Undirected Graphical Models for Sequence Analysis

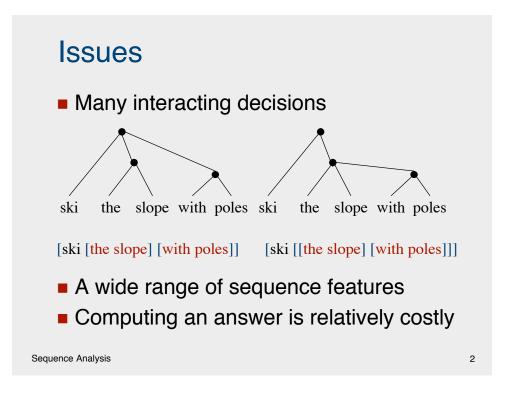
Fernando Pereira University of Pennsylvania

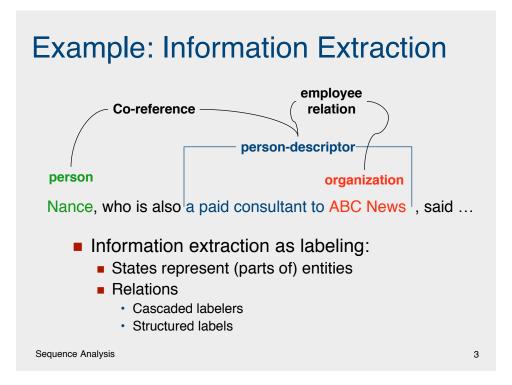
Joint work with John Lafferty, Andrew McCallum, and Fei Sha

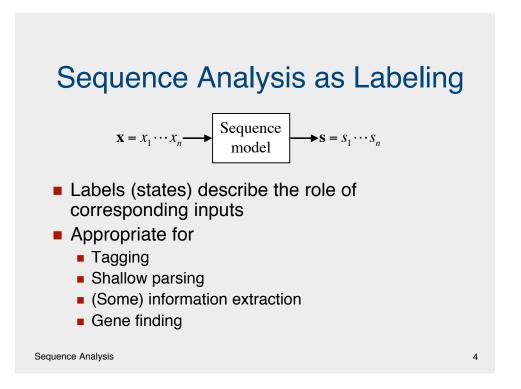


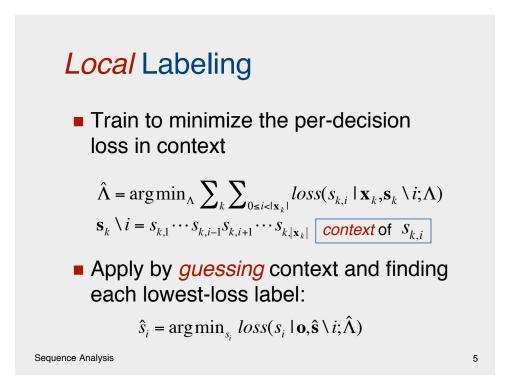
- Language
 - Syntactic structure
 - Sense tagging
 - Information extraction
- Biological sequences
 - Genes, regulatory regions
 - Secondary structure (folding)

Sequence Analysis









6

Global Labeling

Minimize training labeling loss

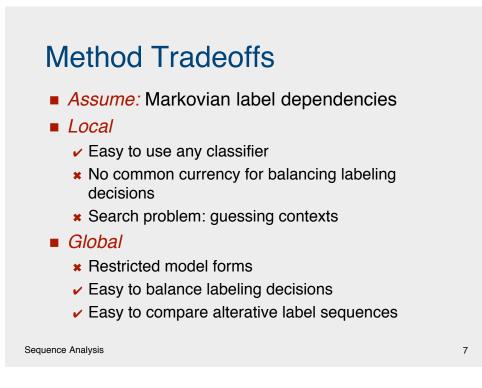
$$\hat{\Lambda} = \arg\min_{\Lambda} \sum_{k} Loss(\mathbf{x}_{k}, \mathbf{s}_{k} \mid \Lambda)$$

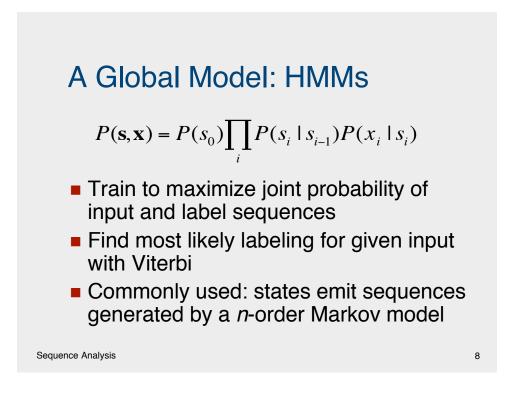
Computing the best labeling:

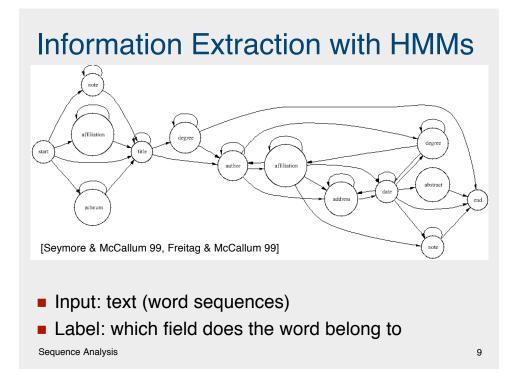
$$\hat{\mathbf{s}} = \arg\min_{\mathbf{s}} Loss(\mathbf{x}, \mathbf{s} \mid \hat{\Lambda})$$

- Efficient minimization requires:
 - A common currency for local labeling decisions
 - Efficient algorithm to combine the decisions

Sequence Analysis







10

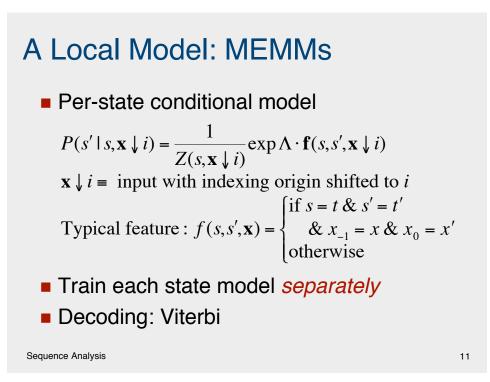
Problems with HMMs

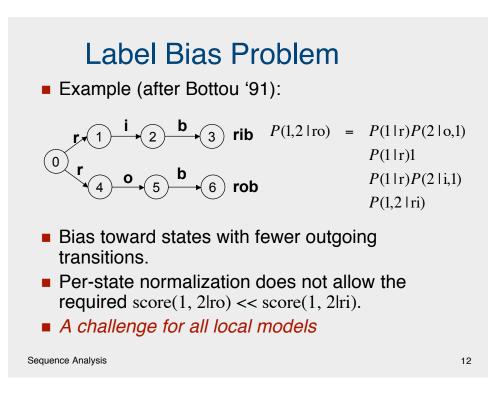
 Applications need richer input representation: multiple overlapping features, whole chunks of text

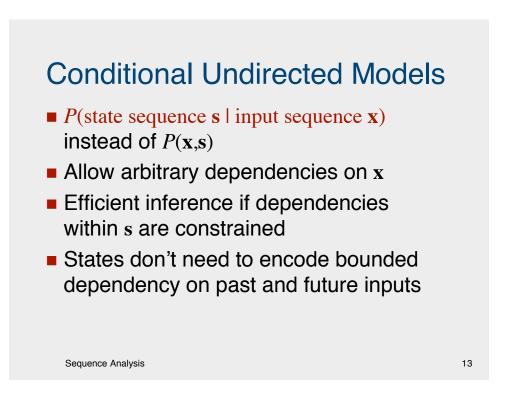
Word features	Context features	
word identity	previous words	
capitalization	next words	
ends in "-tion"	markup	
word in word list	starts sentence	

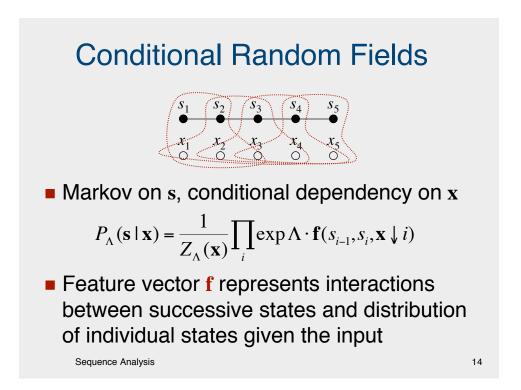
- Generative models do not handle easily overlapping, non-independent features
- Alternative: *conditional* model $P(\mathbf{s}|\mathbf{x})$

Sequence Analysis

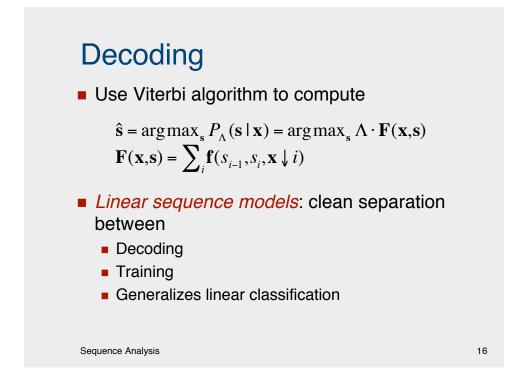








From HMMs to CRFs $s = s_{1} \cdots s_{n} \qquad x = x_{1} \cdots x_{n}$ HMM $P(s \mid x) = \frac{P(s_{0})}{P(x)} \prod_{i} P(s_{i} \mid s_{i-1}) P(x_{i} \mid s_{i})$ CRF $P(s \mid x) = \frac{1}{Z(x)} \prod_{i} \exp \begin{pmatrix} \Lambda \cdot \mathbf{f}(s_{i}, s_{i-1}) \\ + \\ \Omega \cdot \mathbf{g}(s_{i}, x_{i}) \end{pmatrix}$



Efficient Estimation

Matrix notation

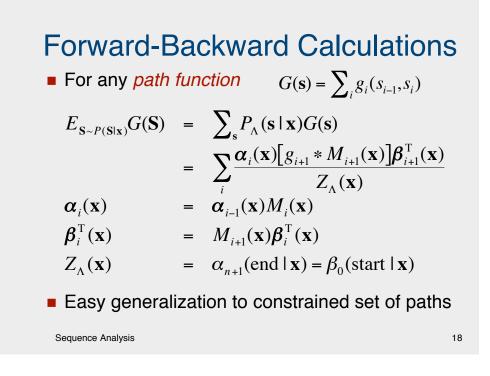
$$M_{i}(s,s' | \mathbf{x}) = \exp \Lambda \cdot \mathbf{f}(s,s',\mathbf{x} \downarrow i)$$

$$P_{\Lambda}(\mathbf{s} | \mathbf{x}) = \frac{1}{Z_{\Lambda}(\mathbf{o})} \prod_{i} M_{i}(s_{i-1},s_{i} | \mathbf{x})$$

$$Z_{\Lambda}(\mathbf{x}) = (M_{1}(\mathbf{x})M_{2}(\mathbf{x})\cdots M_{n+1}(\mathbf{x}))_{\text{start,stop}}$$

Efficient normalization: *forward-backward* algorithm

Sequence Analysis



Training

- Maximize $L(\Lambda) = \sum_{k} \log P_{\Lambda}(\mathbf{s}_{k} | \mathbf{x}_{k})$ Log-likelihood *gradient*

$$\nabla L(\Lambda) = \sum_{k} \left(\mathbf{F}(\mathbf{x}_{k}, \mathbf{s}_{k}) - E_{\mathbf{S} \sim P_{\Lambda}(\mathbf{S}|\mathbf{x}_{k})} \mathbf{F}(\mathbf{x}_{k}, \mathbf{S}) \right)$$

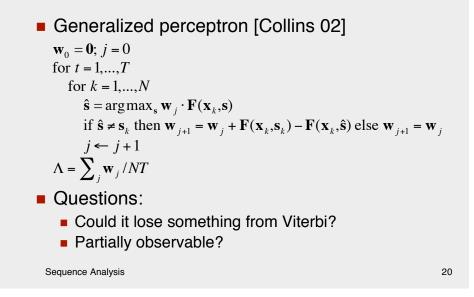
 Methods: iterative scaling, conjugate gradient, L-BFGS

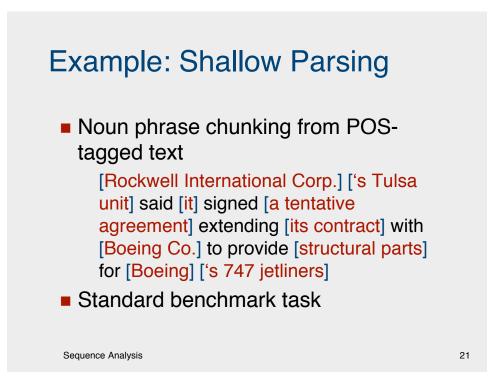
Partially-observable case (labeled states)

$$P_{\Lambda}(\mathbf{y} \mid \mathbf{x}) = \sum_{\mathbf{s}:\ell(\mathbf{s})=\mathbf{y}} P_{\Lambda}(\mathbf{s} \mid \mathbf{x})$$

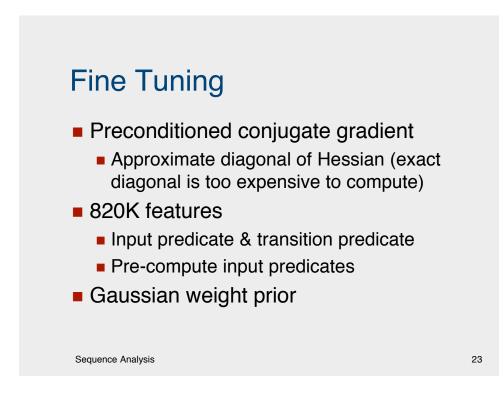
$$\nabla L(\Lambda) = \sum_{k} \begin{pmatrix} E_{\mathbf{S}\sim P_{\Lambda}(\mathbf{S}\mid\mathbf{x}_{k},\ell(\mathbf{S})=\mathbf{y}_{k}) \mathbf{F}(\mathbf{x}_{k},\mathbf{S}) \\ -E_{\mathbf{S}\sim P_{\Lambda}(\mathbf{S}\mid\mathbf{x}_{k})} \mathbf{F}(\mathbf{x}_{k},\mathbf{S}) \end{pmatrix}$$
Sequence Analysis

Alternative Training Method





NP Chunking Results		
Model	F	
24 SVM combination [Kudo & Matsumoto 01]	94.22%	
CRF [Sha & Pereira 03]	94.20%	
Voted perceptron [Collins 02; Sha & Pereira 03]	94.09%	$F = \frac{2PR}{P+R}$
<i>Winnow</i> [Zhang, Damerau & Johnson 02]	93.89%	
MEMM [Sha & Pereira 03]	93.70%	
 Warning: different feature sets 		
Sequence Analysis		22



Further Questions

- Limited by dimensionality (number of features): kernels?
- Generalization bounds
- Parsing
 - Trees instead of chains
 - Inside-outside replaces forward-backward
 - Computationally challenging: large label set
- General graphs using loopy BP
 - Suggestive results for *collective classification* [Taskar & al 02]

Sequence Analysis