PDEEC - Machine Learning 2018/19

Lecture
Context Dependent Classification
Sequential data
Hidden Markov Models

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Introduction



- Sets of data points assumed to be independent and identically distributed (i.i.d) so far
- ▶ i.i.d is a poor assumption for sequential data
 - measurements of time series (rainfall), daily values of a currency exchange rate, acoustic features in speech recognition
 - sequence of nucleotide base pairs along a strand of DNA, sequence of characters in an English sentence

Markov Model

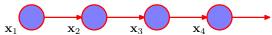
► In general

$$P(\mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_N) = P(\mathbf{x}_1) \prod_{n=2}^{N} P(\mathbf{x}_n | \mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_{n-1})$$

▶ Markov model: Each of the conditional distributions is independent of all previous observations except M most recent

The first-order Markov chain

Homogeneous Markov chain



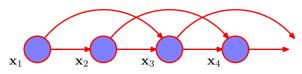
▶ Joint distribution for a sequence of N observations

$$P(\mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_N) = P(\mathbf{x}_1) \prod_{n=2}^{N} P(\mathbf{x}_n | \mathbf{x}_{n-1})$$

$$P(\mathbf{x}_n|\mathbf{x}_1,\mathbf{x}_2,\cdots,\mathbf{x}_{n-1})=P(\mathbf{x}_n|\mathbf{x}_{n-1})$$

A higher-order Markov chain

The second-order Markov chain



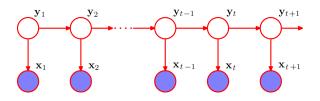
The joint distribution

$$P(\mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_N) = P(\mathbf{x}_1)P(\mathbf{x}_2|\mathbf{x}_1) \prod_{n=3}^{N} P(\mathbf{x}_n|\mathbf{x}_{n-1}, \mathbf{x}_{n-2})$$

- A higher-order Markov chain
- ▶ Suppose the observations are discrete variables having K states
- ▶ first-order: K-1 parameters for each K states $\to K(K-1)$ parameters
- ▶ Mth-order: $K^M(K-1)$ parameters



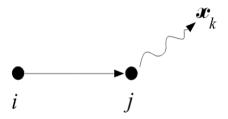
Hidden Markov models (HMM)



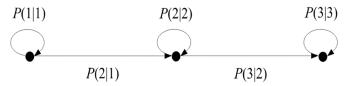
- $ightharpoonup \mathbf{y}_t$ latent variables (discrete)
- \triangleright \mathbf{x}_t observed variables
- ▶ The joint distribution of the state space model

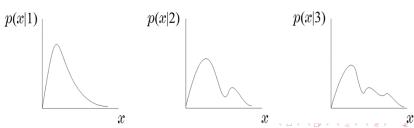
$$P(\mathbf{x}_1,\cdots,\mathbf{x}_T,\mathbf{y}_1,\cdots,\mathbf{y}_T) = P(\mathbf{y}_1) \prod_{t=2}^T P(\mathbf{y}_t|\mathbf{y}_{t-1}) \prod_{t=1}^T P(\mathbf{x}_t|\mathbf{y}_t)$$

- ▶ An HMM is a stochastic finite state automaton, that generates the observation sequence, $\mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_N$
- We assume that: The observation sequence is produced as a result of successive transitions between states, upon arrival at a state:



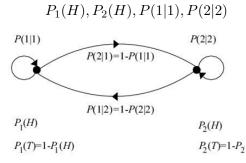
► This type of modeling is used for nonstationary stochastic processes that undergo distinct transitions among a set of different stationary processes.





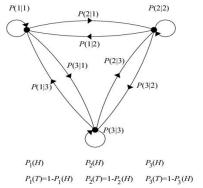
Example of HMM

► The two-coins case: Assume one of two coins is tossed behind a curtain. We observe a sequence of H or T. However, we have no access to know which coin was tossed. Identify one state for each coin. This is an example where states are not observable. H or T can be emitted from either state. The model depends on four parameters.



Example of HMM

▶ The three-coins case example is shown below:



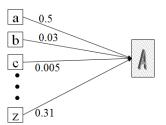
- ► Note that in all previous examples, specifying the model is equivalent to knowing:
 - ► The probability of each observation (H,T) to be emitted from each state.
 - lacktriangle The transition probabilities among states: P(i|j).

Word recognition example(I)

► Typed word recognition, assume all characters are separated.



► Character recognizer outputs probability of the image being particular character, P(image—character).

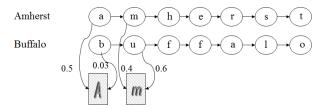


Word recognition example(II)

- Hidden states of HMM = characters.
- Observations = typed images of characters segmented from the image. Note that there is an infinite number of observations.
- ▶ Observation probabilities = character recognizer scores.
- Transition probabilities will be defined differently in two subsequent models.

Word recognition example(III)

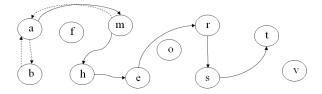
If lexicon is given, we can construct separate HMM models for each lexicon word.



- Here recognition of word image is equivalent to the problem of evaluating few HMM models
- This is an application of Evaluation problem.

Word recognition example(IV)

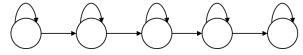
- We can construct a single HMM for all words.
- Hidden states = all characters in the alphabet.
- ► Transition probabilities and initial probabilities are calculated from language model.
- Observations and observation probabilities are as before



- ► Here we have to determine the best sequence of hidden states, the one that most likely produced word image.
- ▶ This is an application of Decoding problem

Character recognition with HMM example

▶ The structure of hidden states is chosen



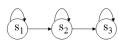
▶ Observations are feature vectors extracted from vertical slices.



- Probabilistic mapping from hidden state to feature vectors:
 - use mixture of Gaussian models.
 - Quantize feature vector space

Exercise: character recognition with HMM(I)

• The structure of hidden states:



- Observation = number of islands in the vertical slice.
- •HMM for character 'A':

Transition probabilities:
$$\{a_{ij}\}=\left(\begin{array}{ccc} .8 & .2 & 0 \\ 0 & .8 & .2 \\ 0 & 0 & 1 \end{array}\right)$$

Observation probabilities:
$$\{b_{jk}\}= \begin{pmatrix} .9 & .1 & 0 \\ .1 & .8 & .1 \\ .9 & .1 & 0 \end{pmatrix}$$



•HMM for character 'B':

Transition probabilities:
$$\{a_{ij}\}=\begin{bmatrix} .8 & .2 & 0 \\ 0 & .8 & .2 \\ 0 & 0 & 1 \end{bmatrix}$$

Observation probabilities:
$$\{b_{jk}\}=$$

$$\begin{bmatrix}
.9 & .1 & 0 \\
0 & .2 & .8 \\
.6 & 4 & 0
\end{bmatrix}$$

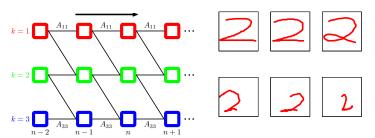


Exercise: character recognition with HMM(II)

- ▶ Suppose that after character image segmentation the following sequence of island numbers in 4 slices was observed: $\{1,3,2,1\}$
- ► What HMM is more likely to generate this observation sequence , HMM for 'A' or HMM for 'B' ?

HMM applications

- Speech recognition
- Natural language modelling
- Analysis of biological sequences (e.g. proteins and DNA)
- ▶ On-line handwriting recognition; Example: Handwritten digits
 - Left-to-right architecture
 - On-line data: each digit represented by the trajectory of the pen as a function of time



The Dishonest Casino !!!

A casino has two dice:

- Fair die P(1) = P(2) = P(3) = P(5) = P(6) = 1/6
- P Loaded die
 P(1) = P(2) = P(3) = P(5) =
 1/10
 P(6) = 1/2

In average, casino player switches back-&-forth between fair and loaded die once every 20 turns





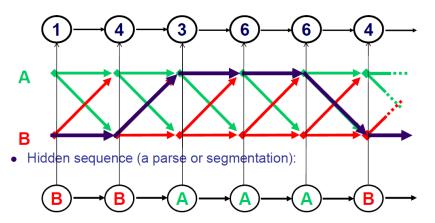
Main Questions Regarding the Dishonest Casino

GIVEN: A sequence of rolls by the casino player 1245526462146146136136661664661636616366163616515615115146123562344 QUESTION

- How likely is this sequence, given our model of how the casino works? (This is the EVALUATION problem)
- What portion of the sequence was generated with the fair die, and what portion with the loaded die? This is the DECODING question
- How "loaded" is the loaded die? How "fair" is the fair die? How often does the casino player change from fair to loaded, and back? This is the LEARNING question

Definition (of HMM)

• Observed sequence:



An HMM is a Stochastic Generative Model

Observation space

Alphabetic set: Euclidean space:

$$C = \{c_1, c_2, \cdots, c_{\kappa}\}$$

Index set of hidden states

$$I = \{1, 2, \cdots, M\}$$

Transition probabilities between any two states

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

or
$$p(y_t \mid y_{t-1}^i = 1) \sim \text{Multinomial}(a_{i,1}, a_{i,2}, \dots, a_{i,M}), \forall i \in \mathbb{I}.$$

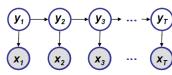
Start probabilities

$$p(y_1) \sim \text{Multinomial}(\pi_1, \pi_2, \dots, \pi_M)$$

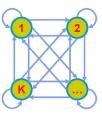
Emission probabilities associated with each state

$$p(x_t \mid y_t^j = 1) \sim \text{Multinomial}(b_{j,1}, b_{j,2}, \dots, b_{j,K}), \forall i \in I.$$
 or in general:

$$p(\mathbf{x}_{t} \mid \mathbf{y}_{t}^{i} = 1) \sim f(\cdot \mid \theta_{i}), \forall i \in I.$$

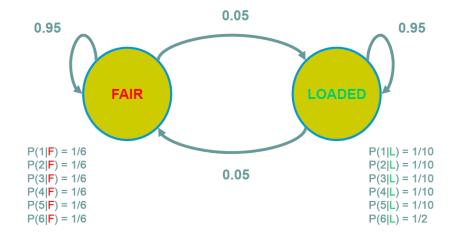


Graphical model



State automata

The Dishonest Casino Model



Three Main Questions on HMMs

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 $P(x \mid \theta)$

ALGO. Baum-Welch (EM)

Joint Probability

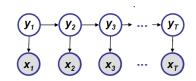
1245526462146146136136661664661636616366163156151515146123562344

▶ When the state-labeling is known, this is easy...

$$P(X,Y) = P(\mathbf{x}_1, \mathbf{x}_2, \cdots, \mathbf{x}_T, \mathbf{y}_1, \mathbf{y}_2, \cdots, \mathbf{y}_T)$$

Probability of a Parse

- Given a sequence x = x₁.....x_T
 and a parse y = y₁,, 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)
= $p(y_1) p(x_1 | y_1) p(y_2 | y_1) p(x_2 | y_2) \dots p(y_T | y_{T-1}) p(x_T | y_T)$
= $p(y_1) P(y_2 | y_1) \dots p(y_T | y_{T-1}) \times p(x_1 | y_1) p(x_2 | y_2) \dots p(x_T | y_T)$

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

Example: the Dishonest Casino

- Let the sequence of rolls be:
 - x = 1, 2, 1, 5, 6, 2, 1, 6, 2, 4



- Then, what is the likelihood of
 - y = Fair, Fair, Fair, Fair, Fair, Fair, Fair, Fair, Fair, Fair?
 (say initial probs a_{0Fair} = ½, a_{01 paded} = ½)

$$\frac{1}{2} \times (\frac{1}{6})^{10} \times (0.95)^9 = .00000000521158647211 = 5.21 \times 10^{-9}$$

Example: the Dishonest Casino

• So, the likelihood the die is fair in all this run is just 5.21×10^{-9}



- OK, but what is the likelihood of
 - π = Loaded, Loaded, Loaded, Loaded, Loaded, Loaded, Loaded, Loaded, Loaded?

$$\frac{1}{2} \times P(1 \mid Loaded) P(Loaded \mid Loaded) \dots P(4 \mid Loaded) =$$

$$\frac{1}{2} \times (\frac{1}{10})^8 \times (\frac{1}{2})^2 (0.95)^9 = .00000000078781176215 = 0.79 \times 10^{-9}$$

 Therefore, it is after all 6.59 times more likely that the die is fair all the way, than that it is loaded all the way

Example: the Dishonest Casino

Let the sequence of rolls be:

•
$$x = 1, 6, 6, 5, 6, 2, 6, 6, 3, 6$$



- Now, what is the likelihood $\pi = F, F, ..., F$?
 - $1/2 \times (1/6)^{10} \times (0.95)^9 = 0.5 \times 10^{-9}$, same as before
- What is the likelihood y = L, L, ..., L?

$$\frac{1}{2} \times (\frac{1}{10})^4 \times (\frac{1}{2})^6 (0.95)^9 = .00000049238235134735 = 5 \times 10^{-7}$$

• So, it is 100 times more likely the die is loaded

Marginal Probability

1245526462146146136136661664661636616366163616515615115146123562344

What if state-labeling Y is not observed

$$P(X) = \sum_{Y} P(X, Y)$$

the wrong way would be to...

The Forward Algorithm

We want to calculate $P(\mathbf{x})$, the likelihood of \mathbf{x} , given the HMM

• Sum over all possible ways of generating x:

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

ullet To avoid summing over an exponential number of paths ${f y}$, define

$$\alpha(\boldsymbol{y}_{t}^{k}=1)=\alpha_{t}^{k}\stackrel{\mathrm{def}}{=}P(\boldsymbol{x}_{1},...,\boldsymbol{x}_{t},\boldsymbol{y}_{t}^{k}=1) \tag{the forward probability}$$

The recursion:

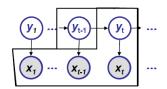
$$\alpha_t^k = p(x_t \mid y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k}$$

$$P(\mathbf{x}) = \sum_k \alpha_t^k$$

The Forward Algorithm

Compute the forward probability:

$$\alpha_t^k = P(x_1, ..., x_{t-1}, x_t, y_t^k = 1)$$



$$\begin{split} &= \sum_{y_{t-1}} P(x_1, ..., x_{t-1}, y_{t-1}) P(y_t^k = 1 \mid y_{t-1}, x_1, ..., x_{t-1}) P(x_t \mid y_t^k = 1, x_1, ..., x_{t-1}, y_{t-1}) \\ &= \sum_{y_{t-1}} P(x_1, ..., x_{t-1}, y_{t-1}) P(y_t^k = 1 \mid y_{t-1}) P(x_t \mid y_t^k = 1) \\ &= P(x_t \mid y_t^k = 1) \sum_{i} P(x_1, ..., x_{t-1}, y_{t-1}^i = 1) P(y_t^k = 1 \mid y_{t-1}^i = 1) \\ &= P(x_t \mid y_t^k = 1) \sum_{i} \alpha_{t-1}^i a_{i,k} \end{split}$$

Chain rule: $P(A, B, C) = P(A)P(B \mid A)P(C \mid A, B)$

The Forward Algorithm

• We can compute α_t^k for all k, t, using dynamic programming!

Initialization:

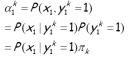
$$\alpha_1^k = P(x_1 \mid y_1^k = 1)\pi_k$$

Iteration:

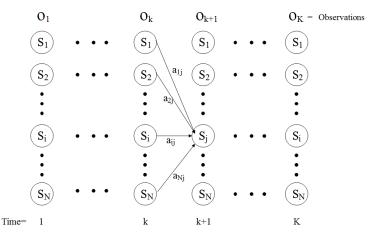
$$\alpha_t^k = P(x_t \mid y_t^k = 1) \sum_i \alpha_{t-1}^i a_{i,k}$$

Termination:

$$P(\mathbf{x}) = \sum_{k} \alpha_{T}^{k}$$



Recognition: Any path method



The Forward Algorithm

- we are exploiting the graph structure (the joint distribution) to do an efficient inference
- the algorithm can be expressed in terms of the propagation of local messages around the graph
- each summation effectively removes a variable from the distribution, which can be viewed as the removal of a node from the graph

The Backward Algorithm

Define the backward probability:

Define the backward probability:
$$\beta_t^k = P(x_{t+1},...,x_T \mid y_t^k = 1) \\ = \sum_{y_{t+1}} P(x_{t+1},...,x_T,y_{t+1} \mid y_t^k = 1) \\ = \sum_{i} P(y_{t+1}^i = 1 \mid y_t^k = 1) P(x_{t+1} \mid y_{t+1}^i = 1,y_t^k = 1) P(x_{t+2},...,x_T \mid x_{t+1},y_{t+1}^i = 1,y_t^k = 1) \\ = \sum_{i} P(y_{t+1}^i = 1 \mid y_t^k = 1) P(x_{t+1} \mid y_{t+1}^i = 1) P(x_{t+2},...,x_T \mid y_{t+1}^i = 1) \\ = \sum_{i} a_{k,i} P(x_{t+1} \mid y_{t+1}^i = 1) \beta_{t+1}^i$$

Chain rule:
$$P(A, B, C \mid \alpha) = P(A \mid \alpha)P(B \mid A, \alpha)P(C \mid A, B, \alpha)$$

The Backward Algorithm

• We can compute β_t^k for all k, t, using dynamic programming!

Initialization:

$$\beta_T^k = 1, \ \forall k$$

Iteration:

$$\beta_t^k = \sum_i a_{k,i} P(x_{t+1} \mid y_{t+1}^i = 1) \beta_{t+1}^i$$

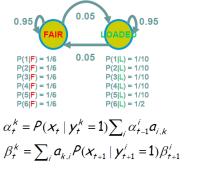
Termination:

$$P(\mathbf{x}) = \sum_{k} \alpha_1^k \beta_1^k$$

Example

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

Alpha (actual)		Beta (actua	Beta (actual)	
0.0833	0.0500	0.0000 0.0	000	
0.0136	0.0052	0.0000 0.0	000	
0.0022	0.0006	0.0000 0.0	000	
0.0004	0.0001	0.0000 0.0	000	
0.0001	0.0000	0.0001 0.0	001	
0.0000	0.0000	0.0007 0.0	006	
0.0000	0.0000	0.0045 0.0	055	
0.0000	0.0000	0.0264 0.0	112	
0.0000	0.0000	0.1633 0.1	033	
0.0000	0.0000	1.0000 1.0	000	



What is the probability of a hidden state prediction?

A single state:

$$P(\mathbf{y}_t|\mathbf{X})$$

▶ What about a hidden state sequence ?

$$P(\mathbf{y}_1,\cdots,\mathbf{y}_T)|\mathbf{X})$$

Posterior decoding

We can now calculate

$$P(\mathbf{y}_t^k = 1 \mid \mathbf{x}) = \frac{P(\mathbf{y}_t^k = 1, \mathbf{x})}{P(\mathbf{x})} = \frac{\alpha_t^k \beta_t^k}{P(\mathbf{x})}$$

- Then, we can ask
 - What is the most likely state at position *t* of sequence **x**:

$$\mathbf{k}_{t}^{*} = \operatorname{arg\,max}_{k} P(\mathbf{y}_{t}^{k} = 1 \mid \mathbf{x})$$

- 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: $\left\{ y_t^{k_t^*} = 1 : t = 1 \cdots T \right\}$
- This is different from MPA of a whole sequence states
- This can be understood as bit error rate
 vs. word error rate

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

of hidden

ormuden			
X	y	P(x,y)	
0	0	0.35	
0	1	0.05	
1	0	0.3	
1	1	0.3	

Viterbi decoding

GIVEN x = x₁, ..., x_T, we want to find y = y₁, ..., y_T, such that P(y|x) is maximized:

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

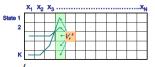
Let

$$V_{t}^{k} = \max_{\{y_{1},...,y_{t-1}\}} P(x_{1},...,x_{t-1},y_{1},...,y_{t-1},x_{t},y_{t}^{k} = 1)$$

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

The recursion:

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



- Underflows are a significant problem $p(x_1,...,x_r,y_1,...,y_r) = \pi_{v_1} a_{v_1,v_2} \cdots a_{v_{r-1},v_r} b_{v_r,x_r} \cdots b_{v_r,x_r}$
 - These numbers become extremely small underflow
 - I hese 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

Define the viterbi probability:

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

The Viterbi Algorithm

▶ Input: $\mathbf{x} = \mathbf{x}_1, \cdots, \mathbf{x}_T$

Initialization:

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

Iteration:

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

$$Ptr(\mathbf{k}, \mathbf{t}) = \arg\max_{i} \mathbf{a}_{i,k} V_{t-1}^{i}$$

Termination:

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

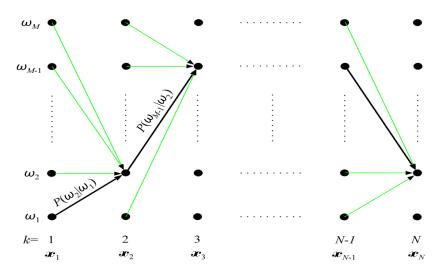
TraceBack:

$$\mathbf{y}_{T}^{*} = \operatorname{arg\,max}_{k} \mathbf{V}_{T}^{k}$$

$$\boldsymbol{y}_{t-1}^* = \operatorname{Ptr}(\boldsymbol{y}_t^*, \boldsymbol{t})$$

Context Dependent Classification

The Viterbi Algorithm



Computational Complexity and implementation details

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

$$\alpha_{t}^{k} = p(x_{t} | y_{t}^{k} = 1) \sum_{i} \alpha_{t-1}^{i} a_{i,k}$$

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

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

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

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 $\theta = (\pi_i, a_{ii}, \eta_{ik})$ that maximize $P(x \mid \theta)$

ALGO. Baum-Welch (EM)

Learning HMM: two scenarios

- Supervised learning: estimation when the "right answer" is known
 - Examples: GIVEN: a genomic region $x = x_1 \cdots x_1 \ _{000 \ 000}$ 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: 10000 rolls of the casino player, but we don't see when he changes dice
- ▶ QUESTION: Update the parameters θ of the model to maximize $P(\mathbf{X}|\theta)$ Maximal likelihood (ML) estimation

Supervised ML estimation

- Given $x = x_1...x_N$ for which the true state path $y = y_1...y_N$ 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:

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

$$b_{ik}^{ML} = \frac{\#(i \to k)}{\#(i \to \bullet)} = \frac{\sum_{n} \sum_{t=1}^{T} y_{n,t}^{j} x_{n,t}^{k}}{\sum_{n} \sum_{t=1}^{T} y_{n,t}^{j}} = \frac{B_{ik}}{\sum_{k} B_{ik}}$$

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

Supervised ML estimation

• 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 <u>overfitting</u>:
 - P(x|θ) is maximized, but θ 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

• Then:
$$a_{FF} = 1$$
; $a_{FL} = 0$
 $b_{F1} = b_{F3} = .2$;
 $b_{E2} = .3$; $b_{E4} = 0$; $b_{E5} = b_{E6} = .1$

Pseudocounts

- 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 \mathbf{y} emits k in \mathbf{x} + S_{ik}
```

- R_{ij} , S_{ij} 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 ⇒ strong prior belief
- Small total pseudocounts: just to avoid 0 probabilities --smoothing

Unsupervised ML estimation

• Given $x = x_1...x_N$ for which the true state path $y = y_1...y_N$ is unknown,

EXPECTATION MAXIMIZATION

- Starting with our best guess of a model M, parameters θ .
- 1. Estimate A_{ii} , B_{ik} in the training data
 - How? $A_{ij} = \sum_{n,t} \langle y'_{n,t-1} y'_{n,t} \rangle$ $B_{ik} = \sum_{n,t} \langle y'_{n,t} \rangle x'_{n,t}$,
 - Update θ according to A_{ij} , B_{ik}
 - Now a "supervised learning" problem
- 2. Repeat 1 & 2, until convergence

This is called the Baum-Welch Algorithm

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

Unsupervised ML estimation

The Baum Welch algorithm

• The complete log likelihood

$$\ell_{c}(\theta; \mathbf{x}, \mathbf{y}) = \log p(\mathbf{x}, \mathbf{y}) = \log \prod_{n} \left(p(y_{n,1}) \prod_{t=2}^{T} p(y_{n,t} \mid y_{n,t-1}) \prod_{t=1}^{T} p(x_{n,t} \mid x_{n,t}) \right)$$

The expected complete log likelihood

$$\ell_{e}(\boldsymbol{\theta}; \mathbf{x}, \mathbf{y}) \rangle = \sum_{n} \left(\left\langle \boldsymbol{y}_{n,1}^{i} \right\rangle_{\rho(y_{n,1}|\mathbf{x}_{n})} \log \pi_{i} \right) + \sum_{n} \sum_{t=2}^{T} \left(\left\langle \boldsymbol{y}_{n,t-1}^{i} \boldsymbol{y}_{n,t}^{j} \right\rangle_{\rho(y_{n,t-1},y_{n})\mathbf{x}_{n})} \log \boldsymbol{a}_{i,j} \right) + \sum_{n} \sum_{t=1}^{T} \left(\boldsymbol{x}_{n,t}^{k} \left\langle \boldsymbol{y}_{n,t}^{i} \right\rangle_{\rho(y_{n,t}|\mathbf{x}_{n})} \log \boldsymbol{b}_{i,k} \right)$$

- EM
 - The E step

$$\begin{aligned} \gamma_{n,t}^{i} &= \left\langle \mathbf{y}_{n,t}^{i} \right\rangle = p(\mathbf{y}_{n,t}^{i} = 1 \mid \mathbf{x}_{n}) \\ \xi_{n,t}^{i,j} &= \left\langle \mathbf{y}_{n,t-1}^{i} \mathbf{y}_{n,t}^{j} \right\rangle = p(\mathbf{y}_{n,t-1}^{i} = 1, \mathbf{y}_{n,t}^{j} = 1 \mid \mathbf{x}_{n}) \end{aligned}$$

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

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

Unsupervised ML estimation

The Baum Welch algorithm

Maximum likelihood for the HMM

- ▶ We have observed a data set $\{\mathbf{x}_1, \dots, \mathbf{x}_T\}$ (or possibly multiples sequences)
- so we can determine the parameters of an HMM

$$\theta = \{\pi, A, \phi\}$$

by using maximum likelihood (ϕ represent the parameters of the emission probabilities).

▶ The likelihood function is

$$p(\mathbf{X}|\theta) = \sum_{\mathbf{y}} p(\mathbf{X}, \mathbf{y}|\theta)$$

Unsupervised ML estimation

The Baum Welch algorithm

Maximizing the likelihood function Expectation maximization algorithm (EM)

- ▶ Initial selection for the model parameters: θ^{old}
- ► E step:
 - ▶ Posterior distribution of the latent variables $p(y|X, \theta^{\text{old}})$

$$Q(\theta, \theta^{\mathsf{old}}) = \sum_{\mathbf{y}} p(\mathbf{y}|\mathbf{X}, \theta^{\mathsf{old}}) \ln p(\mathbf{X}, \mathbf{y}|\theta)$$

Unsupervised ML estimation

The Baum Welch algorithm

Maximizing the likelihood function: EM E step:

$$Q(\theta, \theta^{\mathsf{old}}) = \sum_{k=1}^{K} \gamma(y_{1k}) \ln \pi_k + \sum_{t=2}^{T} \sum_{j=1}^{K} \sum_{k=1}^{K} \xi(\mathbf{y}_{t-1,j}, \mathbf{y}_{tk}) \ln A_{jk}$$
$$+ \sum_{t=1}^{T} \sum_{k=1}^{K} \gamma(\mathbf{y}_{tk}) \ln p(\mathbf{x}_t | \phi_k)$$

▶ The marginal posterior distribution of a latent variable γ and the joint posterior distribution of two successive latent variables ξ

$$\gamma(\mathbf{y}_t) = p(\mathbf{y}_t | \mathbf{X}, \theta^{\mathsf{old}})$$
$$\xi(\mathbf{y}_{t-1}, \mathbf{y}_t) = p(\mathbf{y}_{t-1}, \mathbf{y}_t | \mathbf{X}, \theta^{\mathsf{old}})$$

Unsupervised ML estimation

The Baum Welch algorithm

Maximizing the likelihood function: EM

M step:

Maximize $Q(\theta, \theta^{\mathsf{old}})$ with respect to parameters $\theta = \{\pi, A, \phi\}$, treat $\gamma(\mathbf{y}_t)$ and $\xi(\mathbf{y}_{t-1}, \mathbf{y}_t)$ as constant. By using Lagrange multipliers

$$\pi_k = \frac{\gamma(y_{1k})}{\sum_{j=1}^K \gamma(y_{1j})}$$

$$A_{jk} = \frac{\sum_{t=2}^T \xi(y_{t-1,j}, y_{tk})}{\sum_{t=1}^K \sum_{t=2}^T \xi(y_{t-1,j}, y_{tl})}$$

Unsupervised ML estimation

The Baum Welch algorithm

Maximizing the likelihood function: EM M step:

- ightharpoonup Parameters ϕ_k independent
 - \rightarrow for Gaussian emission densities $p(\mathbf{x}|\phi_k) = \mathcal{N}(\mathbf{x}|\mu_k, \Sigma_k)$

$$\mu_k = \frac{\sum_{t=1}^{T} \gamma(y_{tk}) \mathbf{x}_t}{\sum_{t=1}^{T} \gamma(y_{tk})}$$

$$\Sigma_{k} = \frac{\sum_{t=1}^{T} \gamma(y_{tk}) (\mathbf{x}_{t} - \mu_{k}) (\mathbf{x}_{t} - \mu_{k})^{T}}{\sum_{t=1}^{T} \gamma(y_{tk})}$$

Unsupervised ML estimation

The Baum Welch algorithm

$$\gamma(\mathbf{y}_t) = p(\mathbf{y}_t | \mathbf{X}) = \frac{p(\mathbf{X} | \mathbf{y}_t) p(\mathbf{y}_t)}{p(\mathbf{X})} = \frac{\alpha(\mathbf{y}_t) \beta(\mathbf{y}_t)}{p(\mathbf{X})}$$
$$\xi(\mathbf{y}_{t-1}, \mathbf{y}_t) = p(\mathbf{y}_{t-1}, \mathbf{y}_t | \mathbf{X}) = \frac{p(\mathbf{X} | y_{t-1}, \mathbf{y}_t) p(\mathbf{y}_{t-1}, \mathbf{y}_t)}{p(\mathbf{X})} = \frac{\alpha(\mathbf{y}_{t-1}) p(\mathbf{x}_t | \mathbf{y}_t) p(\mathbf{y}_t | \mathbf{y}_{t-1}) \beta(\mathbf{y}_t)}{p(\mathbf{X})}$$

Unsupervised ML estimation

The Baum Welch algorithm - comments

Time Complexity: # iterations $\times \mathcal{O}(K^2N)$

- 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: Over-fitting

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