



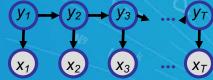
Probabilistic Graphical Models

01010001 Ω

Case Studies: HMM and CRF

Eric Xing Lecture 6, February 3, 2020

Reading: see class homepage

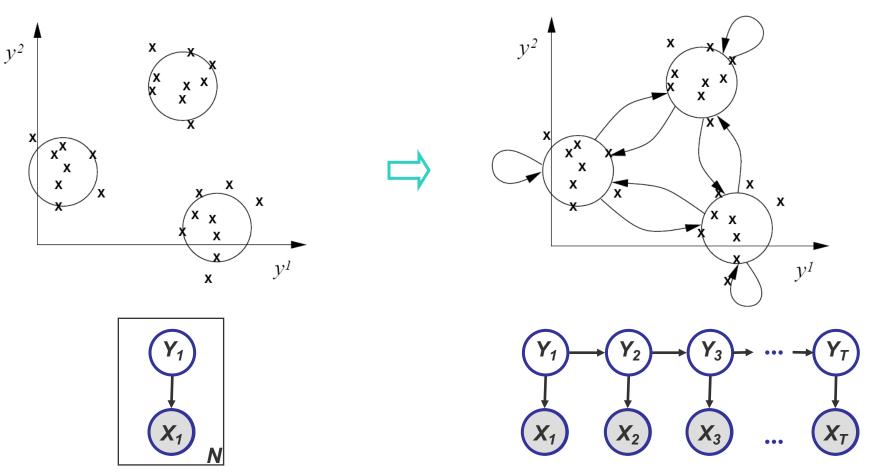


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Hidden Markov Model: from static to dynamic mixture models

Static mixture

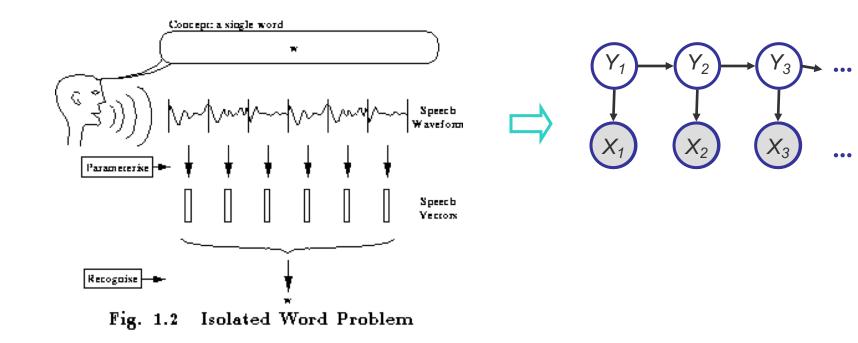
Dynamic mixture







Speech recognition





3

 X_T

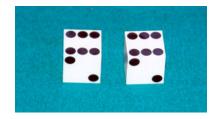


Some early applications of HMMs

- 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







- Observation space
 Alphabetic set:
 Euclidean space:
- □ Index set of hidden states $I = \{1, 2, \dots, M\}$
- Transition probabilities between any two states

$$p(y_{t}^{j} = 1 | y_{t-1}^{i} = 1) = a_{i,j},$$

$$p(y_{t} | y_{t-1}^{i} = 1) \sim \text{Multinomial}(a_{i,1}, a_{i,1}, \dots, a_{i,M}), \forall i \in \mathbb{I}.$$

 $\mathbf{C} = \left\{ \boldsymbol{c}_1, \boldsymbol{c}_2, \cdots, \boldsymbol{c}_{\mathcal{K}} \right\}$

• Start probabilities

or

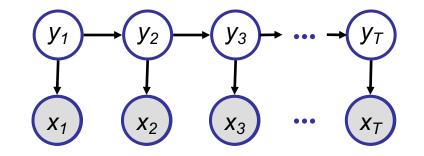
 $p(\gamma_1) \sim \text{Multinomial}(\pi_1, \pi_2, \dots, \pi_M).$

Emission probabilities associated with each state

$$p(\mathbf{x}_t | \mathbf{y}_t^i = \mathbf{1}) \sim \text{Multinomial}(\mathbf{b}_{i,1}, \mathbf{b}_{i,1}, \dots, \mathbf{b}_{i,k}), \forall i \in \mathbf{I}.$$

or in general:

 $p(\mathbf{x}_t | \mathbf{y}_t^i = 1) \sim f(\cdot | \theta_i), \forall i \in I.$

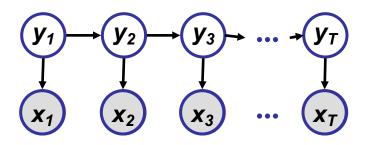






- Given a sequence $\mathbf{x} = \mathbf{x}_1, \dots, \mathbf{x}_T$ and a parse $\mathbf{y} = \mathbf{y}_1, \dots, \mathbf{y}_T$,
- To find how likely is the parse:
 (given our HMM and the sequence)

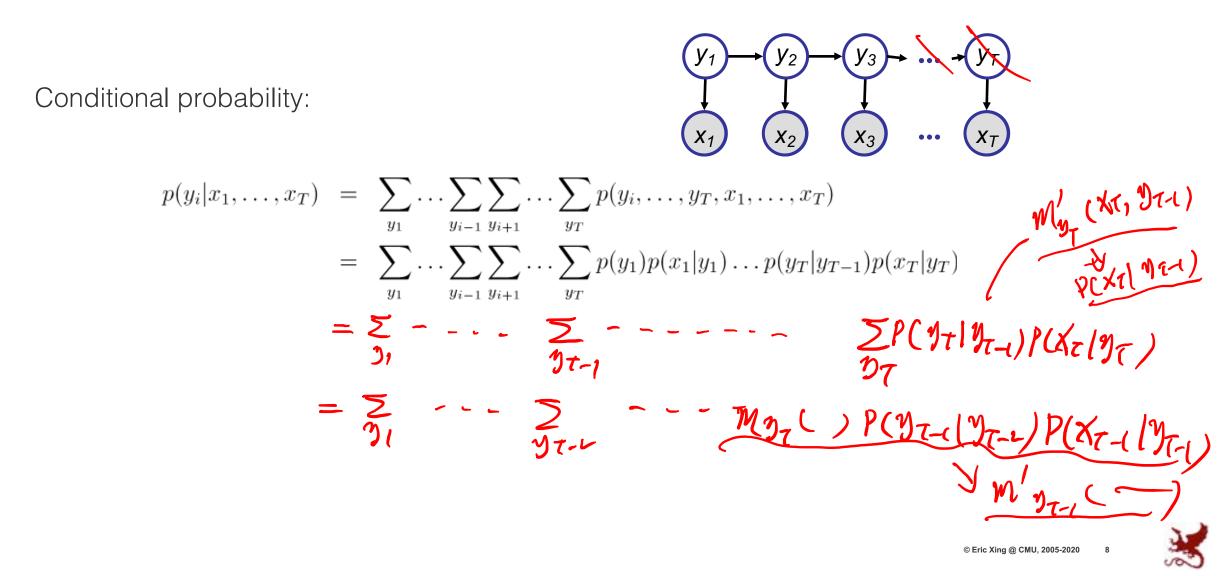
 $p(\mathbf{x}, \mathbf{y}) = p(x_1, \dots, x_T, y_1, \dots, y_T)$ (Joint probability) = $p(y_1) p(x_1 | y_1) p(y_2 | y_1) p(x_2 | y_2) \dots p(y_T | y_{T-1}) p(x_T | y_T)$ = $p(y_1) P(y_2 | y_1) \dots p(y_T | y_{T-1}) \times p(x_1 | y_1) p(x_2 | y_2) \dots p(x_T | y_T)$ = $p(y_1, \dots, y_T) p(x_1, \dots, x_T | y_1, \dots, y_T)$



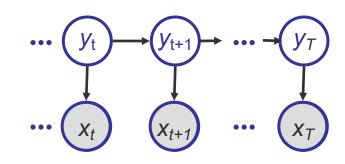


Variable Elimination on Hidden Markov Model **y**3 **X**3 XT X_2 $p(\mathbf{x}, \mathbf{y}) = p(x_1, \dots, x_T, y_1, \dots, y_T)$ $= p(y_1) p(x_1 | y_1) p(y_2 | y_1) p(x_2 | y_2) \dots p(y_T | y_{T-1}) p(x_T | y_T)$ My, (x, N)) = P(N2 x) Conditional probability: $p(y_i|x_1,\ldots,x_T) \,\, \bigotimes \,\, \sum \ldots \sum p(y_i,\ldots,y_T,x_1,\ldots,x_T)$ $y_{i-1}y_{i+1}$ $\sum \dots \sum p(y_1)p(x_1|y_1)\dots p(y_T|y_{T-1})p(x_T|y_T)$ $y_{i-1} \ y_{i+1}$ y_1 $\geq P(\mathcal{Y}_1) P(X_1|\mathcal{Y}_1) P(\mathcal{Y}_2|\mathcal{Y}_1)$ $\sum_{m} M(K_{1}y_{1}) = M(K_{1}y_{1}) P(y_{1}y_{2}) = Fric Xing @ CMU, 2005-2020 7$

Variable Elimination on Hidden Markov Model







- We want to calculate P(x), the likelihood of x, given the HMM
 - Sum over all possible ways of generating **x**:

$$p(\mathbf{x}) = \sum_{\mathbf{y}} p(\mathbf{x}, \mathbf{y}) = \sum_{y_1} \sum_{y_2} \cdots \sum_{y_N} \pi_{y_1} \prod_{t=2}^{T} a_{y_{t-1}, y_t} \prod_{t=1}^{T} p(x_t \mid y_t)$$

D To avoid summing over an exponential number of paths **y**, define

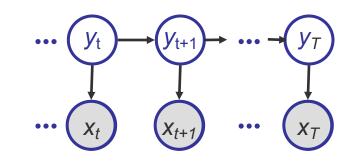
$$\alpha(y_t^k = \mathbf{1}) = \alpha_t^k \stackrel{\text{def}}{=} P(x_1, \dots, x_t, y_t^k = \mathbf{1}) \quad \text{(the forward probability)}$$

• The recursion:

$$\alpha_t^k = p(x_t \mid y_t^k = \mathbf{1}) \sum_i \alpha_{t-1}^i a_{i,k}$$
$$P(\mathbf{x}) = \sum_k \alpha_T^k$$



The Backward Algorithm



• We want to compute $P(y_t^k = \mathbf{1} | \mathbf{x})$,

the posterior probability distribution on the \mathbf{T}^{th} position, given \mathbf{x}

We start by computing

$$P(\mathbf{y}_{t}^{k} = \mathbf{1}, \mathbf{x}) = P(\mathbf{x}_{1}, ..., \mathbf{x}_{t}, \mathbf{y}_{t}^{k} = \mathbf{1}, \mathbf{x}_{t+1}, ..., \mathbf{x}_{T})$$

= $P(x_{1}, ..., x_{t}, y_{t}^{k} = \mathbf{1})P(x_{t+1}, ..., x_{T} | x_{1}, ..., x_{t}, y_{t}^{k} = \mathbf{1})$
= $P(x_{1}...x_{t}, y_{t}^{k} = \mathbf{1})P(x_{t+1}...x_{T} | y_{t}^{k} = \mathbf{1})$

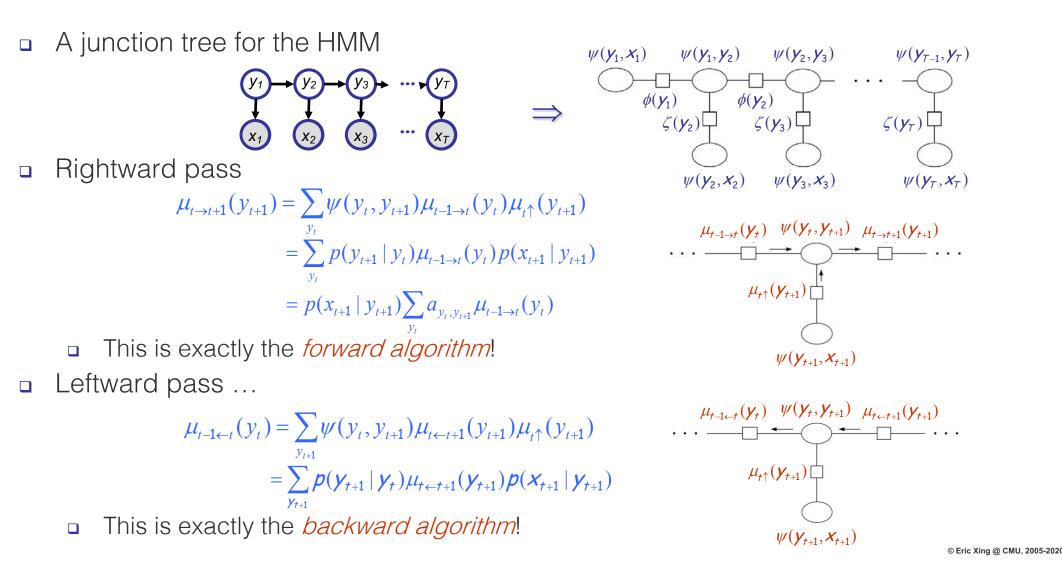
Forward, α_t^{k}

Backward, $\beta_t^k = P(x_{t+1}, ..., x_T | y_t^k = 1)$

The recursion:



The junction tree algorithm: message passing for HMM





• Forward algorithm

 $\alpha_t^k \stackrel{\text{def}}{=} \mu_{t-1 \to t}(k) = P(x_1, \dots, x_{t-1}, x_t, y_t^k = 1)$ $\alpha_t^k = p(x_t \mid y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k}$ • Backward algorithm

 $\beta_{t}^{k} = \sum_{i} a_{k,i} p(\mathbf{x}_{t+1} \mid y_{t+1}^{i} = 1) \beta_{t+1}^{i}$ $\beta_{t}^{k} \stackrel{\text{def}}{=} \mu_{t-1 \leftarrow t}(\mathbf{k}) = P(\mathbf{x}_{t+1}, ..., \mathbf{x}_{T} \mid \mathbf{y}_{t}^{k} = 1)$

$$\gamma_{t}^{i} \stackrel{\text{def}}{=} p(y_{t}^{i} = 1 \mid x_{1:T}) \propto \alpha_{t}^{i} \beta_{t}^{i} = \sum_{j} \xi_{t}^{i,j}$$

$$\xi_{t}^{i,j} \stackrel{\text{def}}{=} p(y_{t}^{i} = 1, y_{t+1}^{j} = 1, x_{1:T})$$

$$\propto \mu_{t-1 \to t} (y_{t}^{i} = 1) \mu_{t \leftarrow t+1} (y_{t+1}^{j} = 1) p(x_{t+1} \mid y_{t+1}) p(y_{t+1} \mid y_{t})$$

$$\xi_{t}^{i,j} = \alpha_{t}^{i} \beta_{t+1}^{j} a_{i,j} p(x_{t+1} \mid y_{t+1}^{i} = 1)$$

The matrix-vector form: $B_{t}(i) \stackrel{\text{def}}{=} p(x_{t} | y_{t}^{i} = 1)$ $A(i, j) \stackrel{\text{def}}{=} p(y_{t+1}^{j} = 1 | y_{t}^{i} = 1)$ $\alpha_{t} = (A^{T} \alpha_{t-1}) \cdot B_{t}$ $\beta_{t} = A(\beta_{t+1} \cdot B_{t+1})$ $\xi_{t} = (\alpha_{t} (\beta_{t+1} \cdot B_{t+1})^{T}) \cdot A$ $\gamma_{t} = \alpha_{t} \cdot B_{t}$





• 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 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?
- $\left\{ \boldsymbol{y}_{t}^{\boldsymbol{k}_{t}^{*}} = \boldsymbol{1}: \boldsymbol{t} = \boldsymbol{1} \cdots \boldsymbol{\mathcal{T}} \right\}$ Posterior Decoding:
- This is different from MPA of a whole sequence hidden states **Example:**
- This can be understood as *bit error rate* vs. word error rate

P(x,y)X Y 0.35 0 0 0 1 0.05 1 0 0.3 MPA of (X, Y)? 0.3 1

MPA of X?



Viterbi decoding

• GIVEN $\mathbf{x} = \mathbf{x}_1, \dots, \mathbf{x}_T$, we want to find $\mathbf{y} = \mathbf{y}_1, \dots, \mathbf{y}_T$, such that $\mathbf{P}(\mathbf{y}|\mathbf{x})$ is maximized:

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

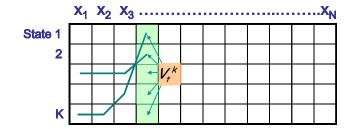
• Let
$$V_t^k = \max_{\{y_1, \dots, y_{t-1}\}} P(x_1, \dots, x_{t-1}, y_1, \dots, y_{t-1}, x_t, y_t^k = 1)$$

= Probability of most likely <u>sequence of states</u> ending at state $y_t = k$

- The recursion: $V_{t}^{k} = p(x_{t} | y_{t}^{k} = 1) \max_{i,k} V_{t-1}^{i}$
- Underflows are a significant problem

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

- These numbers become extremely small underflow
- Solution: Take the logs of all values:



 $V_{t}^{k} = \log p(x_{t} | y_{t}^{k} = 1) + \max_{i} (\log(a_{i,k}) + V_{t-1}^{i})$



The Viterbi Algorithm – derivation

Define the viterbi probability:

$$\begin{aligned} \mathcal{V}_{t+1}^{k} &= \max_{\{y_{1},\dots,y_{t}\}} \mathcal{P}(x_{1},\dots,x_{t},y_{1},\dots,y_{t},x_{t+1},y_{t+1}^{k}=1) \\ &= \max_{\{y_{1},\dots,y_{t}\}} \mathcal{P}(x_{t+1},y_{t+1}^{k}=1 \mid x_{1},\dots,x_{t},y_{1},\dots,y_{t}) \mathcal{P}(x_{1},\dots,x_{t},y_{1},\dots,y_{t}) \\ &= \max_{\{y_{1},\dots,y_{t}\}} \mathcal{P}(x_{t+1},y_{t+1}^{k}=1 \mid y_{t}) \mathcal{P}(x_{1},\dots,x_{t-1},y_{1},\dots,y_{t-1},x_{t},y_{t}) \\ &= \max_{i} \mathcal{P}(x_{t+1},y_{t+1}^{k}=1 \mid y_{t}^{i}=1) \max_{\{y_{1},\dots,y_{t-1}\}} \mathcal{P}(x_{1},\dots,x_{t-1},y_{1},\dots,y_{t-1},x_{t},y_{t}^{i}=1) \\ &= \max_{i} \mathcal{P}(x_{t+1},\mid y_{t+1}^{k}=1) a_{i,k} V_{t}^{i} \\ &= \mathcal{P}(x_{t+1},\mid y_{t+1}^{k}=1) \max_{i,k} a_{i,k} V_{t}^{i} \end{aligned}$$



Computational Complexity and implementation details

• What is the running time, and space required, for Forward, and Backward? $\alpha_{i}^{k} = p(x_{i} \mid y_{i}^{k} = 1) \sum \alpha_{i}^{i} \cdot a_{i}$

$$\alpha_{t} = p(\mathbf{x}_{t} | \mathbf{y}_{t} = \mathbf{I}) \sum_{i} \alpha_{t-1} a_{i,k}$$

$$\beta_{t}^{k} = \sum_{i} a_{k,i} p(x_{t+1} | y_{t+1}^{i} = \mathbf{I}) \beta_{t+1}^{i}$$

$$V_{t}^{k} = p(\mathbf{x}_{t} | \mathbf{y}_{t}^{k} = \mathbf{I}) \max_{i} a_{i,k} V_{t-1}^{i}$$

Time: **O(K² N)**; Space: O(**KN)**.

- Useful implementation technique to avoid underflows
 - Viterbi: sum of logs
 - □ Forward/Backward: rescaling at each position by multiplying by a constant



Learning HMM: two scenarios

- Supervised learning: estimation when the "right answer" is known
 - Examples:

GIVEN: a genomic region $x = x_1...x_{1,000,000}$ 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

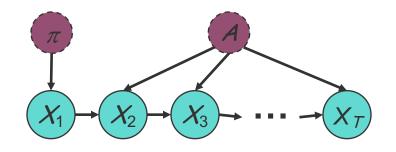
• Examples:

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 θ of the model to maximize $P(x|\theta)$ ----Maximal likelihood (ML) estimation







- Consider a time-invariant (stationary) 1st-order Markov model
 - Initial state probability vector:
 - State transition probability matrix:
- The joint:

$$p(X_{1:T} \mid \theta) = p(x_1 \mid \pi) \prod_{t=2}^{T} \prod_{t=2} p(X_t \mid X_{t-1})$$

- The log-likelihood:
 - $\ell(\theta; D) = \sum \log p(x_{n,1} | \pi) + \sum_{n} \sum_{t=2}^{T} \log p(x_{n,t} | x_{n,t-1}, A)$ Again, we optimize each parameter separately

 $A_{ii} \stackrel{\text{def}}{=} p(X_t^j = \mathbf{1} | X_{t-1}^i = \mathbf{1})$

 $\pi_k = p(X_1^k = \mathbf{1})$

- - π is a multinomial frequency vector, and we've seen it before
 - What about A?

Learning a Markov chain transition matrix

• *A* is a stochastic matrix:

$$\sum\nolimits_{j} A_{ij} = \mathbf{1}$$

- Each row of A is multinomial distribution.
- So MLE of A_{ij} is the fraction of transitions from *i* to *j*

$$A_{ij}^{ML} = \frac{\#(i \to j)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=2}^{T} x_{n,t-1}^{i} x_{n,t}^{j}}{\sum_{n} \sum_{t=2}^{T} x_{n,t-1}^{i}}$$

- Application:
 - \Box if the states X_{t} represent words, this is called a *bigram language model*
- Sparse data problem:
 - □ If $i \rightarrow j$ did not occur in data, we will have $A_{ij} = 0$, then any future sequence with word pair $i \rightarrow j$ will have zero probability.
 - A standard hack: *backoff smoothing* or *deleted interpolation*

$$\widetilde{\mathcal{A}}_{i\to\bullet} = \lambda \eta_t + (1-\lambda) \mathcal{A}_{i\to\bullet}^{ML}$$

Supervised ML estimation for "Hidden" MM

- Given x = x₁...x_N for which the true state path y = y₁...y_N is known,
 Define:
 - $A_{ij} = \#$ times state transition $i \rightarrow j$ occurs in y
 - B_{ik} = # times state *i* in y emits *k* in x
 - We can show that the maximum likelihood parameters θ are:

$$a_{ij}^{ML} = \frac{\#(i \to j)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=2}^{T} y_{n,t-1}^{i} y_{n,t}^{j}}{\sum_{n} \sum_{t=2}^{T} y_{n,t-1}^{i}} = \frac{A_{ij}}{\sum_{j'} A_{ij'}}$$
$$b_{ik}^{ML} = \frac{\#(i \to k)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=1}^{T} y_{n,t}^{i} x_{n,t}^{k}}{\sum_{n} \sum_{t=1}^{T} y_{n,t}^{i}} = \frac{B_{ik}}{\sum_{k'} B_{ik'}}$$

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



Supervised ML estimation, ctd.

Intuition:

• When we know the underlying states, the best estimate of θ 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 θ is unreasonable: 0 probabilities VERY BAD

Example:

Given 10 casino rolls, we observe

• Then:

$$\mathbf{x} = 2, 1, 5, 6, 1, 2, 3, 6, 2, 3$$

$$\mathbf{y} = \mathbf{F}, \mathbf{F}$$



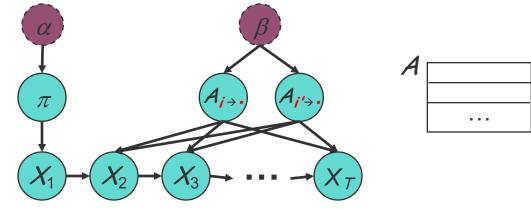


- Solution for small training sets:
 - Add pseudocounts
 - $A_{ij} = \#$ times state transition $i \rightarrow j$ occurs in $\mathbf{y} + R_{ij}$
 - B_{ik} = # times state *i* in y emits *k* in x + S_{ik}
 - \square R_{ij} , S_{ij} are pseudocounts representing our prior belief
 - **D** Total pseudocounts: $\mathbf{R}_i = \Sigma_j \mathbf{R}_{ij}$, $\mathbf{S}_i = \Sigma_k \mathbf{S}_{ik}$,
 - --- "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
- This is equivalent to Bayesian est. under a uniform prior with "parameter strength" equals to the pseudocounts





Global and local parameter independence



- The posterior of $A_{i \rightarrow}$ and $A_{i' \rightarrow}$ is factorized despite v-structure on X_{t} , because X_{t-1} acts like a multiplexer
- Assign a Dirichlet prior β_i to each row of the transition matrix:

$$A_{ij}^{Bayes} \stackrel{\text{def}}{=} p(j \mid i, D, \beta_i) = \frac{\#(i \to j) + \beta_{i,k}}{\#(i \to \bullet) + |\beta_i|} = \lambda_i \beta_{i,k}' + (\mathbf{1} - \lambda_i) A_{ij}^{ML}, \text{ where } \lambda_i = \frac{|\beta_i|}{|\beta_i| + \#(i \to \bullet)}$$

 We could consider more realistic priors, e.g., mixtures of Dirichlets to account for types of words (adjectives, verbs, etc.)





• Supervised learning: estimation when the "right answer" is known

• Examples:

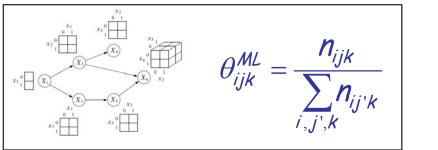
GIVEN:a genomic region $x = x_1 \dots x_{1,000,000}$ where we have good (experimental)annotations of the CpG islandsGIVEN:GIVEN:the casino player allows us to observe him one evening,as he changes diceand produces 10,000 rolls

- Unsupervised learning: estimation when the "right answer" is unknown
 - <u>Examples:</u> 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 θ of the model to maximize $P(x|\theta)$ ----Maximal likelihood (ML) estimation





- Supervised learning: if only we knew the true state path then ML parameter estimation would be trivial
 - E.g., recall that for complete observed tabular BN:



$$a_{ij}^{ML} = \frac{\#(i \to j)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=2}^{T} Y_{n,t-1}^{i} Y_{n,t}^{j}}{\sum_{n} \sum_{t=2}^{T} Y_{n,t-1}^{i}}$$
$$b_{ik}^{ML} = \frac{\#(i \to k)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=1}^{T} Y_{n,t}^{i} X_{n,t}^{k}}{\sum_{n} \sum_{t=1}^{T} Y_{n,t}^{i}}$$

• What if y is continuous? We can treat $\{(x_{n,t}, y_{n,t}): t = 1:T, n = 1:N\}$ as N' T observations of, e.g., a GLIM, and apply learning rules for GLIM ...

- Unsupervised learning: when the true state path is unknown, we can fill in the missing values using inference recursions.
 - □ The Baum Welch algorithm (i.e., EM)
 - Guaranteed to increase the log likelihood of the model after each iteration
 - Converges to local optimum, depending on initial conditions

The Baum Welch algorithm

- The complete log likelihood $\ell_{c}(\boldsymbol{\theta}; \mathbf{x}, \mathbf{y}) = \log p(\mathbf{x}, \mathbf{y}) = \log \prod_{n} \left(p(\boldsymbol{y}_{n,1}) \prod_{t=2}^{T} p(\boldsymbol{y}_{n,t} \mid \boldsymbol{y}_{n,t-1}) \prod_{t=1}^{T} p(\boldsymbol{x}_{n,t} \mid \boldsymbol{x}_{n,t}) \right)$
- $\Box \text{ The expected complete log likelihood} \\ \langle \ell_c(\mathbf{\theta}; \mathbf{x}, \mathbf{y}) \rangle = \sum_n \left(\left\langle \mathbf{y}_{n,1}^i \right\rangle_{p(y_{n,1}|\mathbf{x}_n)} \log \pi_i \right) + \sum_n \sum_{t=2}^T \left(\left\langle \mathbf{y}_{n,t-1}^i \mathbf{y}_{n,t}^j \right\rangle_{p(y_{n,t-1},y_{n,t}|\mathbf{x}_n)} \log \mathbf{a}_{i,j} \right) + \sum_n \sum_{t=1}^T \left(\mathbf{x}_{n,t}^k \left\langle \mathbf{y}_{n,t}^i \right\rangle_{p(y_{n,t}|\mathbf{x}_n)} \log \mathbf{b}_{i,k} \right)$

The E step

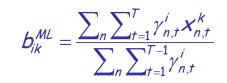
$$\gamma_{n,t}^{i} = \left\langle \mathbf{y}_{n,t}^{i} \right\rangle = \mathbf{p}(\mathbf{y}_{n,t}^{i} = \mathbf{1} | \mathbf{x}_{n})$$

$$\xi_{n,t}^{i,j} = \left\langle \mathbf{y}_{n,t-1}^{i} \mathbf{y}_{n,t}^{j} \right\rangle = \mathbf{p}(\mathbf{y}_{n,t-1}^{i} = \mathbf{1}, \mathbf{y}_{n,t}^{j} = \mathbf{1} | \mathbf{x}_{n})$$

 $a_{ij}^{ML} = \frac{\#(i \to j)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=2}^{T} Y_{n,t-1}^{i} Y_{n,t}^{j}}{\sum_{n} \sum_{t=2}^{T} Y_{n,t-1}^{i}}$ $b_{ik}^{ML} = \frac{\#(i \to k)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=1}^{T} Y_{n,t}^{i} X_{n,t}^{k}}{\sum_{n} \sum_{t=1}^{T} Y_{n,t}^{i}}$

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

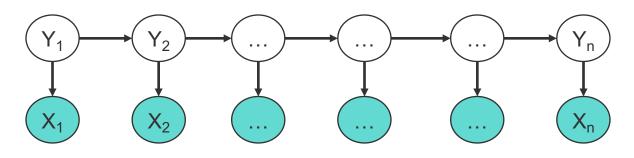
$$\pi_{i}^{ML} = \frac{\sum_{n} \gamma_{n,1}^{i}}{N} \qquad a_{ij}^{ML} = \frac{\sum_{n} \sum_{t=2}^{T} \xi_{n,t}^{i,j}}{\sum_{n} \sum_{t=1}^{T-1} \gamma_{n,t}^{i}}$$



Conditional Random Fields



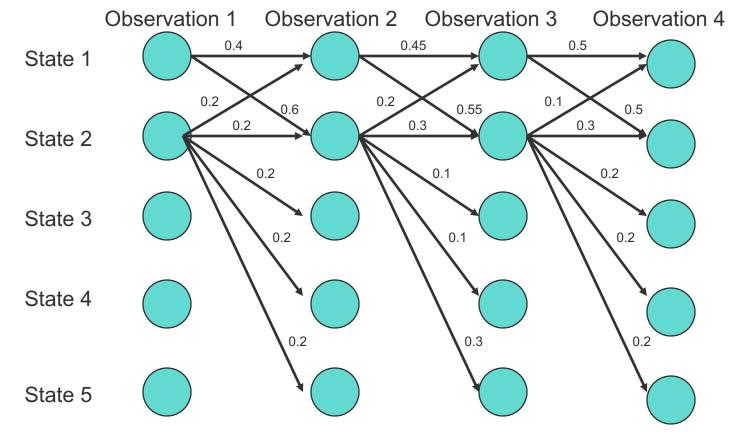
Shortcomings of Hidden Markov Model (1): locality of features



- 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)



Shortcomings of HMM (2): the Label bias problem

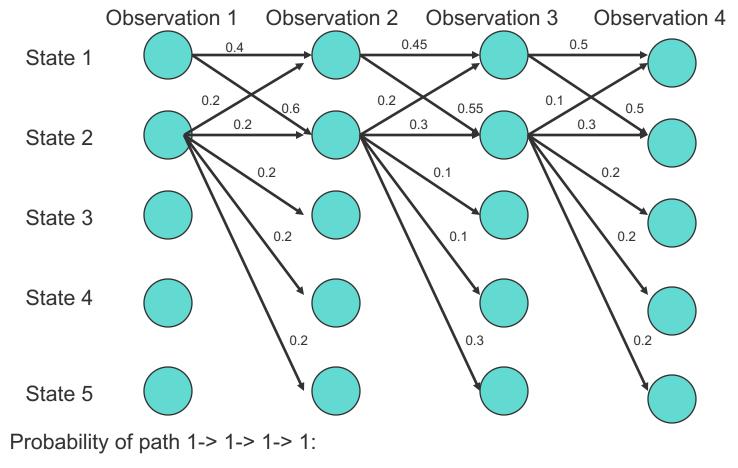


What the local transition probabilities say:

- State 1 almost always prefers to go to state 2
- State 2 almost always prefer to stay in state 2



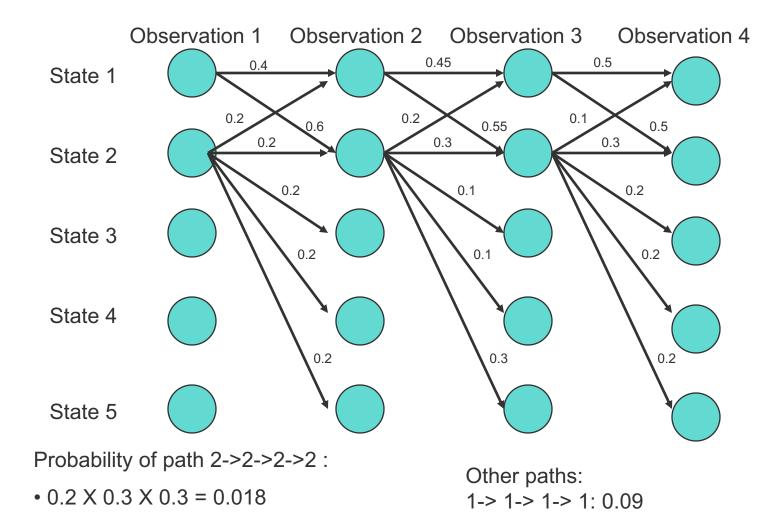




• $0.4 \times 0.45 \times 0.5 = 0.09$

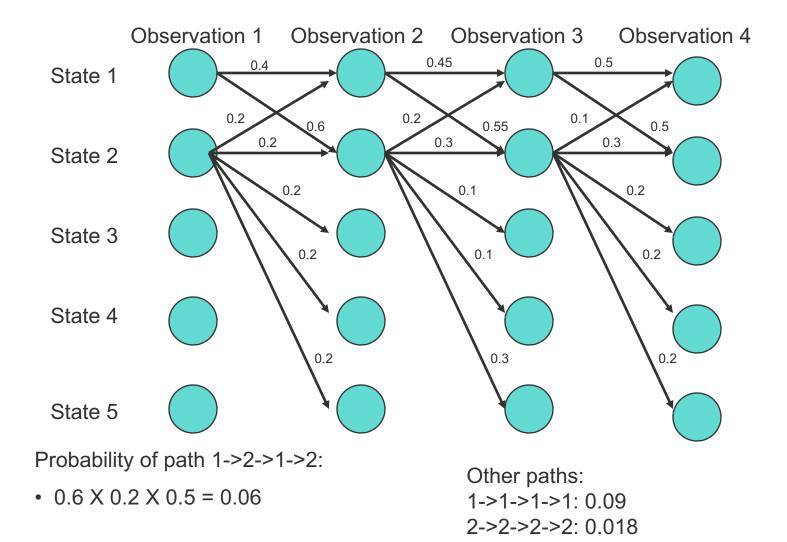






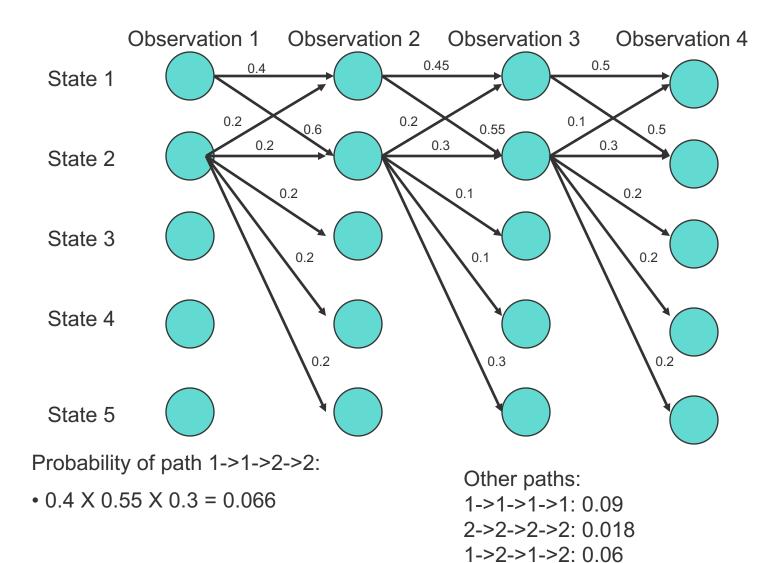








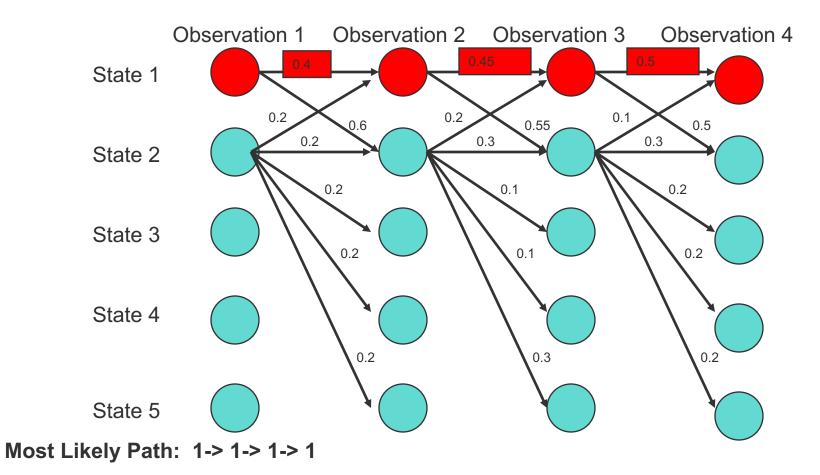




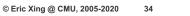
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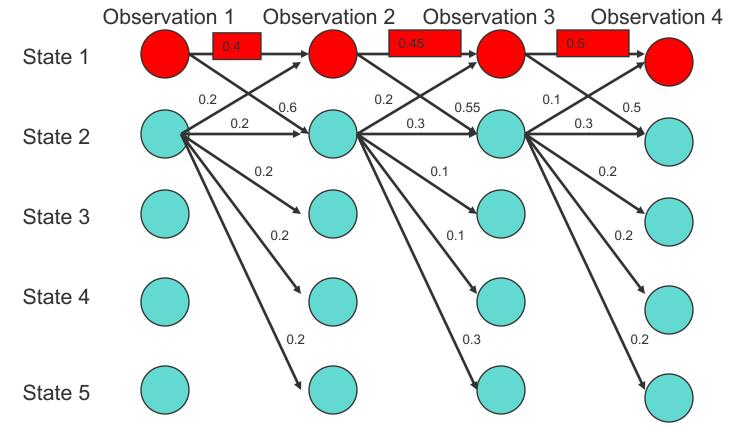


• Although locally it seems state 1 wants to go to state 2 and state 2 wants to remain in state 2.



[•] why?





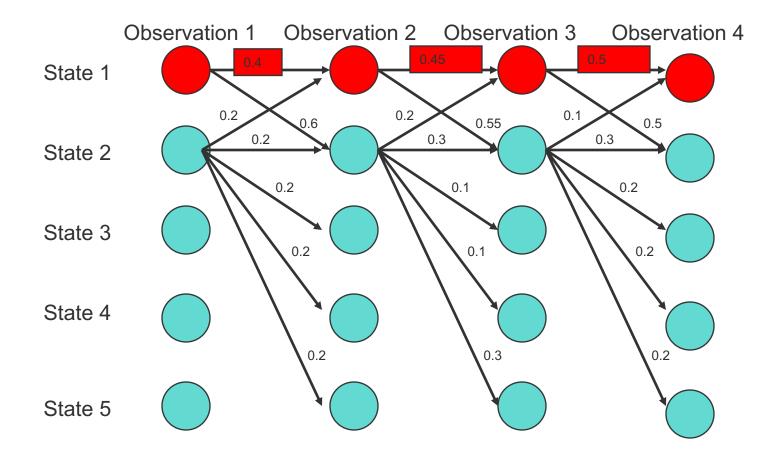
Most Likely Path: 1-> 1-> 1-> 1

• State 1 has only two transitions but state 2 has 5:

Average transition probability from state 2 is lower





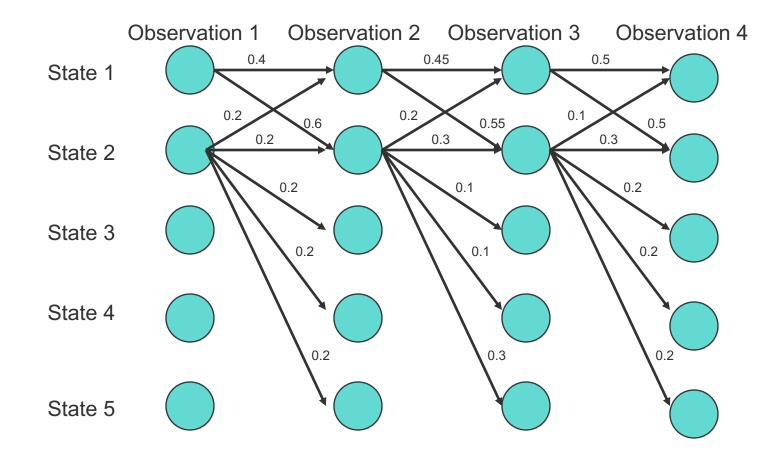


Label bias problem in HMM:

• Preference of states with lower number of transitions over others



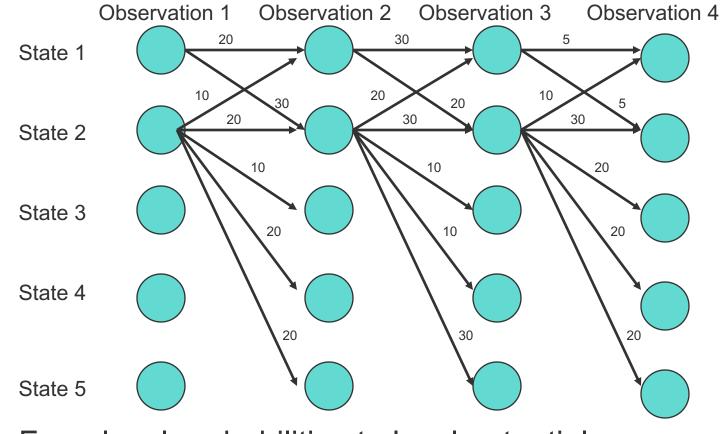
Solution: Do not normalize probabilities locally



From local probabilities



Solution: Do not normalize probabilities locally

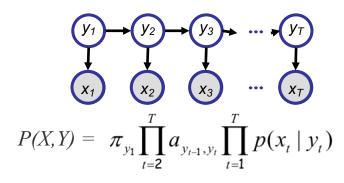


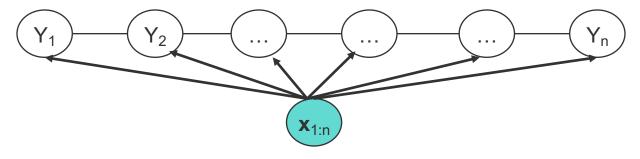
From local probabilities to local potentials

• States with lower transitions do not have an unfair advantage!









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

- CRF is a partially directed model
 - Discriminative model, unlike HMM
 - Usage of global normalizer Z(x) overcomes the label bias problem of HMM
 - Models the dependence between each state and the entire observation sequence





General parametric form:

 Y1
 Y2
 ...

$$P(\mathbf{y}|\mathbf{x}) = \frac{1}{Z(\mathbf{x},\lambda,\mu)} \exp\left(\sum_{i=1}^{n} \left(\sum_{k} \lambda_{k} f_{k}(y_{i},y_{i-1},\mathbf{x}) + \sum_{l} \mu_{l} g_{l}(y_{i},\mathbf{x})\right)\right)$$
$$= \frac{1}{Z(\mathbf{x},\lambda,\mu)} \exp\left(\sum_{i=1}^{n} \left(\lambda^{T} \mathbf{f}(y_{i},y_{i-1},\mathbf{x}) + \mu^{T} \mathbf{g}(y_{i},\mathbf{x})\right)\right)$$

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

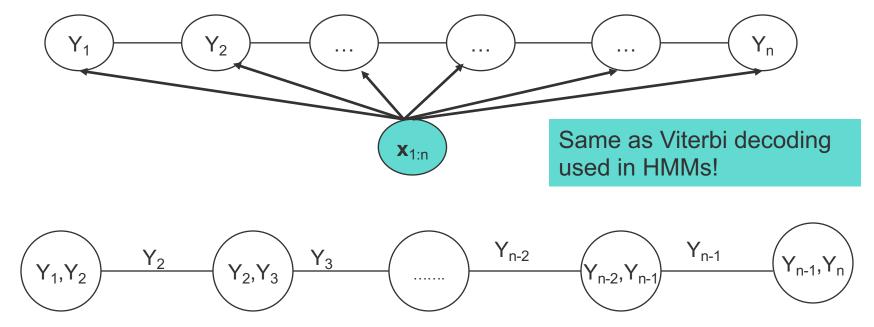




Given CRF parameters λ and μ , find the y^* that maximizes P(y|x)

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

- 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:







• Given $\{(\mathbf{x}_d, \mathbf{y}_d)\}_{d=1}^N$, find λ^* , μ^* such that

$$\lambda *, \mu * = \arg \max_{\lambda,\mu} L(\lambda,\mu) = \arg \max_{\lambda,\mu} \prod_{d=1}^{N} P(\mathbf{y}_{d} | \mathbf{x}_{d}, \lambda, \mu)$$

$$= \arg \max_{\lambda,\mu} \prod_{d=1}^{N} \frac{1}{Z(\mathbf{x}_{d}, \lambda, \mu)} \exp(\sum_{i=1}^{n} (\lambda^{T} \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_{d}) + \mu^{T} \mathbf{g}(y_{d,i}, \mathbf{x}_{d})))$$

$$= \arg \max_{\lambda,\mu} \sum_{d=1}^{N} (\sum_{i=1}^{n} (\lambda^{T} \mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_{d}) + \mu^{T} \mathbf{g}(y_{d,i}, \mathbf{x}_{d})) - \log Z(\mathbf{x}_{d}, \lambda, \mu))$$

• Computing the gradient w.r.t λ :

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

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



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

$$\square \text{ Computing the model expectations:}$$

Requires exponentially large number of summations: Is it intractable?

$$\sum_{\mathbf{y}} (P(\mathbf{y}|\mathbf{x}_d) \sum_{i=1}^n \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d)) = \sum_{i=1}^n (\sum_{\mathbf{y}} \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d) P(\mathbf{y}|\mathbf{x}_d))$$
$$= \sum_{i=1}^n \sum_{y_i, y_{i-1}} \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d) P(y_i, y_{i-1}|\mathbf{x}_d)$$
$$Expectation of \mathbf{f} \text{ over the corresponding marginal probability of neighboring nodes!!}$$

Can compute marginals using the sum-product algorithm on the chain





• Computing marginals using junction-tree calibration:

X_{1:n}

 Y_2

Junction Tree Initialization:

 $(Y_{1},Y_{2}) \xrightarrow{Y_{2}} (Y_{2},Y_{3}) \xrightarrow{Y_{3}} (Y_{n-2},Y_{n-1}) \xrightarrow{Y_{n-1}} (Y_{n-1},Y_{n})$

• After calibration:

$$P(y_i, y_{i-1} | \mathbf{x}_d) \propto \alpha(y_i, y_{i-1})$$
Also called
forward-backward algorithm

$$\Rightarrow P(y_i, y_{i-1} | \mathbf{x}_d) = \frac{\alpha(y_i, y_{i-1})}{\sum_{y_i, y_{i-1}} \alpha(y_i, y_{i-1})} = \alpha'(y_i, y_{i-1})$$

 $\alpha^0(y_i, y_{i-1}) = \exp(\lambda^T \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d))$





• Computing feature expectations using calibrated potentials:

$$\sum_{y_i, y_{i-1}} \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d) P(y_i, y_{i-1} | \mathbf{x}_d) = \sum_{y_i, y_{i-1}} \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d) \alpha'(y_i, y_{i-1})$$

• Now we know how to compute $r_{\lambda}L(\lambda,\mu)$:

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

=
$$\sum_{d=1}^{N} \left(\sum_{i=1}^{n} \left(\mathbf{f}(y_{d,i}, y_{d,i-1}, \mathbf{x}_d) - \sum_{y_i, y_{i-1}} \alpha'(y_i, y_{i-1}) \mathbf{f}(y_i, y_{i-1}, \mathbf{x}_d) \right) \right)$$

• Learning can now be done using gradient ascent:

$$\lambda^{(t+1)} = \lambda^{(t)} + \eta \nabla_{\lambda} L(\lambda^{(t)}, \mu^{(t)})$$

$$\mu^{(t+1)} = \mu^{(t)} + \eta \nabla_{\mu} L(\lambda^{(t)}, \mu^{(t)})$$



 In practice, we use a Gaussian Regularizer for the parameter vector to improve generalizability

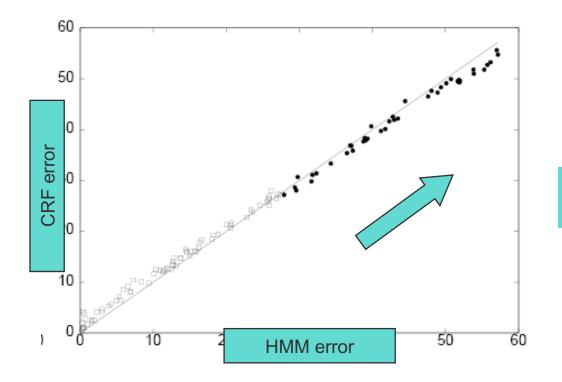
$$\lambda *, \mu * = \arg \max_{\lambda, \mu} \sum_{d=1}^{N} \log P(\mathbf{y}_d | \mathbf{x}_d, \lambda, \mu) - \frac{1}{2\sigma^2} (\lambda^T \lambda + \mu^T \mu)$$

- In practice, gradient ascent has very slow convergence
 - Alternatives:
 - Conjugate Gradient method
 - Limited Memory Quasi-Newton Methods





Comparison of error rates on synthetic data



Data is increasingly higher order in the direction of arrow

CRFs achieve the lowest error rate for higher order data





Parts of Speech tagging

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

⁺Using spelling features

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

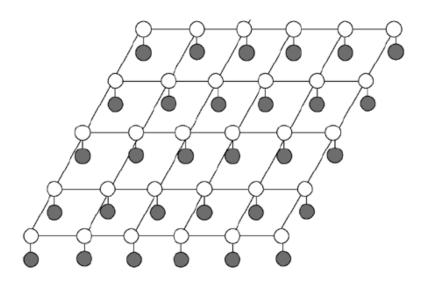


Supplementary





- So far we have discussed only 1dimensional chain CRFs
 - Inference and learning: exact
- We could also have CRFs for arbitrary graph structure
 - E.g: Grid CRFs
 - Inference and learning no longer tractable
 - Approximate techniques used
 - MCMC Sampling
 - Variational Inference
 - Loopy Belief Propagation
 - We will discuss these techniques SOON







Stereo Matching

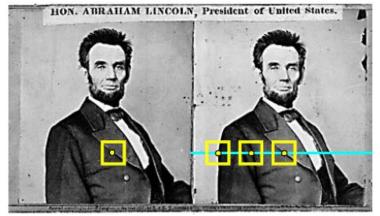


Image Segmentation



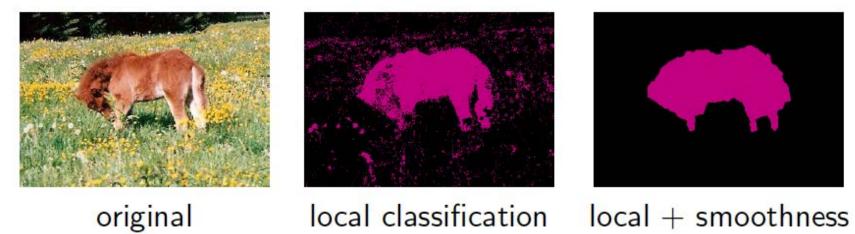
Image Restoration





Application: Image Segmentation

 $\begin{array}{l} \phi_i(y_i,x) \in \mathbb{R}^{\approx 1000} \colon \text{local image features, e.g. bag-of-words} \\ \rightarrow \quad \langle w_i, \phi_i(y_i,x) \rangle \colon \text{local classifier (like logistic-regression)} \\ \phi_{i,j}(y_i,y_j) = \llbracket y_i = y_j \rrbracket \in \mathbb{R}^1 \colon \text{test for same label} \\ \rightarrow \quad \langle w_{ij}, \phi_{ij}(y_i,y_j) \rangle \colon \text{penalizer for label changes (if } w_{ij} > 0) \\ \text{combined: } \operatorname{argmax}_y p(y|x) \text{ is smoothed version of local cues} \end{array}$



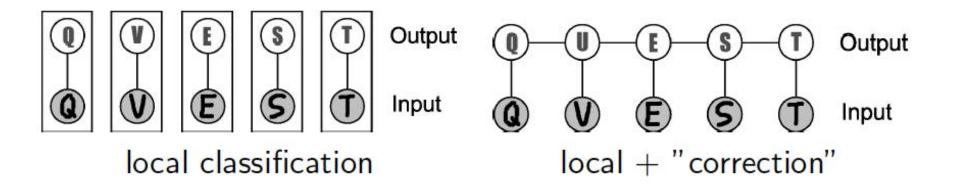


Application: Handwriting Recognition

 $\phi_i(y_i, x) \in \mathbb{R}^{\approx 1000}$: image representation (pixels, gradients) $\rightarrow \langle w_i, \phi_i(y_i, x) \rangle$: local classifier if x_i is letter y_i

 $\phi_{i,j}(y_i, y_j) = e_{y_i} \otimes e_{y_j} \in \mathbb{R}^{26 \cdot 26}$: letter/letter indicator $\rightarrow \langle w_{ij}, \phi_{ij}(y_i, y_j) \rangle$: encourage/suppress letter combinations

combined: $\operatorname{argmax}_y p(y|x)$ is "corrected" version of local cues



Application: Pose Estimation

 $\begin{array}{l} \phi_i(y_i,x) \in \mathbb{R}^{\approx 1000} \text{: local image representation, e.g. HoG} \\ \rightarrow \quad \langle w_i, \phi_i(y_i,x) \rangle \text{: local confidence map} \\ \phi_{i,j}(y_i,y_j) = \textit{good_fit}(y_i,y_j) \in \mathbb{R}^1 \text{: test for geometric fit} \\ \rightarrow \quad \langle w_{ij}, \phi_{ij}(y_i,y_j) \rangle \text{: penalizer for unrealistic poses} \end{array}$

together: $\operatorname{argmax}_{y} p(y|x)$ is sanitized version of local cues



original



local classification



local + geometry

Feature Functions for CRF in Vision

 $\phi_i(y_i, x)$: local representation, high-dimensional $\rightarrow \langle w_i, \phi_i(y_i, x) \rangle$: local classifier

 $\phi_{i,j}(y_i, y_j)$: prior knowledge, low-dimensional $\rightarrow \langle w_{ij}, \phi_{ij}(y_i, y_j) \rangle$: penalize outliers

learning adjusts parameters:

- unary w_i : learn local classifiers and their importance
- binary w_{ij} : learn importance of smoothing/penalization

 $\operatorname{argmax}_{y} p(y|x)$ is cleaned up version of local prediction





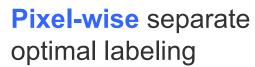
□ Image segmentation (FG/BG) by modeling of interactions btw RVs

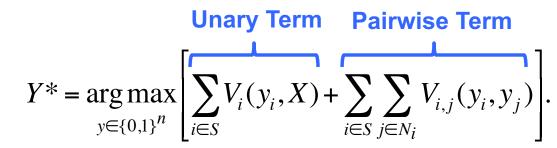
- Images are noisy.
- Objects occupy continuous regions in an image.



Input image







[Nowozin,Lampert 2012]



Locally-consistent joint optimal labeling

Y: labelsX: data (features)S: pixelsN_i: neighbors of pixel i



Discriminative Random Fields

- A special type of CRF
 - The unary and pairwise potentials are designed using local discriminative classifiers.
 - Posterior

$$P(Y \mid X) = \frac{1}{Z} \exp(\sum_{i \in S} A_i(y_i, X) + \sum_{i \in S} \sum_{j \in N_i} I_{ij}(y_i, y_j, X))$$

- Association Potential
 - Local discriminative model for site *i*: using logistic link with GLM.

 $A_i(y_i, X) = \log P(y_i | f_i(X)) \qquad P(y_i = 1 | f_i(X)) = \frac{1}{1 + \exp(-(w^T f_i(X)))} = \sigma(w^T f_i(X))$

- Interaction Potential
 - Measure of how likely site *i* and *j* have the same label given $X_{I_{ij}}(y_i, y_j, X) = ky_i y_j + (1-k)(2\sigma(y_i y_j \mu_{ij}(X)) - 1))$

(1) Data-independent smoothing term (2) Data-dependent pairwise logistic function

S. Kumar and M. Hebert. Discriminative Random Fields. IJCV, 2006.



Task: Detecting man-made structure in natural scenes.

■ Each image is divided in non-overlapping 16x16 tile blocks.

An example



Logistic: No smoothness in the labels

MRF: Smoothed False positive. Lack of neighborhood interaction of the data

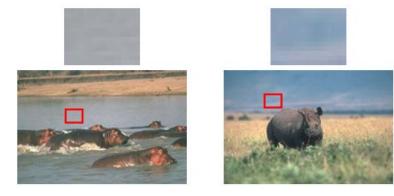
S. Kumar and M. Hebert. Discriminative Random Fields. IJCV, 2006.

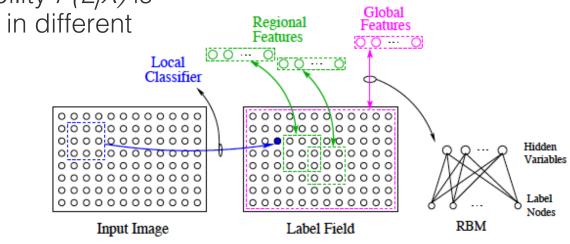


Multiscale Conditional Random Fields

- Considering features in different scales
 - Local Features (site)
 - Regional Label Features (small patch)
 - Global Label Features (big patch or the whole image)
- The conditional probability P(L|X) is formulated by features in different $\Re(a||e^{x}) = \frac{1}{Z} \prod_{s} P_{s}(L|X)$

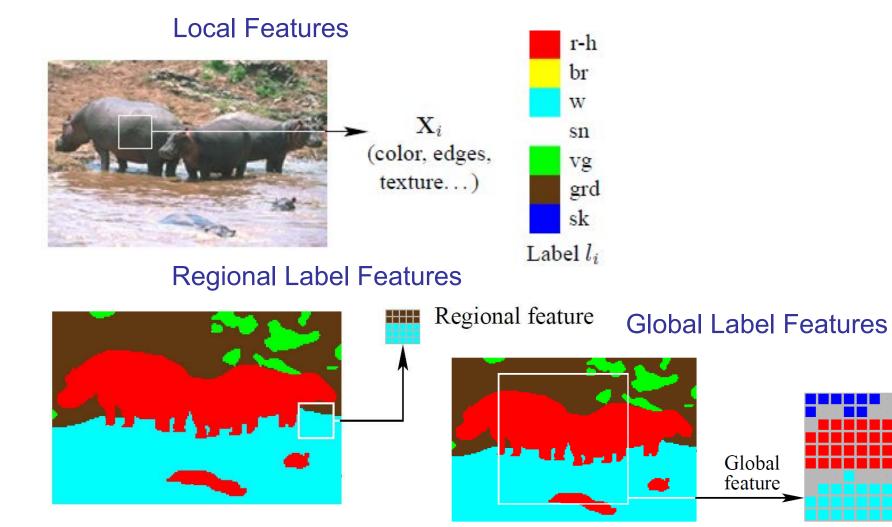
$$Z = \sum_{L} \prod_{s} P_{s}(L \mid X)$$



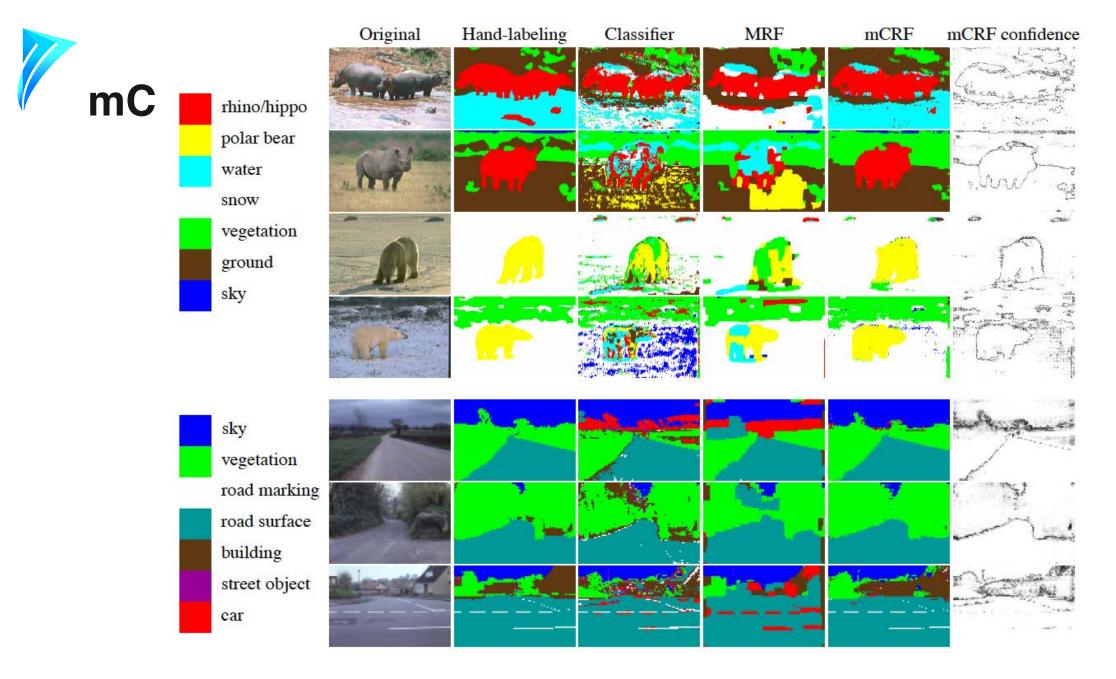




Multiscale Conditional Random Fields



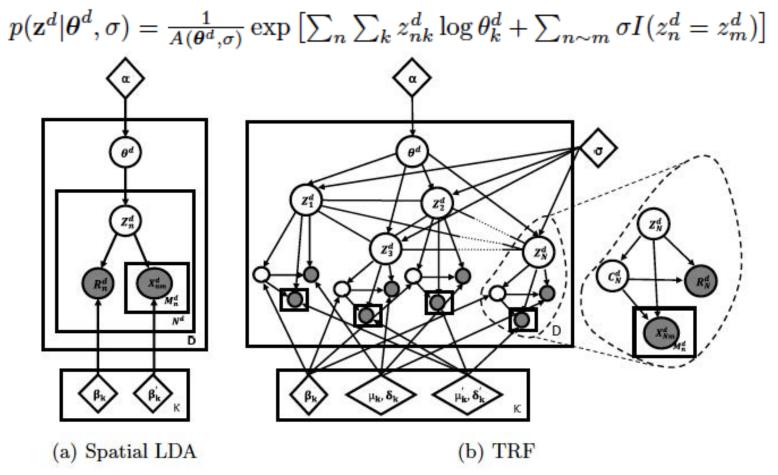
He, X. et. al.: Multiscale conditional random fields for image labeling. CVPR 2004 © Eric Xing @ CMU, 2005-2020



He, X. et. al.: Multiscale conditional random fields for image labeling. CVPR 2004



Spatial MRF over topic assignments

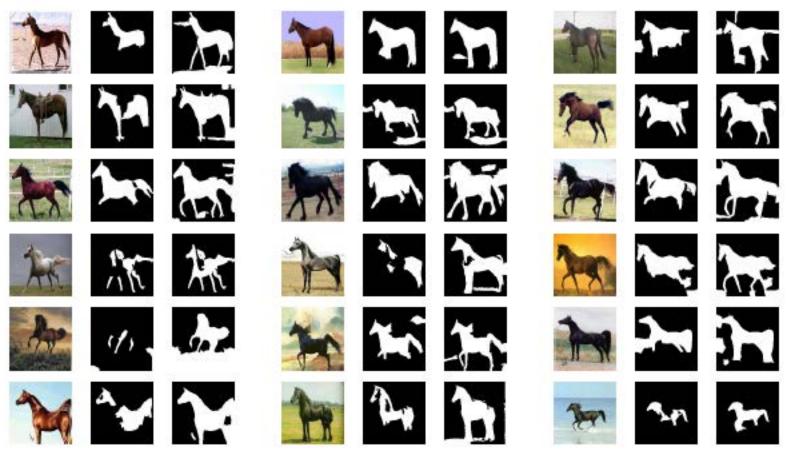


Zhao, B. et. al.: Topic random fields for image segmentation. ECCV 2010





Spatial LDA vs. Topic Random Fields



Zhao, B. et. al.: Topic random fields for image segmentation. ECCV 2010



- Conditional Random Fields are partially directed discriminative models
- They overcome the label bias problem of HMM by using a global normalizer
- Inference for 1-D chain CRFs is exact
 - Same as Max-product or Viterbi decoding
- Learning also is exact
 - globally optimum parameters can be learned
 - Requires using sum-product or forward-backward algorithm
- CRFs involving arbitrary graph structure are intractable in general
 - E.g.: Grid CRFs
 - Inference and learning require approximation techniques
 - MCMC sampling
 - Variational methods
 - Loopy BP

