{n,t-1}^i y_{n,t}^j \rangle = p(y_{n,t-1}^i = 1, y_{n,t}^j = 1 | x_n)$$

- The M step ("symbolically" identical to MLE)

$$\pi_i^{ML} = \frac{\sum_n \gamma_{n,1}^i}{N}$$

$$a_{ij}^{ML} = \frac{\sum_n \sum_{t=2}^T \xi_{n,t}^{i,j}}{\sum_n \sum_{t=1}^{T-1} \gamma_{n,t}^i}$$

$$b_k^{ML} = \frac{\sum_n \sum_{t=1}^T \gamma_{n,t}^i x_{n,t}^k}{\sum_n \sum_{t=1}^{T-1} \gamma_{n,t}^i}$$

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## The Baum-Welch algorithm -- comments



Time Complexity:

$$\# \text{ iterations} \times O(K^2N)$$

- Guaranteed to increase the log likelihood of the model
- Not guaranteed to find globally best parameters
- Converges to local optimum, depending on initial conditions
- Too many parameters / too large model: Overt-fitting

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## Summary: the HMM algorithms



Questions:

- **Evaluation:** What is the probability of the observed sequence? **Forward**
- **Decoding:** What is the probability that the state of the 3rd roll is loaded, given the observed sequence? **Forward-Backward**
- **Decoding:** What is the most likely die sequence? **Viterbi**
- **Learning:** Under what parameterization are the observed sequences most probable? **Baum-Welch (EM)**

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## Applications of HMMs

- Some early applications of HMMs

- finance, but we never saw them
- speech recognition
- modelling ion channels

- In the mid-late 1980s HMMs entered genetics and molecular biology, and they are now firmly entrenched.

- Some current applications of HMMs to biology

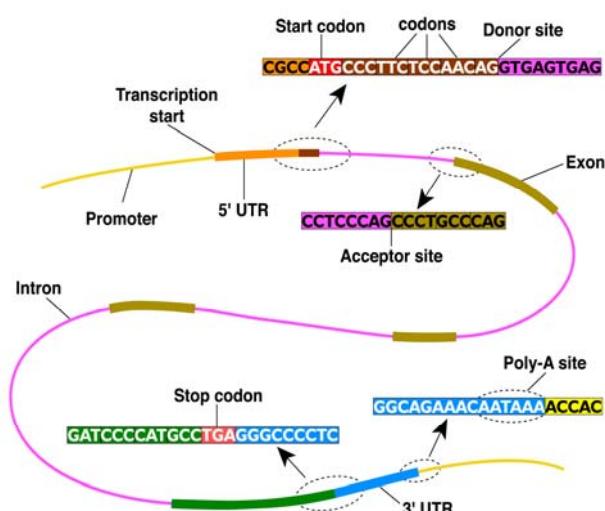
- mapping chromosomes
- aligning biological sequences
- predicting sequence structure
- inferring evolutionary relationships
- finding genes in DNA sequence

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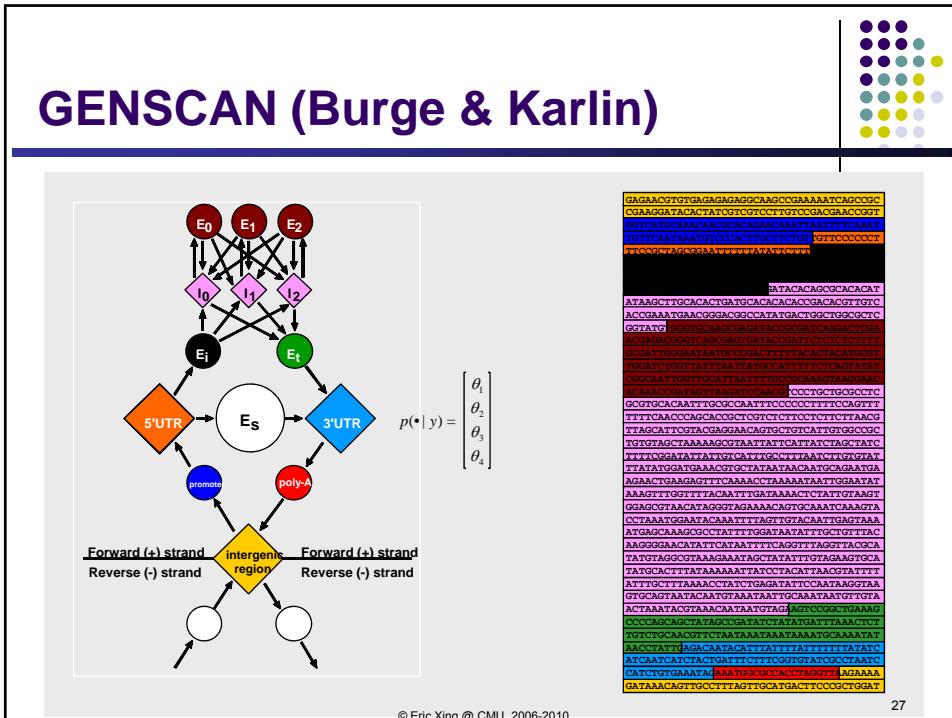
## Typical structure of a gene



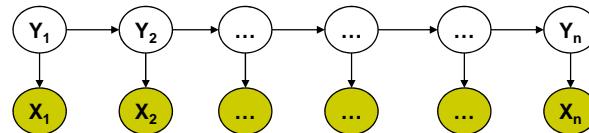
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## GENSCAN (Burge & Karlin)



## Shortcomings of Hidden Markov Model



- HMM models capture dependences between each state and **only its** corresponding observation
  - NLP example: In a sentence segmentation task, each segmental state may depend not just on a single word (and the adjacent segmental stages), but also on the (non-local) features of the whole line such as line length, indentation, amount of white space, etc.
- Mismatch between learning objective function and prediction objective function
  - HMM learns a joint distribution of states and observations  $P(Y, X)$ , but in a prediction task, we need the conditional probability  $P(Y|X)$

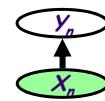
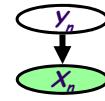
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## Recall Generative vs. Discriminative Classifiers



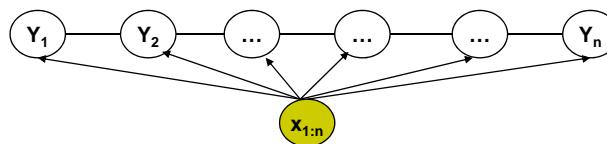
- Goal: Wish to learn  $f: X \rightarrow Y$ , e.g.,  $P(Y|X)$
- Generative classifiers (e.g., Naïve Bayes):
  - Assume some functional form for  $P(X|Y), P(Y)$   
This is a '**generative**' model of the data!
  - Estimate parameters of  $P(X|Y), P(Y)$  directly from training data
  - Use Bayes rule to calculate  $P(Y|X=x)$
- Discriminative classifiers (e.g., logistic regression)
  - Directly assume some functional form for  $P(Y|X)$   
This is a '**discriminative**' model of the data!
  - Estimate parameters of  $P(Y|X)$  directly from training data



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## Structured Conditional Models



- Conditional probability  $P(\text{label sequence } y \mid \text{observation sequence } x)$  rather than joint probability  $P(y, x)$ 
  - Specify the probability of possible label sequences given an observation sequence
- Allow arbitrary, non-independent features on the observation sequence  $X$
- The probability of a transition between labels may depend on **past** and **future** observations
- Relax strong independence assumptions in generative models

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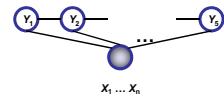
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## Conditional Distribution

- If the graph  $G = (V, E)$  of  $\mathbf{Y}$  is a tree, the conditional distribution over the label sequence  $\mathbf{Y} = \mathbf{y}$ , given  $\mathbf{X} = \mathbf{x}$ , by the Hammersley Clifford theorem of random fields is:

$$p_{\theta}(\mathbf{y} | \mathbf{x}) \propto \exp \left( \sum_{e \in E, k} \lambda_k f_k(e, \mathbf{y}|_e, \mathbf{x}) + \sum_{v \in V, k} \mu_k g_k(v, \mathbf{y}|_v, \mathbf{x}) \right)$$

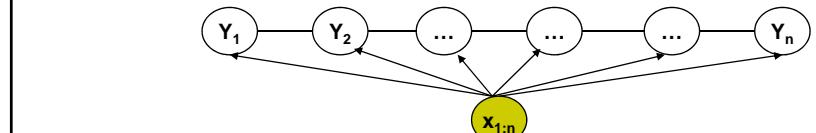
- $\mathbf{x}$  is a data sequence
- $\mathbf{y}$  is a label sequence
- $v$  is a vertex from vertex set  $V$  = set of label random variables
- $e$  is an edge from edge set  $E$  over  $V$
- $f_k$  and  $g_k$  are given and fixed.  $g_k$  is a Boolean vertex feature;  $f_k$  is a Boolean edge feature
- $k$  is the number of features
- $\theta = (\lambda_1, \lambda_2, \dots, \lambda_n; \mu_1, \mu_2, \dots, \mu_n); \lambda_k$  and  $\mu_k$  are parameters to be estimated
- $\mathbf{y}|_e$  is the set of components of  $\mathbf{y}$  defined by edge  $e$
- $\mathbf{y}|_v$  is the set of components of  $\mathbf{y}$  defined by vertex  $v$



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## Conditional Random Fields



$$P(\mathbf{y}_{1:n} | \mathbf{x}_{1:n}) = \frac{1}{Z(\mathbf{x}_{1:n})} \prod_{i=1}^n \phi(y_i, y_{i-1}, \mathbf{x}_{1:n}) = \frac{1}{Z(\mathbf{x}_{1:n}, \mathbf{w})} \prod_{i=1}^n \exp(\mathbf{w}^T \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_{1:n}))$$

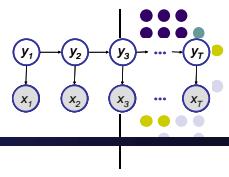
- CRF is a partially directed model

- Discriminative model
- Usage of global normalizer  $Z(\mathbf{x})$
- Models the dependence between each state and the entire observation sequence

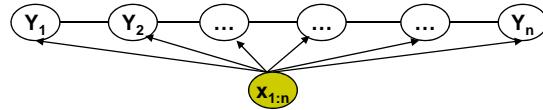
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## Conditional Random Fields



- General parametric form:



$$\begin{aligned} P(\mathbf{y}|\mathbf{x}) &= \frac{1}{Z(\mathbf{x}, \lambda, \mu)} \exp\left(\sum_{i=1}^n \left(\sum_k \lambda_k f_k(y_i, y_{i-1}, \mathbf{x}) + \sum_l \mu_l g_l(y_i, \mathbf{x})\right)\right) \\ &= \frac{1}{Z(\mathbf{x}, \lambda, \mu)} \exp\left(\sum_{i=1}^n (\lambda^T \mathbf{f}(y_i, y_{i-1}, \mathbf{x}) + \mu^T \mathbf{g}(y_i, \mathbf{x}))\right) \end{aligned}$$

$$\text{where } Z(\mathbf{x}, \lambda, \mu) = \sum_{\mathbf{y}} \exp\left(\sum_{i=1}^n (\lambda^T \mathbf{f}(y_i, y_{i-1}, \mathbf{x}) + \mu^T \mathbf{g}(y_i, \mathbf{x}))\right)$$

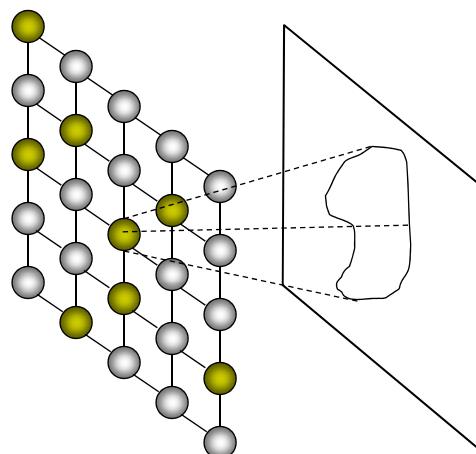
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## Conditional Random Fields



$$p_\theta(\mathbf{y}|\mathbf{x}) = \frac{1}{Z(\theta, \mathbf{x})} \exp\left\{\sum_c \theta_c f_c(\mathbf{x}, \mathbf{y}_c)\right\}$$



- Allow arbitrary dependencies on input
- Clique dependencies on labels
- Use approximate inference for general graphs

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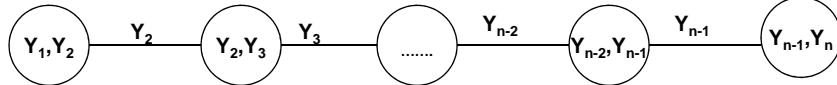
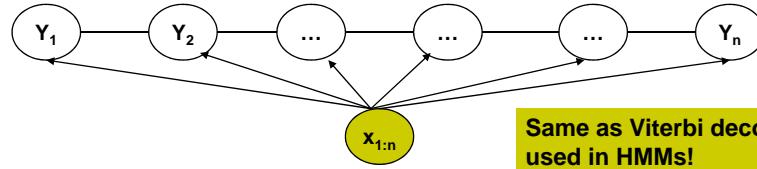
## CRFs: Inference

- Given CRF parameters  $\lambda$  and  $\mu$ , find the  $\mathbf{y}^*$  that maximizes  $P(\mathbf{y}|\mathbf{x})$

$$\mathbf{y}^* = \arg \max_{\mathbf{y}} \exp \left( \sum_{i=1}^n (\lambda^T \mathbf{f}(y_i, y_{i-1}, \mathbf{x}) + \mu^T \mathbf{g}(y_i, \mathbf{x})) \right)$$

• Can ignore  $Z(\mathbf{x})$  because it is not a function of  $\mathbf{y}$

- Run the max-product algorithm on the junction-tree of CRF:



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## CRF learning

- Given  $\{(\mathbf{x}_d, \mathbf{y}_d)\}_{d=1}^N$ , find  $\lambda^*, \mu^*$  such that

$$\begin{aligned} \lambda^*, \mu^* &= \arg \max_{\lambda, \mu} L(\lambda, \mu) = \arg \max_{\lambda, \mu} \prod_{d=1}^N P(\mathbf{y}_d | \mathbf{x}_d, \lambda, \mu) \\ &= \arg \max_{\lambda, \mu} \prod_{d=1}^N \frac{1}{Z(\mathbf{x}_d, \lambda, \mu)} \exp \left( \sum_{i=1}^n (\lambda^T \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_d) + \mu^T \mathbf{g}(y_{d,i}, \mathbf{x}_d)) \right) \\ &= \arg \max_{\lambda, \mu} \sum_{d=1}^N \left( \sum_{i=1}^n (\lambda^T \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_d) + \mu^T \mathbf{g}(y_{d,i}, \mathbf{x}_d)) - \log Z(\mathbf{x}_d, \lambda, \mu) \right) \end{aligned}$$

Gradient of the log-partition function in an exponential family is the expectation of the sufficient statistics.

- Computing the gradient w.r.t  $\lambda$ :

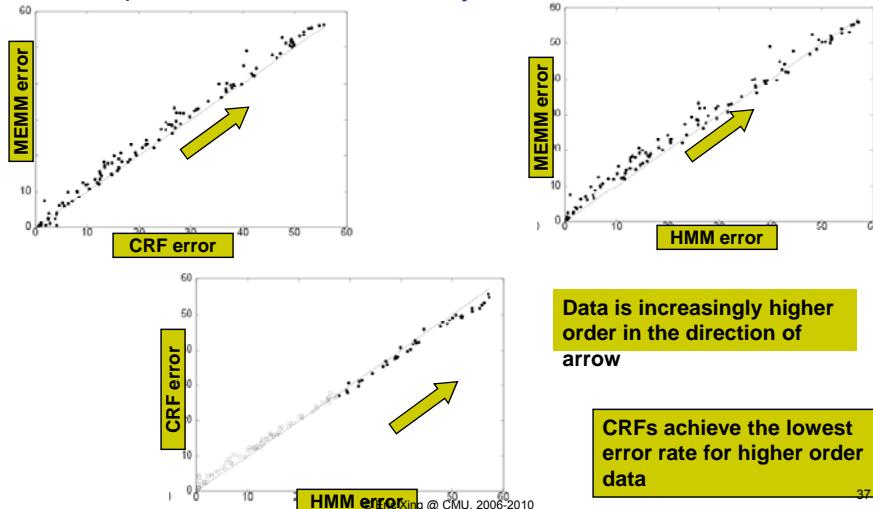
$$\nabla_\lambda L(\lambda, \mu) = \sum_{d=1}^N \left( \sum_{i=1}^n \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_d) - \sum_{\mathbf{y}} (P(\mathbf{y} | \mathbf{x}_d) \sum_{i=1}^n \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_d)) \right)$$

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## CRFs: some empirical results

- Comparison of error rates on synthetic data



## CRFs: some empirical results

- Parts of Speech tagging

model	error	oov error
HMM	5.69%	45.99%
MEMM	6.37%	54.61%
CRF	5.55%	48.05%
MEMM <sup>+</sup>	4.81%	26.99%
CRF <sup>+</sup>	4.27%	23.76%

+ Using spelling features

- Using same set of features: HMM >= CRF > MEMM
- Using additional overlapping features: CRF<sup>+</sup> > MEMM<sup>+</sup> >> HMM

## Summary

- Conditional Random Fields is a discriminative Structured Input Output model!
- HMM is a generative structured I/O model
- Complementary strength and weakness:

- 1.
- 2.
- 3.
- ...

?



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