# CMPT 413 Computational Linguistics

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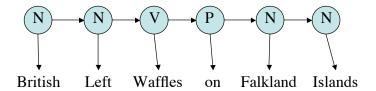
# Sequence Learning

- British Left Waffles on Falkland Islands
  - -(N, N, V, P, N, N)
  - -(N, V, N, P, N, N)
- Segmentation 中国十四个边境开放城市经济建设成就显著
  - -(b, i, b, i, b, b, i, b, i, b, i, b, i, b, i, b, i, b, i)

中国 十四 个 边境 开放 城市 经济 建设 成就 显著

China 's 14 open border cities marked economic achievements

# Sequence Learning



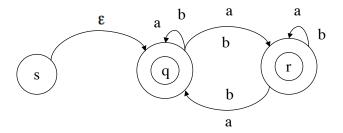
3 states: N, V, P

**Observation sequence**:  $(o_1, \dots o_6)$ 

**State sequence** (6+1): (*Start*, *N*, *N*, *V*, *P*, *N*, *N*)

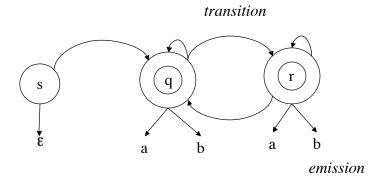
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# Finite State Machines



Mealy Machine

#### Finite State Machines



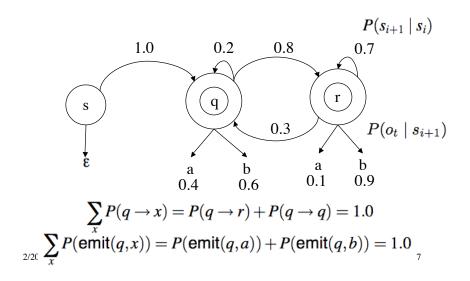
Moore Machine

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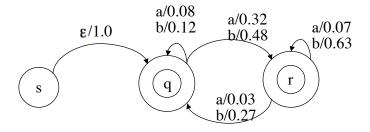
# Probabilistic FSMs

- Each transition is associated with a *transition probability*
- Each emission is associated with an *emission probability*
- Two conditions:
  - All outgoing transition arcs from a state must sum to 1
  - All emission arcs from a state must sum to 1

# Probabilistic FSMs



#### Probabilistic FSMs



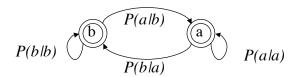
#### Hidden Markov Models

- There are n states  $s_1, ..., s_i, ..., s_n$
- The emissions are observed (input data)
- Observation sequence  $\mathbf{O} = (o_1, ..., o_t, ..., o_T)$
- The states are not directly observed (hidden)
- Data does not directly tell us which state  $X_t$  is linked with observation  $o_t$

$$X_t \in \{s_1,\ldots,s_n\}$$

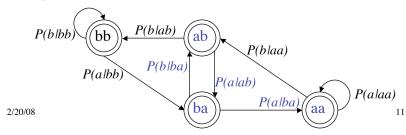
#### Markov Chains vs. HMMs

- For observation sequence babaa i.e:  $o_1$ =b,  $o_2$ =a, ...,  $o_5$ =a
- Compute P(babaa) using a bigram model P(b)\*P(a|b)\*P(b|a)\*P(a|b)\*P(a|a)
- Equivalent Markov chain:



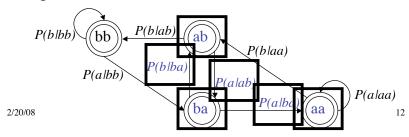
#### Markov Chains vs. HMMs

- For observation sequence *babaa* i.e:  $o_1$ =b,  $o_2$ =a, ...,  $o_5$ =a
- Compute P(babaa) using a trigram model P(ba)\*P(b|ba)\*P(a|ab)\*P(a|ba)
- Equivalent Markov chain:



#### Markov Chains vs. HMMs

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- Equivalent Markov chain:



#### Markov Chains vs. HMMs

• Given an observation sequence

$$\mathbf{O} = (o_1, ..., o_t, ..., o_T)$$

• An *n*th order Markov Chain or *n*-gram model computes the probability

$$P(o_1, ..., o_t, ..., o_T)$$

• An HMM computes the probability  $P(X_1, ..., X_{T+1}, o_1, ..., o_T)$  where the state sequence is *hidden* 

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# Properties of HMMs

• Markov assumption

$$P(X_t = s_i \mid \dots, X_{t-1} = s_j)$$

• Stationary distribution

$$P(X_t = s_i | X_{t-1} = s_i) = P(X_{t+1} = s_i | X_{t+l-1} = s_i)$$

#### **HMM** Algorithms

- HMM as language model: compute probability of given observation sequence
- HMM as parser: compute the best sequence of states for a given observation sequence
- HMM as learner: given a set of observation sequences, learn its distribution, i.e. learn the transition and emission probabilities

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# **HMM** Algorithms

- HMM as language model: compute probability of given observation sequence
- Compute  $P(o_1, ..., o_T)$  from the probability

$$P(X_{I}, ..., X_{T+I}, o_{I}, ..., o_{T})$$

$$= \prod_{t=1}^{T} P(X_{t+1} = s_{j} \mid X_{t} = s_{i}) \times P(o_{t} = k \mid X_{t+1} = s_{j})$$

$$P(o_1, ..., o_T) = \sum_{X_1,...,X_{T+1}} P(X_1,...,X_{T+1},o_1,...,o_T)$$

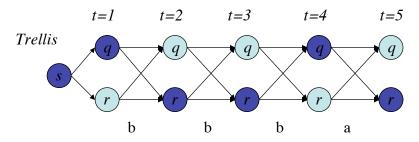
#### **HMM** Algorithms

- HMM as parser: compute the best sequence of states for a given observation sequence
- Compute best path  $X_1, ..., X_{T+1}$  from the probability  $P(X_1, ..., X_{T+1}, o_1, ..., o_T)$ Best state sequence  $X_1^*, ..., X_{T+1}^*$

$$= rgmax_{X_1,\ldots,X_{T+1}} P(X_1,\ldots,X_{T+1},o_1,\ldots,o_T)$$

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# Best Path (Viterbi) Algorithm



- Key Idea 1: storing just the best path doesn't work
- Key Idea 2: store the best path upto each state

#### Viterbi Algorithm

```
function viterbi (edges, input, obs): returns best path
edges = transition probability
input = emission probability
T = length of obs, the observation sequence
num-states = number of states in the HMM
Create a path-matrix: viterbi[num-states+1, T+1] # init to all 0s
for each state s: viterbi[s, 0] = \pi[s]
for each time step t from 0 to T:
  for each state s from 0 to num-states:
     for each s' where edges[s,s'] is a transition probability:
       new-score = viterbi[s,t] * edges[s,s'] * input[s',obs[t]]
       if (viterbi[s',t+1] == 0) or (new-score > viterbi[s', t+1]):
          viterbi[s', t+1] = new-score
          back-pointer[s',t+1] = s
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                                                                           19
```

#### Viterbi Algorithm

#### # finding the best path

```
best-final-score = best-final-state = 0
for each state s from 0 to num-states:
    if (viterbi[s,T+1] > best-final-score):
        best-final-state = s
        best-final-score = viterbi[s,T+1]
# start with the last state in the sequence
x = best-final-state
state-sequence.push(x)
for t from T+1 downto 0:
    state-sequence.push(back-pointer[x,t])
    x = back-pointer[x,t]
return state-sequence
```

#### Forward-Backward Algorithm

- Algorithm that finds the transition and emission probabilities using training data that *does not have* hidden states provided
- Set the probabilities (for all parameters in the HMM) so that the training data T is assigned highest P(T) value (or lowest H(T), entropy value)
- This is called the maximum likelihood value over all possible hidden state sequences for the training data
- Exploits the fact that some transitions and resulting observations will occur more frequently 2/20/than others in the training data

#### Forward-backward Algorithm

- Consider input  $o_1,..., o_t,..., o_T$  where each  $o_t$  is from a set of symbols  $V = \{1,..k,..K\}$
- Let  $\pi_i$  be the probability of state *i* being a start state (for simplicity,  $\pi_i$  is not discussed further)
- Let  $a_{i,j}$  be the transition probability:  $P(X_{t+1} = s_i \mid X_t = s_i) \quad |S|^2 \text{ distinct } a_{i,j} \text{ values}$
- Let  $b_{j,k}$  be the emission probability:  $P(o_t = k \mid X_{t+1} = s_j) \quad |S| \times |V| \text{ distinct } b_{j,k} \text{ values}$
- Probability of going from state  $s_i$  to state  $s_j$  while observing input  $o_t$  is simply  $a_{i,j} \times b_{j,k}$

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#### Forward-backward Algorithm

- The algorithm starts with an initial setting for the probabilities in *a* and *b*
- We are provided with training data which consists of observation sequence(s):  $o_1,...,o_t,...,o_T$
- The probability  $P(o_1,...,o_T)$  depends on the values in a and b
- For given observation sequence(s), different transitions/emissions will be visited with different frequencies

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# Forward-backward Algorithm

- For every path through the HMM, we count how many transitions occurred from state i to state j on observation o<sub>t</sub>
- Then (loosely speaking) we reward those transitions (and emissions) which have high *expected* frequency and penalize the competing transitions
- Expected frequency means we multiply the frequency with the current probability (taken from *a* and *b*)

#### Forward-backward Algorithm

- $P(o_1,...,o_T)$  is the expected frequency of visiting all transitions and so the new frequency is the expected occurrence of a transition divided by  $P(o_1,...,o_T)$
- This gives us new values for all probabilities: a' and b' and we set a and b to these new values
- Compute  $P(o_1,...,o_T)$ . If the value is unchanged from before iteration then stop (convergence)
- Otherwise iterate (the entire procedure) with new values for *a* and *b*

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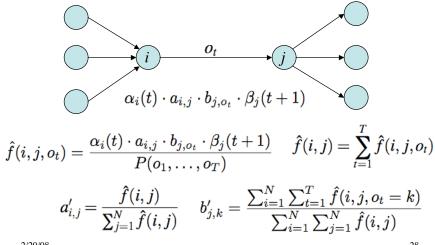
# Forward-backward Algorithm

- How to compute expected frequency over all paths efficiently (reuse dynamic programming idea from Viterbi algorithm)
- For input  $o_1,..., o_r,..., o_T$  where  $o_t \in V = \{1,..k,..K\}$
- For every path from a start state to state i we can compute the probability of observing  $o_1, ..., o_{t-1}$
- Let  $\alpha_i(t)$  be the sum of all these probabilities
- For every path from state j to a final state we can compute the probability of observing  $o_{t+1},...,o_T$
- Let  $\beta_j(t+1)$  be the sum of all these probabilities

## Forward-Backward Algorithm

$$lpha_k(t-1)$$
  $k$   $m$   $eta_m(t+2)$   $a_i(t) \cdot a_{i,j} \cdot b_{j,o_t} \cdot eta_j(t+1)$   $a_i(t) = \sum_{k=1}^N a_{k,i} \cdot b_{i,o_{t-1}} \cdot lpha_k(t-1)$   $\beta_j(t+1) = \sum_{m=1}^N a_{j,m} \cdot b_{m,o_{t+1}} \cdot eta_m(t+2)$   $a_{j,m} \cdot b_{m,o_{t+1}} \cdot \beta_m(t+2)$   $a_{j,m} \cdot b_{m,o_{t+1}} \cdot \beta_m(t+2)$ 

# Forward-Backward Algorithm



# Forward-Backward Algorithm

- Each iteration provides new values for all the *parameters*
- But are the new parameters any better? How can we tell?
- Compute probability of the training data
- For HMMs, Baum 1977 shows that the probability will always be non-decreasing (later generalized to the more general EM algorithm)
- Same as cross-entropy is non-increasing

$$KL(\mu_{i+1} \mid\mid D) \leq KL(\mu_i \mid\mid D)$$

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