CS344: Introduction to Artificial Intelligence (associated lab: CS386)

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Lecture 9: Viterbi; forward and backward probabilities

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HMM Definition

- Set of states: S where |S|=N
- Start state $S_0 / *P(S_0) = 1*/$
- Output Alphabet: O where |O|=M
- Transition Probabilities: A= {a_{ij}} /*state i to state j*/
- Emission Probabilities : $B = \{b_j(o_k)\} / *prob.$ of emitting or absorbing o_k from state j*/
- Initial State Probabilities: $\Pi = \{p_1, p_2, p_3, ..., p_N\}$
- Each $p_i = P(o_0 = \varepsilon, S_i | S_0)$

Markov Processes

- Properties
 - Limited Horizon: Given previous t states, a state i, is independent of preceding 0 to tk+1 states.
 - $P(X_t=i/X_{t-1}, X_{t-2}, ..., X_0) = P(X_t=i/X_{t-1}, X_{t-2}, ..., X_{t-k})$
 - Order k Markov process
 - Time invariance: (shown for k=1)
 - $P(X_t=i/X_{t-1}=j) = P(X_1=i/X_0=j) \dots = P(X_n=i/X_{n-1}=j)$

Three basic problems (contd.)

- Problem 1: Likelihood of a sequence
 - Forward Procedure
 - Backward Procedure
- Problem 2: Best state sequence
 - Viterbi Algorithm
- Problem 3: Re-estimation
 - Baum-Welch (Forward-Backward Algorithm)

Probabilistic Inference

- O: Observation Sequence
- S: State Sequence
- Given O find S* where $S^* = \arg \max_{S} p(S/O)$ called Probabilistic Inference
- Infer "Hidden" from "Observed"
- How is this inference different from logical inference based on propositional or predicate calculus?

Essentials of Hidden Markov Model

- 1. Markov + Naive Bayes
- 2. Uses both transition and observation probability

$$p(S_k \to^{O_k} S_{k+1}) = p(O_k / S_k) p(S_{k+1} / S_k)$$

3. Effectively makes Hidden Markov Model a Finite State Machine (FSM) with probability

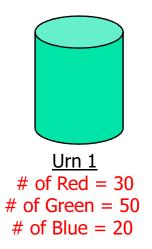
Probability of Observation Sequence

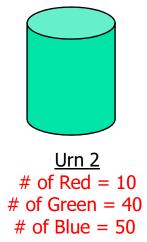
$$p(O) = \sum_{S} p(O, S)$$
$$= \sum_{S} p(S) p(O/S)$$

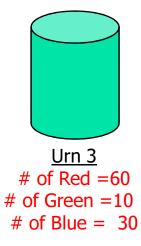
- Without any restriction,
 - Search space size= |S||0|

Continuing with the Urn example

Colored Ball choosing







Example (contd.)

Transition Probability

G	İν	e	n	:

	U_1	U ₂	U_3
U_1	0.1	0.4	0.5
U_2	0.6	0.2	0.2
U_3	0.3	0.4	0.3

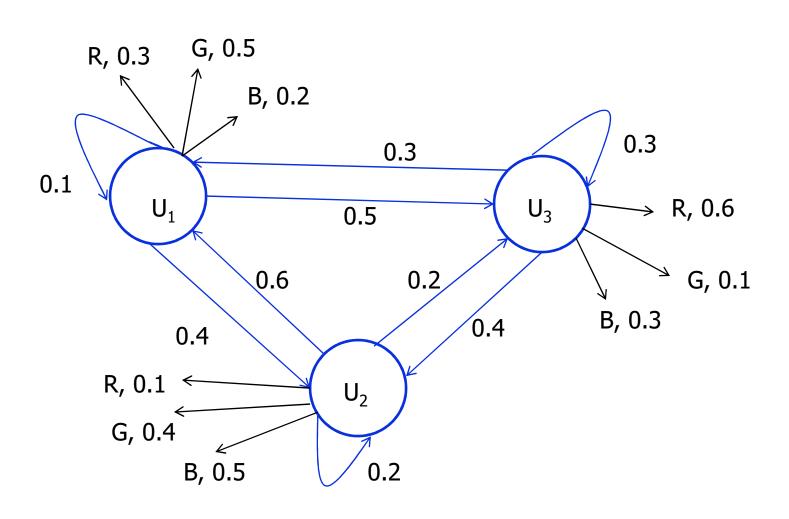
and

	R	G	В
U_1	0.3	0.5	0.2
U_2	0.1	0.4	0.5
U_3	0.6	0.1	0.3

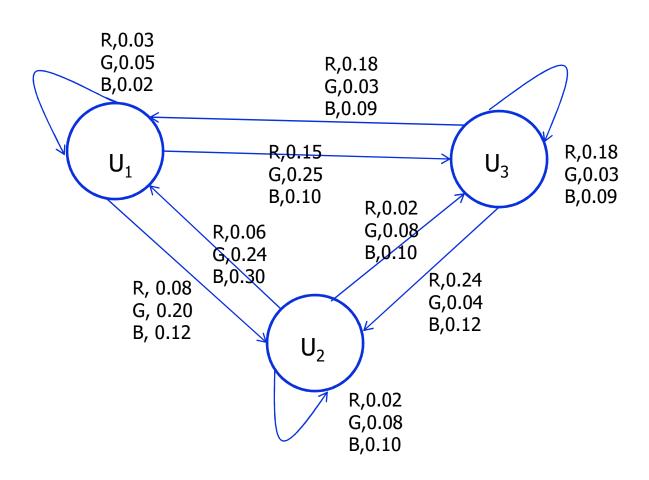
Observation: RRGGBRGR

What is the corresponding state sequence?

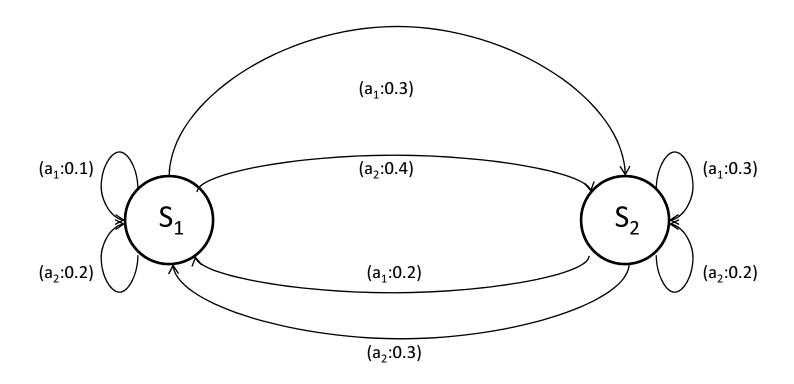
Diagrammatic representation (1/2)



Diagrammatic representation (2/2)



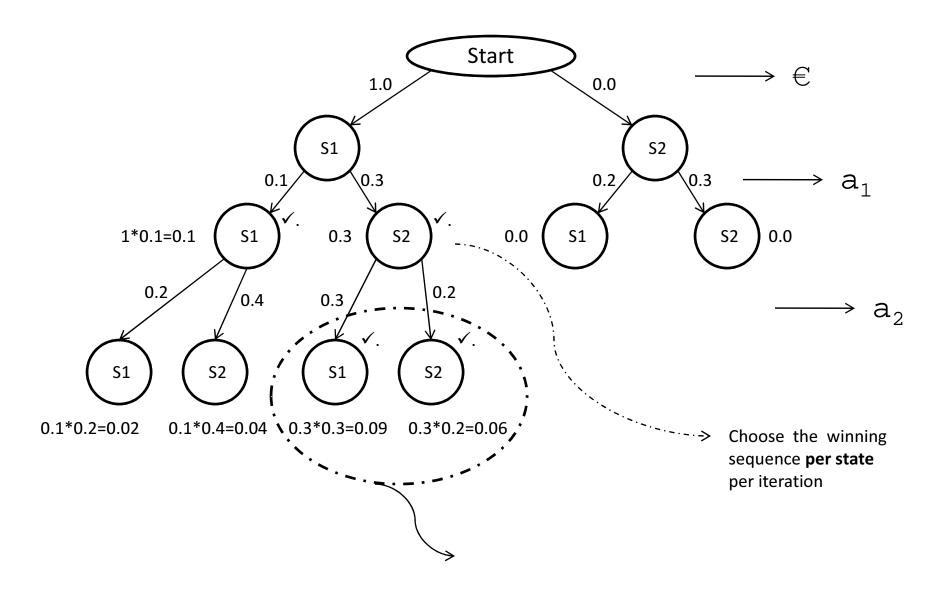
Probabilistic FSM



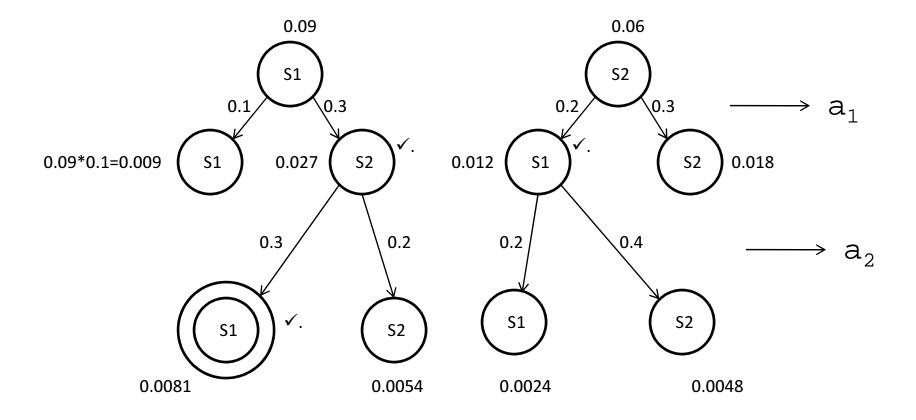
The question here is:

"what is the most likely state sequence given the output sequence seen"

Developing the tree



Tree structure contd...



The problem being addressed by this tree is $S^* = \arg\max_s P(S \mid a_1 - a_2 - a_1 - a_2, \mu)$

a1-a2-a1-a2 is the output sequence and μ the model or the machine

Tabular representation of the tree

Latest symbol observed Ending state	€	a_1	a ₂	a ₁	a ₂
S_1	1.0	(1.0*0.1,0.0*0.2)=(0.1 ,0.0)	(0.02, 0.09)	(0.009, 0.012)	(0.0024, 0,0081)
S ₂	0.0	(1.0*0.3,0.0*0.3)=(0.3 ,0.0)	(0.04, 0.0 6)	(0.027 ,0.018)	(0.0048,0.005 4)

Note: Every cell records the winning probability ending in that state

Final winner

The bold faced values in each cell shows the sequence probability ending in that state. Going backward from final winner sequence which ends in state S_2 (indicated By the 2^{nd} tuple), we recover the sequence.

Algorithm

(following James Alan, Natural Language Understanding (2nd edition), Benjamin Cummins (pub.), 1995

Given:

- 1. The HMM, which means:
 - a. Start State: S₁
 - b. Alphabet: $A = \{a_1, a_2, ... a_p\}$
 - Set of States: $S = \{S_1, S_2, ... S_n\}$
 - d. Transition probability $P(S_i \xrightarrow{a_k} S_j)$ $\forall_{i, j, k}$ which is equal to $P(S_j, a_k \mid S_i)$
- 2. The output string $a_1 a_2 ... a_T$

To find:

The most likely sequence of states $C_1C_2...C_T$ which produces the given output sequence, *i.e.*, $C_1C_2...C_T = \underset{C}{\operatorname{arg\,max}}[P(C \mid a_1, a_2, ...a_T, \mu]$

Algorithm contd...

Data Structure:

- A N*T array called SEQSCORE to maintain the winner sequence always (N=#states, T=length of o/p sequence)
- 2. Another N*T array called BACKPTR to recover the path.

Three distinct steps in the Viterbi implementation

- Initialization
- Iteration
- 3. Sequence Identification

1. Initialization

```
SEQSCORE(1,1)=1.0
BACKPTR(1,1)=0
For(i=2 to N) do
SEQSCORE(i,1)=0.0
[expressing the fact that first state is S_1]
```

2. Iteration

```
For(t=2 to T) do

For(i=1 to N) do

SEQSCORE(i,t) = Max_{(j=1,N)}

[SEQSCORE (j,(t-1)) * P(Sj \longrightarrow Si)]

BACKPTR(I,t) = index j that gives the MAX above
```

3. Seq. Identification

```
C(T) = i that maximizes SEQSCORE(i,T)
For i from (T-1) to 1 do
C(i) = BACKPTR[C(i+1),(i+1)]
```

Optimizations possible:

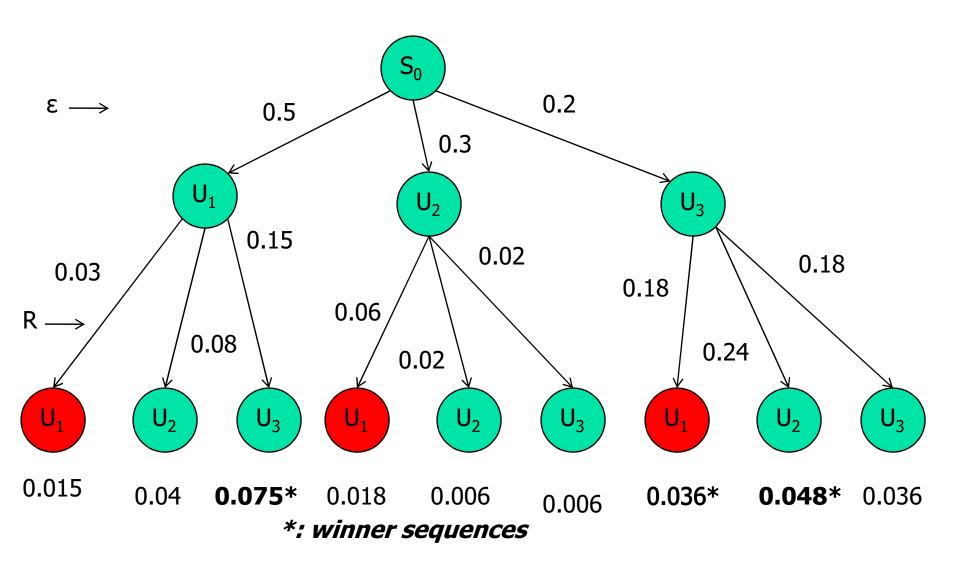
- 1. BACKPTR can be 1*T
- 2. SEQSCORE can be T*2

Homework:- Compare this with A*, Beam Search [Homework]

Reason for this comparison:

Both of them work for finding and recovering sequence

Viterbi Algorithm for the Urn problem (first two symbols)



Markov process of order>1 (say 2)

```
O_1 O_2 O_3
                   O_4
                        O_5 O_6
                                 O_7
                                      O_8
  O_0
Obs: ε R R G G B
                             R G
                                      R
              S_3
                   S_4
                        S<sub>5</sub>
                                       S_8
State: S_0 S_1 S_2
                             S_6
                                         S
```

Same theory works P(S).P(O|S)

=
$$P(O_0|S_0).P(S_1|S_0).$$

 $[P(O_1|S_1). P(S_2|S_1S_0)].$
 $[P(O_2|S_2). P(S_3|S_2S_1)].$
 $[P(O_3|S_3).P(S_4|S_3S_2)].$
 $[P(O_4|S_4).P(S_5|S_4S_3)].$
 $[P(O_5|S_5).P(S_6|S_5S_4)].$

$$[P(O_7|S_7).P(S_8|S_7S_6)].$$

 $[P(O_6|S_6).P(S_7|S_6S_5)].$

 $[P(O_8|S_8).P(S_9|S_8S_7)].$

We introduce the states S_0 and S_9 as initial and final states respectively.

After S_8 the next state is S_9 with probability 1, i.e., $P(S_9|S_8S_7)=1$ O_0 is ϵ -transition

Adjustments

- Transition probability table will have tuples on rows and states on columns
- Output probability table will remain the same
- In the Viterbi tree, the Markov process will take effect from the 3rd input symbol (εRR)
- There will be 27 leaves, out of which only 9 will remain
- Sequences ending in same tuples will be compared
 - Instead of U1, U2 and U3
 - U₁U₁, U₁U₂, U₁U₃, U₂U₁, U₂U₂, U₂U₃, U₃U₁, U₃U₂, U₃U₃

Forward and Backward Probability Calculation

Forward probability *F(k,i)*

- Define F(k,i)= Probability of being in state S_i having seen $o_0o_1o_2...o_k$
- $F(k,i)=P(o_0o_1o_2...o_k, S_i)$
- With m as the length of the observed sequence
- $P(observed\ sequence) = P(o_0o_1o_2...o_m)$ $= \Sigma_{p=0,N} P(o_0o_1o_2...o_m, S_p)$ $= \Sigma_{p=0,N} F(m, p)$

Forward probability (contd.)

$$F(k, q)$$
= $P(o_0o_1o_2..o_k, S_q)$
= $P(o_0o_1o_2..o_k, S_q)$
= $P(o_0o_1o_2..o_{k-1}, o_k, S_q)$
= $\Sigma_{p=0,N} P(o_0o_1o_2..o_{k-1}, S_p, o_k, S_q)$
= $\Sigma_{p=0,N} P(o_0o_1o_2..o_{k-1}, S_p)$.
 $P(o_m, S_q/o_0o_1o_2..o_{k-1}, S_p)$.
= $\Sigma_{p=0,N} F(k-1,p). P(o_k, S_q/S_p)$

Backward probability B(k,i)

- Define B(k,i)= Probability of seeing $o_k o_{k+1} o_{k+2} ... o_m$ given that the state was S_i
- $B(k,i) = P(o_k o_{k+1} o_{k+2} ... o_m \mid S_i)$
- With m as the length of the observed sequence
- $P(observed\ sequence) = P(o_0o_1o_2...o_m)$ = $P(o_0o_1o_2...o_m | S_0)$ = B(0,0)

Backward probability (contd.)

$$\begin{split} B(k, p) &= P(o_k o_{k+1} o_{k+2} ... o_m \mid S_p) \\ &= P(o_{k+1} o_{k+2} ... o_m, o_k \mid S_p) \\ &= \Sigma_{q=0,N} P(o_{k+1} o_{k+2} ... o_m, o_k, S_q \mid S_p) \\ &= \Sigma_{q=0,N} P(o_k, S_q \mid S_p) \\ &= P(o_{k+1} o_{k+2} ... o_m \mid S_q, S_p) \\ &= \Sigma_{q=0,N} P(o_{k+1} o_{k+2} ... o_m \mid S_q, S_p) \\ &= \Sigma_{q=0,N} P(o_{k+1} o_{k+2} ... o_m \mid S_q). P(o_k, S_q \mid S_p) \\ &= \Sigma_{q=0,N} P(o_k, S_q \mid S_p) \\ &= \Sigma_{q=0,N} P(o_k, S_q \mid S_q). \end{split}$$