# 9: Viterbi Algorithm for HMM Decoding Machine Learning and Real-world Data

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#### Last session: estimating parameters of an HMM

- The dishonest casino, dice edition.
- Two hidden states: L (loaded dice), F (fair dice).
- You don't know which dice is currently in use. You can only observe the numbers that are thrown.
- You estimated transition and emission probabilities (Problem 1 from last time).
- We are now turning to Problem 4.
- We want the HMM to find out when the fair dice was out, and when the loaded dice was out.
- We need to write a decoder.

#### Decoding: finding the most likely path

■ Definition of decoding: Finding the most likely hidden state sequence X that explains the observation O given the HMM parameters  $\mu$ .

$$\hat{X} = \underset{X}{\operatorname{argmax}} P(X, O|\mu)$$

$$= \underset{X}{\operatorname{argmax}} P(O|X, \mu) P(X|\mu)$$

$$= \underset{X_1...X_T}{\operatorname{argmax}} \prod_{t=1}^T P(O_t|X_t) P(X_t|X_{t-1})$$

■ Search space of possible state sequences X is  $O(N^T)$ ; too large for brute force search.

#### Viterbi is a Dynamic Programming Application

(Reminder from Algorithms course) We can use Dynamic Programming if two conditions apply:

- Optimal substructure property
  - An optimal state sequence  $X_1 ... X_j ... X_T$  contains inside it the sequence  $X_1 ... X_j$ , which is also optimal
- Overlapping subsolutions property
  - If both  $X_t$  and  $X_u$  are on the optimal path, with u > t, then the calculation of the probability for being in state  $X_t$  is part of each of the many calculations for being in state  $X_u$ .

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#### The intuition behind Viterbi

- Here's how we can save ourselves a lot of time.
- Because of the Limited Horizon of the HMM, we don't need to keep a complete record of how we arrived at a certain state.
- For the first-order HMM, we only need to record one previous step.
- Just do the calculation of the probability of reaching each state once for each time step.
- Then memoise this probability in a Dynamic Programming table
- This reduces our effort to  $O(N^2T)$ .
- This is for the first order HMM, which only has a memory of one previous state.

#### Viterbi: main data structure

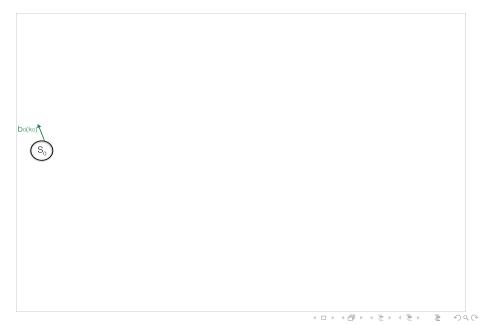
- Memoisation is done using a *trellis*.
- A trellis is equivalent to a Dynamic Programming table.
- The trellis is  $(N+2) \times (T+2)$  in size, with states j as rows and time steps t as columns.
- Each cell j, t records the Viterbi probability  $\delta_j(t)$ , the probability of the most likely path that ends in state  $s_j$  at time t:

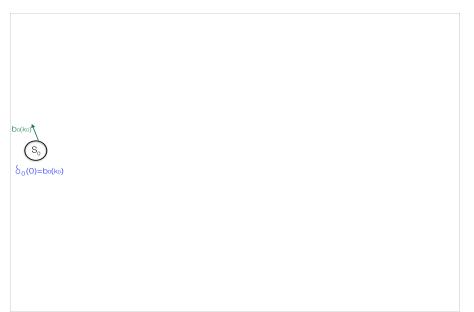
$$\delta_j(t) = \max_{1 \le i \le N} [\delta_i(t-1) a_{ij} b_j(O_t)]$$

- This probability is calculated by maximising over the best ways of going to  $s_i$  for each  $s_i$ .
- $a_{ij}$ : the transition probability from  $s_i$  to  $s_j$
- $b_j(O_t)$ : the probability of emitting  $O_t$  from destination state  $s_j$

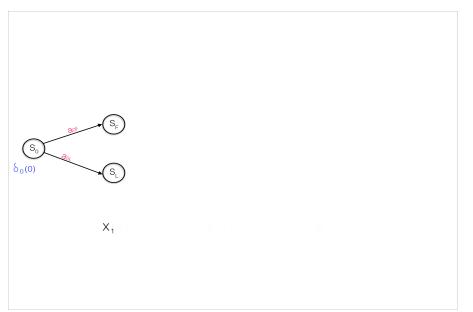
Note: the probability of a state starting the sequence at t=0 is just the probability of it emitting the first symbol.

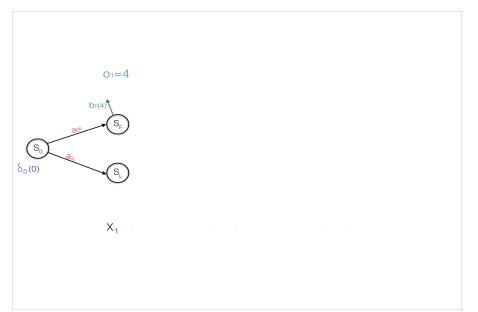


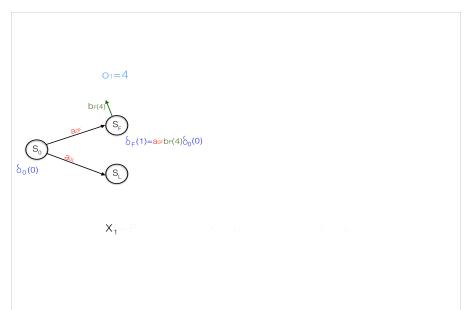




#### Viterbi algorithm, main step





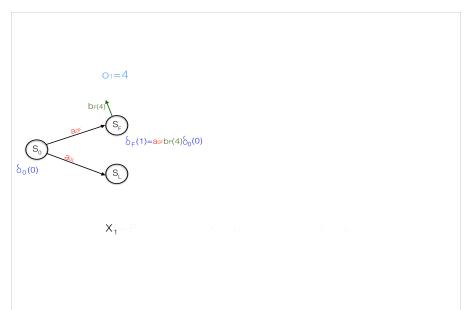


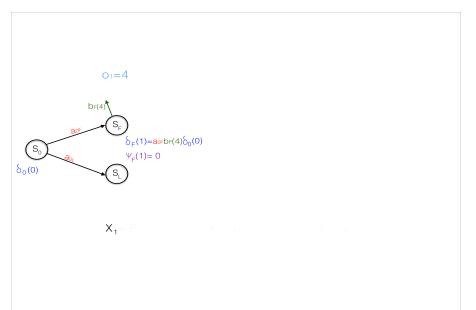
#### Viterbi algorithm, main step, $\psi$

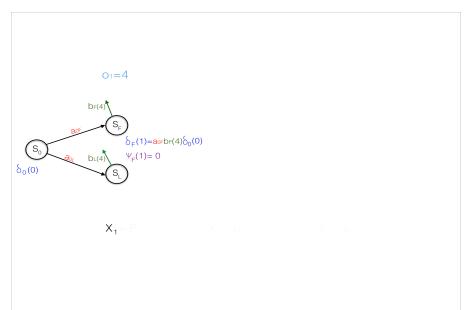
 $\psi_j(t)$  is a helper variable that stores the t-1 state index i on the highest probability path.

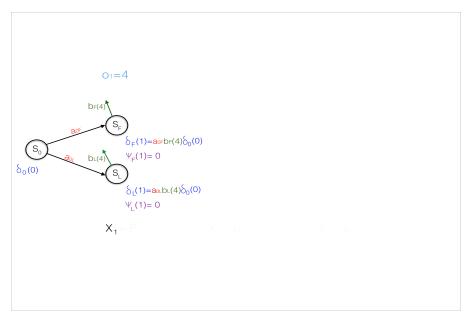
$$\psi_j(t) = \underset{1 < i < N}{\operatorname{argmax}} [\delta_i(t-1) \, a_{ij} \, b_j(O_t)]$$

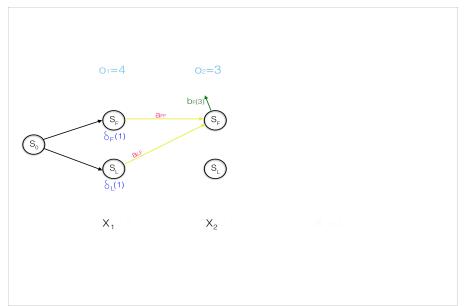
 $\blacksquare$  In the backtracing phase, we will use  $\psi$  to find the previous cell/state in the best path.

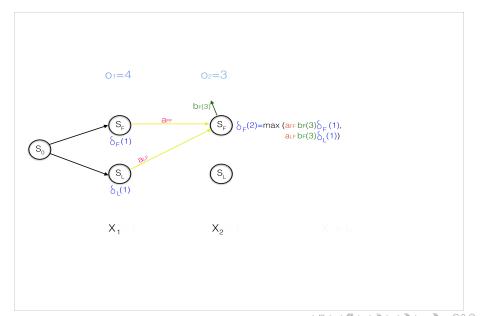


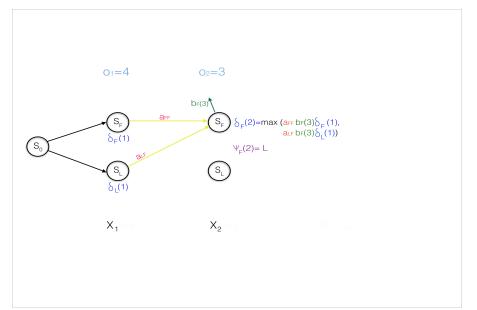


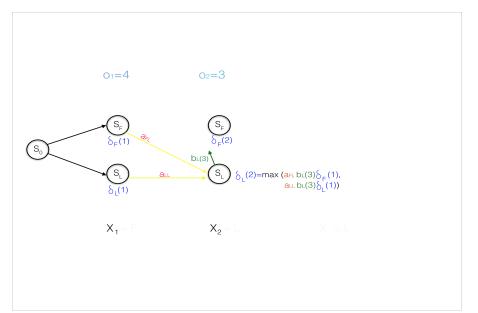


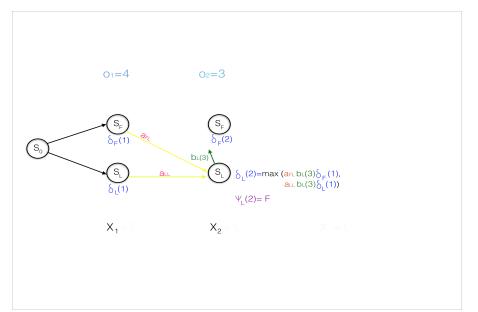


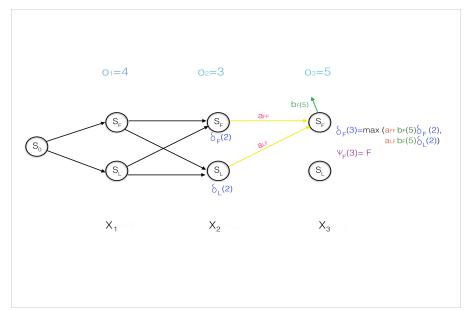


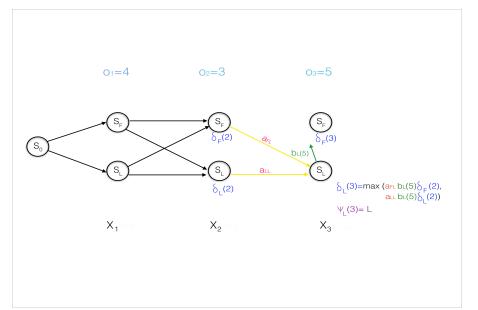




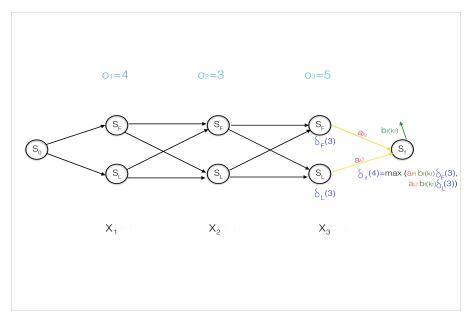




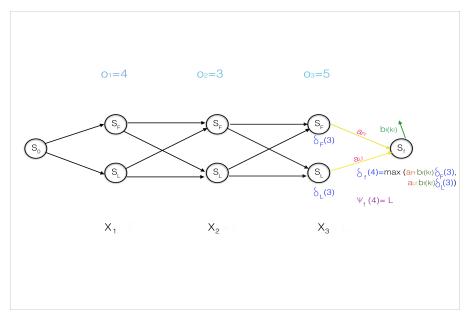


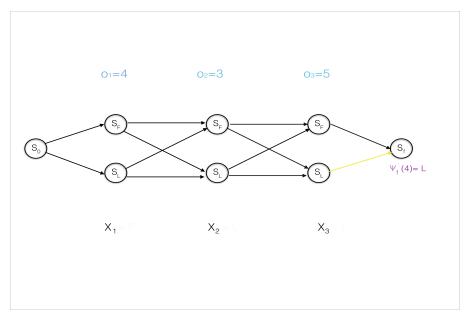


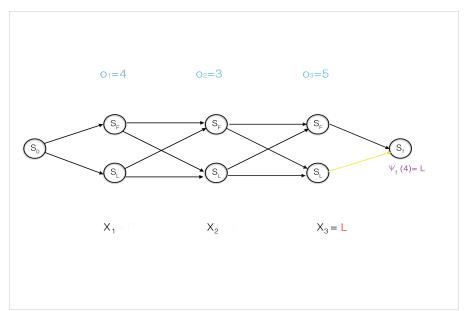
#### Viterbi algorithm, termination

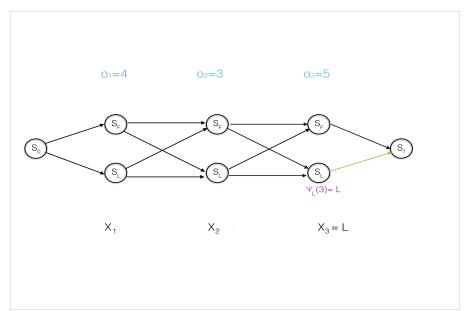


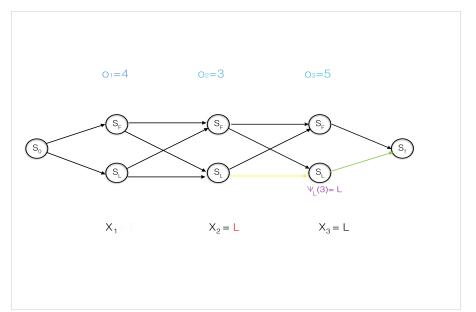
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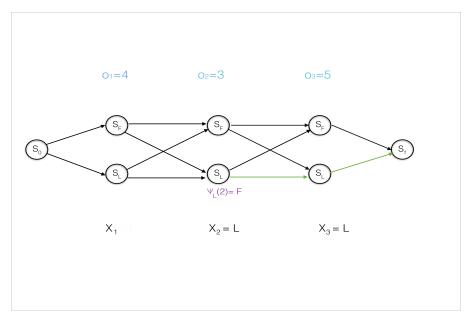


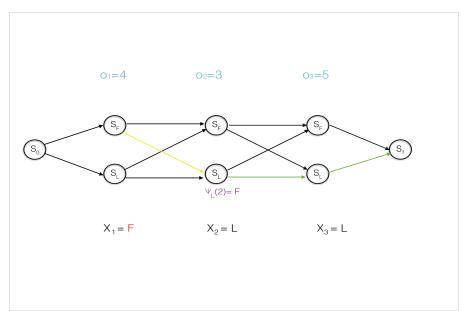


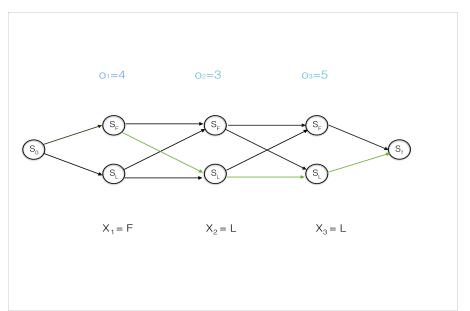


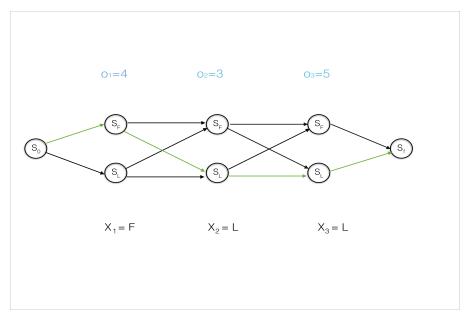












#### Why is it necessary to keep N states at each time step?

- We have convinced ourselves that it's not necessary to keep more than N ("real") states per time step.
- But could we cut down the table to just a one-dimensional table of *T* time slots by choosing the probability of the best path overall ending in that time slot, in any of the states?
  - This would be the greedy choice
  - But think about what could happen in a later time slot.
  - You could encounter a zero or very low probability concerning all paths going through your chosen state  $s_j$  at time t.
  - Now a state  $s_k$  that looked suboptimal in comparison to  $s_j$  at time t becomes the best candidate.
  - As we don't know the future, this could happen to any state, so we need to keep the probabilities for each state at each time slot.
- But thankfully, no more.

#### Precision and Recall

- So far, we have measured system success in accuracy or agreement in Kappa.
- But sometimes it's only one type of instances that we find interesting.
- We don't want a summary measure that averages over interesting and non-interesting instances, as accuracy does.
- In those cases, we use precision, recall and F-measure.
- These metrics are imported from the field of information retrieval, where the difference between interesting and non-interesting examples is particularly high.
- Accuracy doesn't work well when the types of instances are unbalanced

#### Precision and Recall

#### System says:

Truth is:

|       | F   | L   | Total   |
|-------|-----|-----|---------|
| F     | а   | b   | a+b     |
| L     | С   | d   | c+d     |
| Total | a+c | b+d | a+b+c+d |

- Precision of L:  $P_L = \frac{d}{b+d}$
- Recall of L:  $R_L = \frac{d}{c+d}$
- F-measure of L:  $F_L = \frac{2P_LR_L}{P_L + R_L}$
- Accuracy:  $A = \frac{a+d}{a+b+c+d}$

#### Your task today

#### Task 8:

- Implement the Viterbi algorithm.
- Run it on the dice dataset and measure precision of L  $(P_L)$ , recall of L  $(R_L)$  and F-measure of L  $(F_L)$ .

#### Literature

- Manning and Schutze (2000). Foundations of Statistical Natural Language Processing, MIT Press. Chapter 9.3.2.
  - We use a state-emission HMM, but this textbook uses an arc-emission HMM. There is therefore a slight difference in the algorithm as to in which step the initial and final  $b_j(k_t)$  are multiplied in.
- Jurafsky and Martin, 2nd Edition, chapter 6.4
- Smith, Noah A. (2004). Hidden Markov Models: All the Glorious Gory Details.
- Bockmayr and Reinert (2011). Markov chains and Hidden Markov Models. Discrete Math for Bioinformatics WS 10/11.