Hidden Markov models

CSC401/2511 – Natural Language Computing – Winter 2021 Lecture 7 Serena Jeblee, Sean Robertson and Frank Rudzicz University of Toronto

• We've seen this type of model:

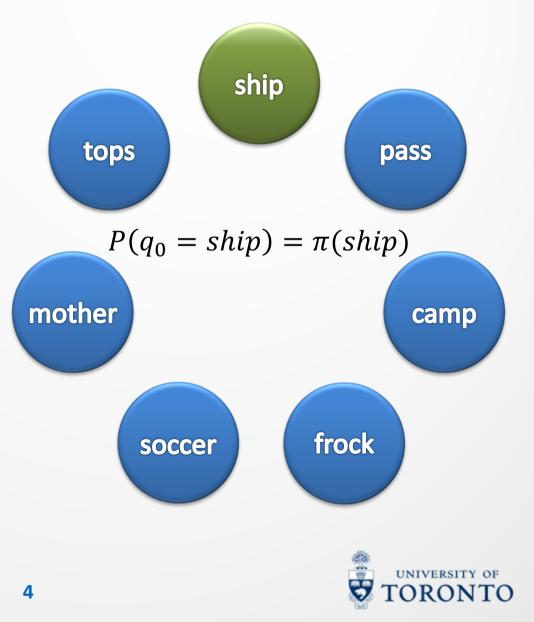
- e.g., consider the 7-word vocabulary: {ship, pass, camp, frock, soccer, mother, tops}
- What is the probability of the **sequence** *ship*, *ship*, *pass*, *ship*, *tops* ?
- Assuming a bigram model (i.e., 1st-order Markov), *P(ship|<s>)P(ship|ship)P(pass|ship)* · *P(ship|pass)P(tops|ship)*



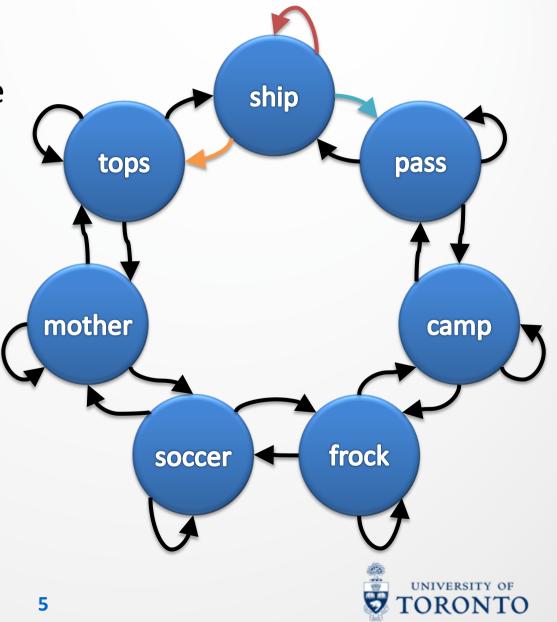
- This can be conceptualized graphically.
- We start with N states, s₁, s₂, ..., s_N that represent unique observations in the world.
- Here, N = 7 and each state represents one of the words we can observe.



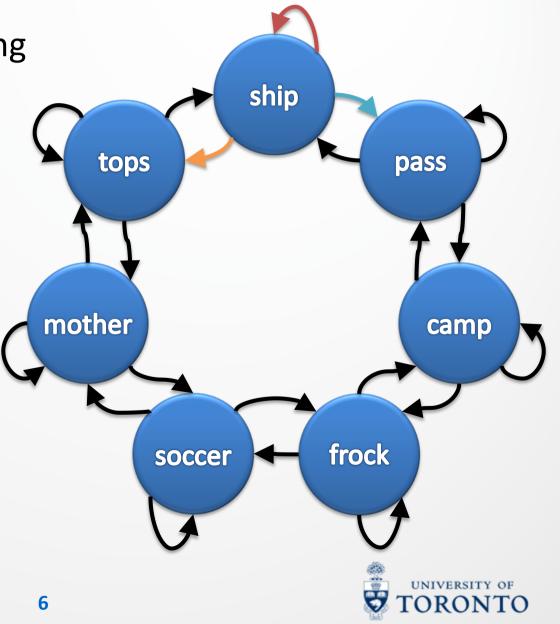
- We have discrete
 timesteps, t = 0, t = 1, ...
- On the tth timestep the system is in exactly one of the available states, q_t.
 - $q_t \in \{s_1, s_2, \dots, s_N\}$
- We could start in any state. The probability of starting with a particular state s is $P(q_0 = s) = \pi(s)$



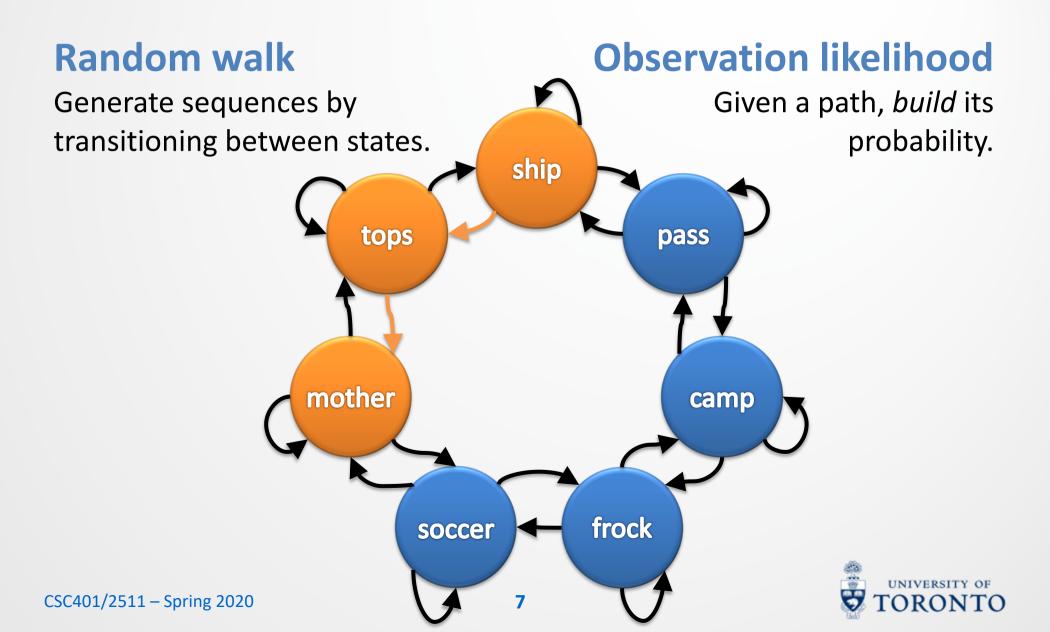
- At each step we must move to a state with some probability.
- Here, an arrow from q_t to q_{t+1} represents $P(q_{t+1}|q_t)$
- P(ship|ship)
- P(tops|ship)
- P(pass|ship)
- P(frock|ship) = 0



- Probabilities on all outgoing arcs must sum to 1.
- P(ship|ship) + P(tops|ship) + P(tops|ship) + P(pass|ship) = 1
- P(ship|tops) +P(tops|tops) +P(mother|tops) = 1



Using the graph



A multivariate system

 What if the probabilities of observing words depended only on some other variable, like mood?

| P(word) | | | |
|---------|--|--|--|
| 0.1 | | | |
| 0.05 | | | |
| 0.05 | | | |
| 0.6 | | | |
| 0.05 | | | |
| 0.1 | | | |
| 0.05 | | | |
| | | | |

| P(word) | | | | |
|---------|--|--|--|--|
| 0.25 | | | | |
| 0.25 | | | | |
| 0.05 | | | | |
| 0.3 | | | | |
| 0.05 | | | | |
| 0.09 | | | | |
| 0.01 | | | | |
| | | | | |

| word | P(word) | | | |
|-------|---------|--|--|--|
| ship | 0.3 | | | |
| pass | 0 | | | |
| camp | 0 | | | |
| frock | 0.2 | | | |
| | | | | |

soccer

mother

tops



0.05

0.05

0.4

A multivariate system

• What if that variable changes over time?

- e.g., I'm happy one second and disgusted the next.
- Here, $state \equiv mood$ observation $\equiv word$.

| > | word | P(word) |
|---|--------|---------|
| | ship | 0.1 |
| | pass | 0.05 |
| | camp | 0.05 |
| | frock | 0.6 |
| | soccer | 0.05 |
| | mother | 0.1 |
| | tops | 0.05 |

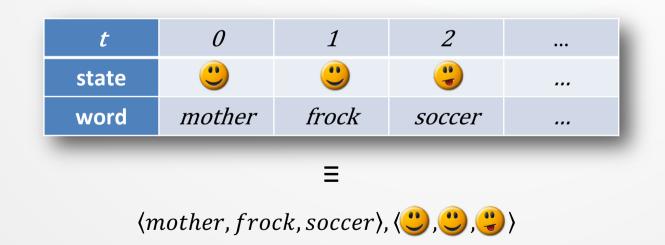
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| frock | 0.2 |
| soccer | 0.05 |
| mother | 0.05 |
| tops | 0.4 |
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Observable multivariate systems

- Imagine you have access to my emotional state somehow.
- All your data are completely observable at every timestep.
- E.g.,

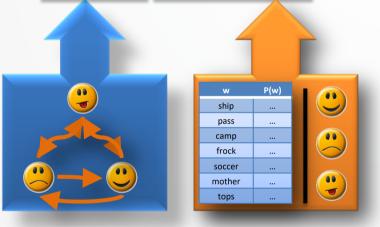




Observable multivariate systems

• What is the probability of a sequence of words and states?

• $P(w_{0:t}, q_{0:t}) = P(q_{0:t})P(w_{0:t}|q_{0:t}) \approx \prod_{i=0}^{t} P(q_i|q_{i-1})P(w_i|q_i)$



• e.g.,

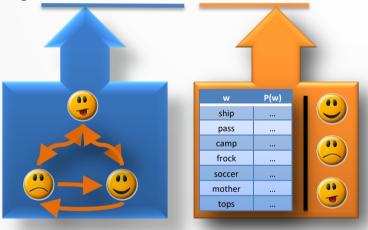
 $P(\langle ship, pass \rangle, \langle \bigcirc, \bigcirc \rangle) = P(q_0 = \bigcirc) P(ship| \bigcirc) P(\bigcirc) P(pass| \bigcirc)$



Observable multivariate systems

• **Q**: How do you **learn** these probabilities?

• $P(w_{0:t}, q_{0:t}) \approx \prod_{i=0}^{t} P(q_i | q_{i-1}) P(w_i | q_i)$



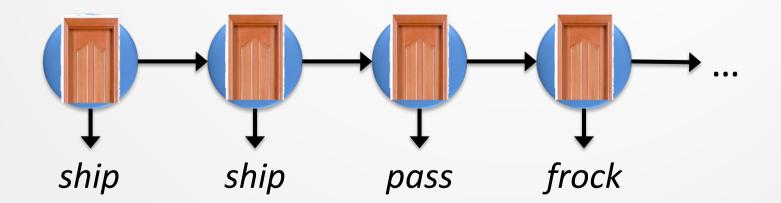
- A: When all data are observed, basically the same as before.
 - $P(q_i|q_{i-1}) = \frac{P(q_{i-1}q_i)}{P(q_{i-1})}$ is learned with MLE from training data. $P(w_i|q_i) = \frac{P(w_i,q_i)}{P(q_i)}$ is also learned with MLE from training data.



Hidden variables

• Q: What if you **don't** know the **states** during *testing*?

- e.g., compute P((ship, ship, pass, frock))
- Q: What if you **don't** know the **states** during *training*?





Examples of hidden phenomena

- We want to represent surface (i.e., observable) phenomena as the output of hidden underlying systems.
 - e.g.,
 - Words are the outputs of hidden parts-of-speech,
 - French phrases are the outputs of hidden English phrases,
 - Speech sounds are the outputs of hidden phonemes.
 - in other fields,
 - Encrypted symbols are the outputs of hidden messages,
 - Genes are the outputs of hidden functional relationships,
 - Weather is the output of hidden climate conditions,
 - Stock prices are the outputs of hidden market conditions,



Definition of an HMM

- A hidden Markov model (HMM) is specified by the 5-tuple $\{S, W, \Pi, A, B\}$:
 - $S = \{S_1, ..., S_N\}$
 - $W = \{w_1, ..., w_k\}$

$$= \{\pi_1, \dots, \pi_N\}$$

• $A = \{a_{ij}\}, i, j \in S$

 $\bullet B = b_i(w), i \in S, w \in W$: state **output** probabilities

- : set of states (e.g., moods)
- : output alphabet (e.g., words)
- : initial state probabilities
- : state transition probabilities

yielding

- $Q = \{q_0, ..., q_{T-1}\}, q_i \in S$: state sequence
- $\mathcal{O} = \{ \sigma_0, \dots, \sigma_{T-1} \}, \sigma_i \in W$: output sequence



A hidden Markov production process

- An HMM is a **representation** of a process in the world.
 - We can synthesize data, as in Shannon's game.
- This is how an HMM generates new sequences:
- $t \coloneqq 0$
- **Start** in state $q_0 = s_i$ with probability π_i
- Emit observation symbol $\sigma_0 = w_k$ with probability $b_i(\sigma_0)$
- While (not forever)
 - **Go** from state $q_t = s_i$ to state $q_{t+1} = s_j$ with probability a_{ij}
 - Emit observation symbol $\sigma_{t+1} = w_k$ with probability $b_j(\sigma_{t+1})$
 - $t \coloneqq t + 1$



Fundamental tasks for HMMs

1. Given a model with particular parameters $\theta = \langle \Pi, A, B \rangle$, how do we efficiently compute the likelihood of a *particular* observation sequence, $P(\mathcal{O}; \theta)$?

We previously computed the probabilities of word sequences using *N*-grams.

The probability of a particular sequence is usually useful as a means to some other end.



Fundamental tasks for HMMs

2. Given an observation sequence O and a model θ , how do we choose a state sequence $Q = \{q_0, \dots, q_{T-1}\}$ that *best explains* the observations?

This is the task of **inference** – i.e., guessing at the best explanation of unknown ('latent') variables given our model.

This is often an important part of **classification**.



Fundamental tasks for HMMs

3. Given a large **observation sequence** O, how do we choose the *best parameters* $\theta = \langle \Pi, A, B \rangle$ that explain the data O?

This is the task of **training**.

As before, we want our parameters to be set so that the available training data is maximally likely, But doing so will involve guessing unseen information.



Task 1: Computing $P(\mathcal{O}; \theta)$

We've seen the probability of a joint sequence of observations and states:

$$P(\mathcal{O}, Q; \theta) = P(\mathcal{O}|Q; \theta) P(Q; \theta)$$

= $\pi_{q_0} b_{q_0}(\sigma_0) a_{q_0 q_1} b_{q_1}(\sigma_1) a_{q_1 q_2} b_{q_2}(\sigma_2) \dots$

• To get the probability of our observations **without** seeing the state, we must **sum over all possible state sequences**:

$$P(\mathcal{O};\theta) = \sum_{Q} P(\mathcal{O},Q;\theta) = \sum_{Q} P(\mathcal{O}|Q;\theta) P(Q;\theta).$$



Computing $P(O; \theta)$ naïvely

 To get the total probability of our observations, we could directly sum over all possible state sequences:

$$P(\mathcal{O};\theta) = \sum_{Q} P(\mathcal{O}|Q;\theta) P(Q;\theta).$$

- For observations of length *T*, each state sequence involves 2*T* multiplications (1 for each state transition, 1 for each observation, 1 for the start state, minus 1).
- There are up to N^T possible state sequences of length T given N states.

 $\therefore \sim (1 + T + T - 1) \cdot N^T$ multiplications \bigcirc



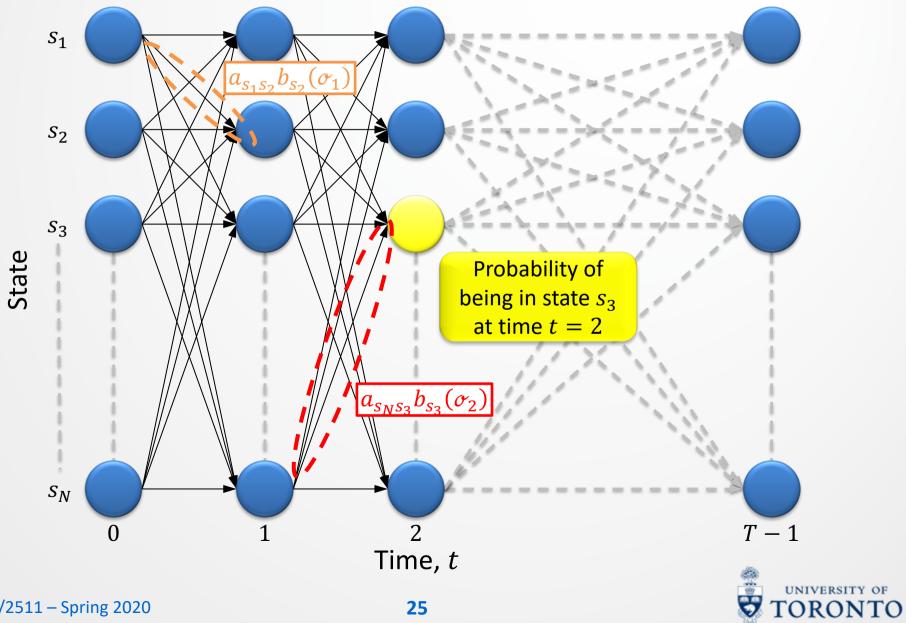
Computing $P(\mathcal{O}; \theta)$ **cleverly**

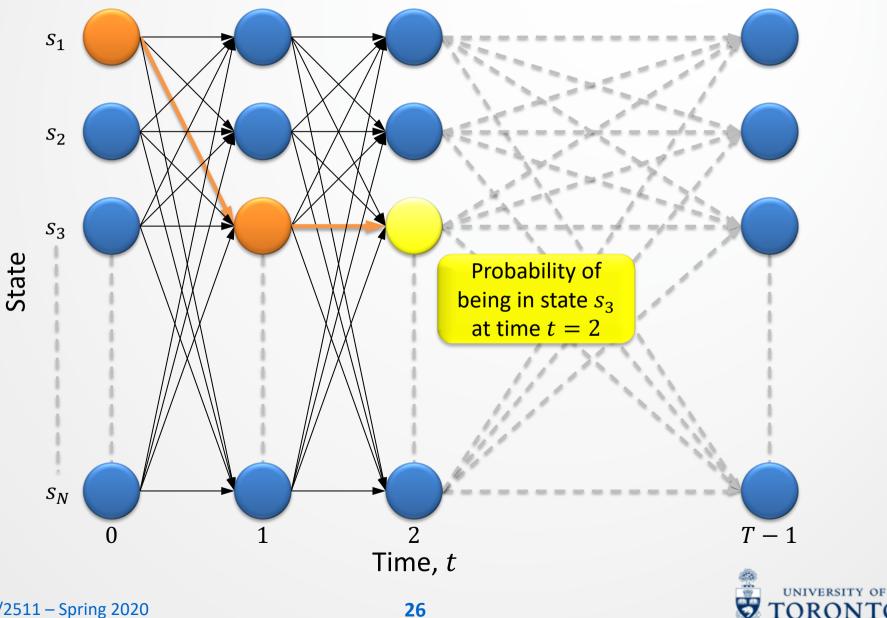
- To avoid this complexity, we use dynamic programming; we remember, rather than recompute, partial results.
- We make a trellis which is an array of states vs. time.
 The element at (i, t) is α_i(t) the probability of being in state i at time t after seeing all observations to that point:

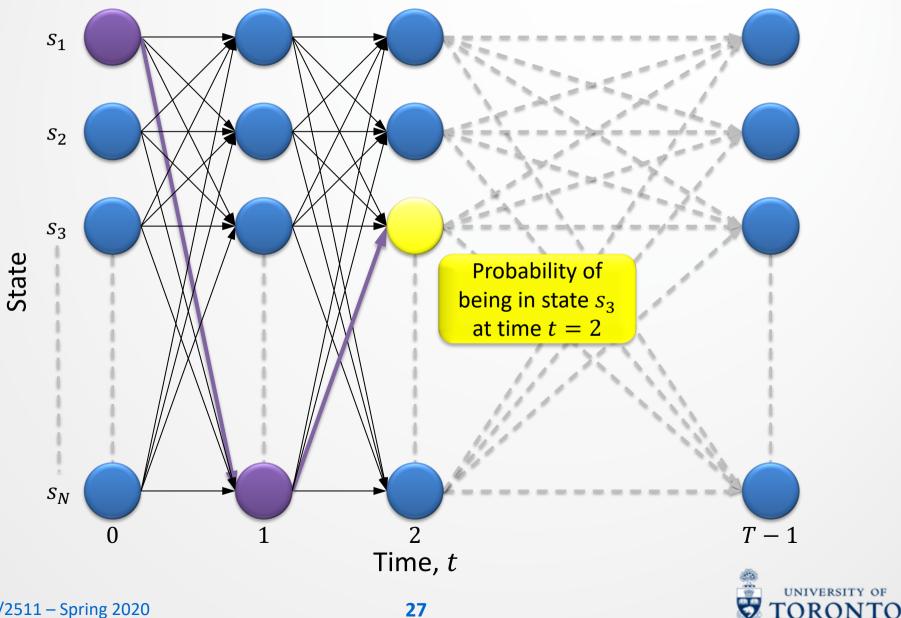
 $P(\sigma_{o:t}, q_t = s_i; \theta)$

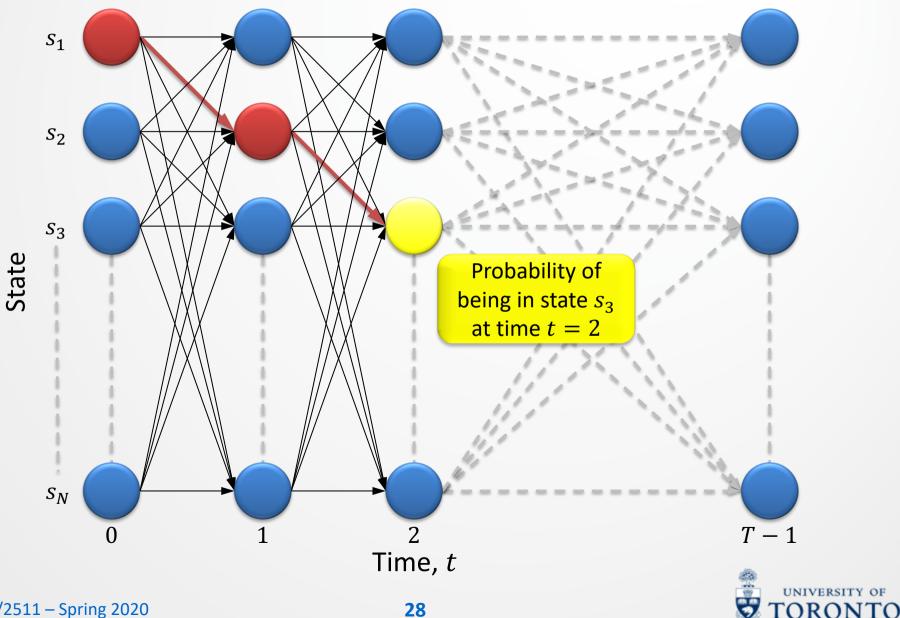


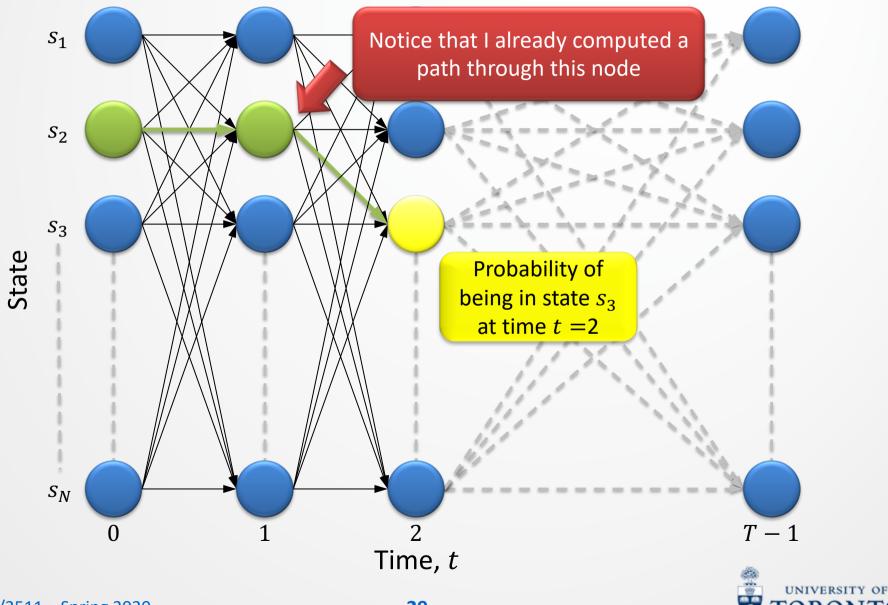
Trellis

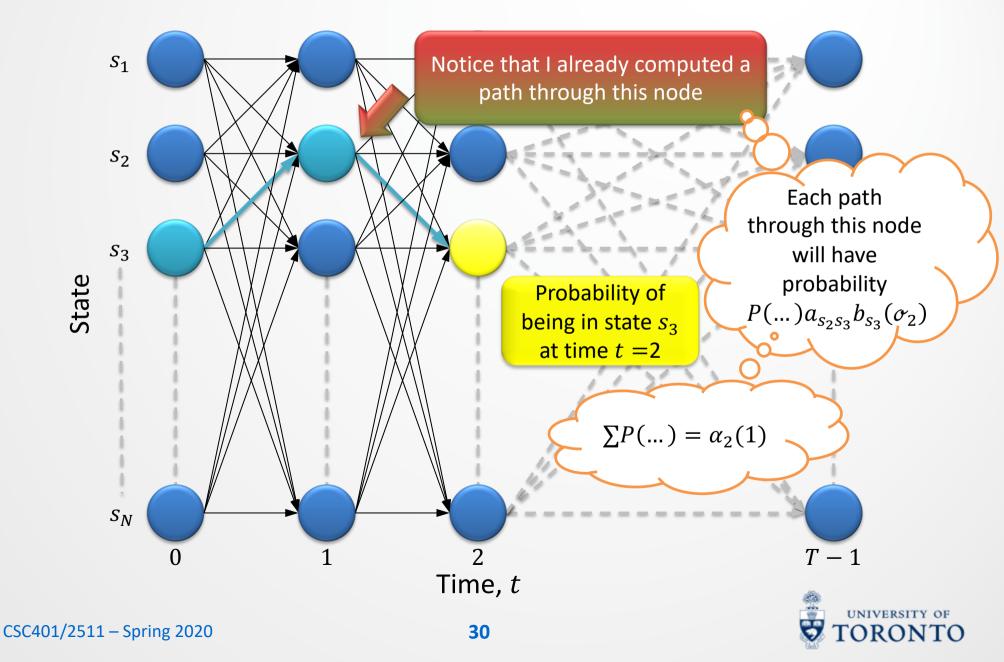








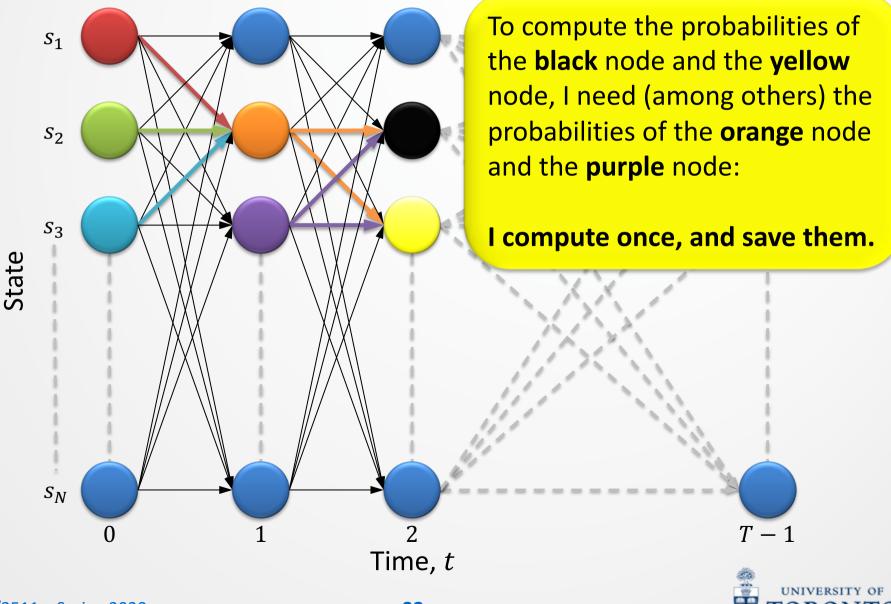




AND SO ON...



Trellis



The Forward procedure

• To compute

$$\alpha_i(t) = P(\sigma_{0:t}, q_t = s_i; \theta)$$

we can compute $\alpha_j(t-1)$ for possible *previous* states s_j , then use our knowledge of a_{ji} and $b_i(\sigma_t)$

 We compute the trellis left-to-right (because of the convention of time) and top-to-bottom ('just because').

• **Remember**: σ_t is fixed and known. $\alpha_i(t)$ is agnostic of the future.

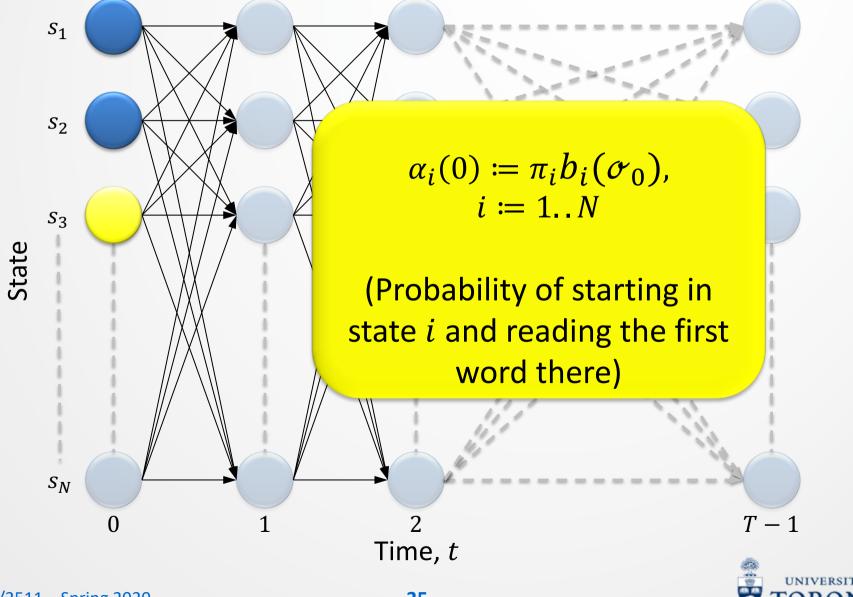


The Forward procedure

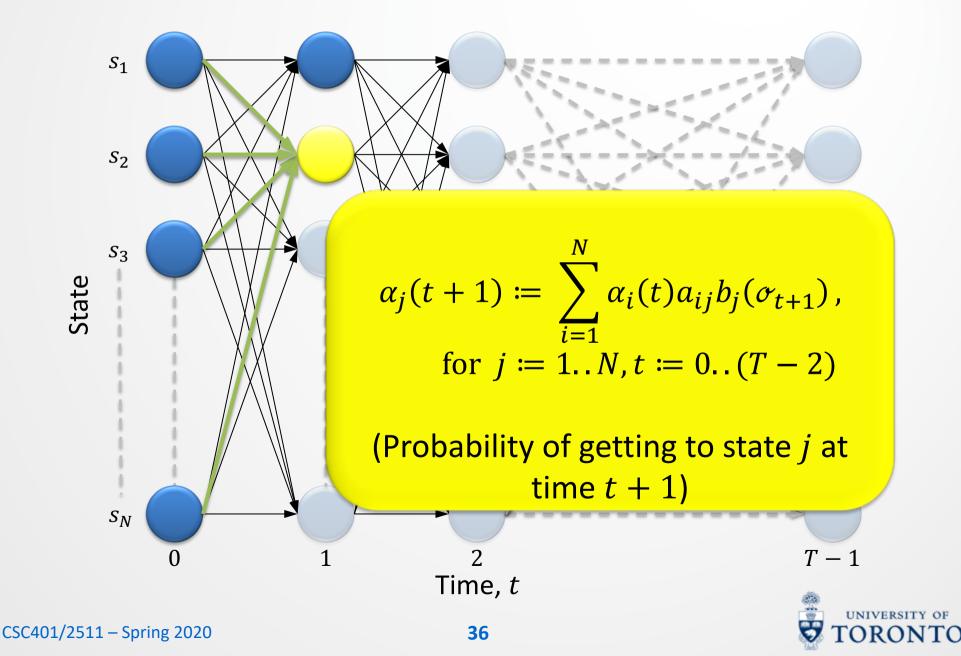
- The trellis is computed left-to-right and top-to-bottom.
- There are three steps in this procedure:
 - Initialization: Compute the nodes in the *first* column of the trellis (t = 0).
 - Induction: Iteratively compute the nodes in the *rest* of the trellis $(1 \le t < T)$.
 - Conclusion: Sum o
- Sum over the nodes in the *last* column of the trellis (t = T 1).



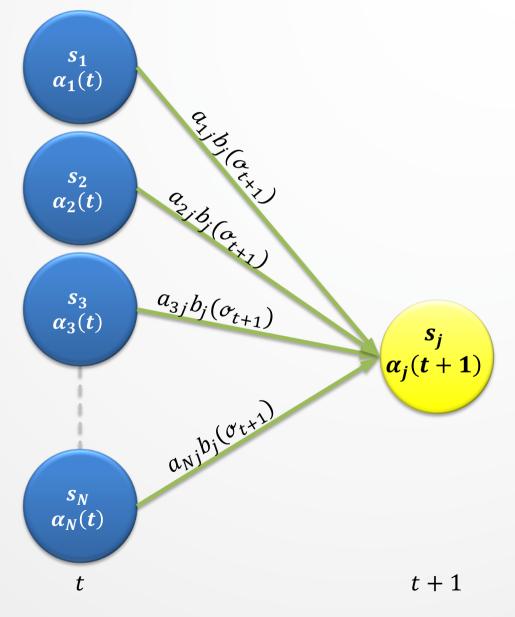
Initialization of Forward procedure



Induction of Forward procedure

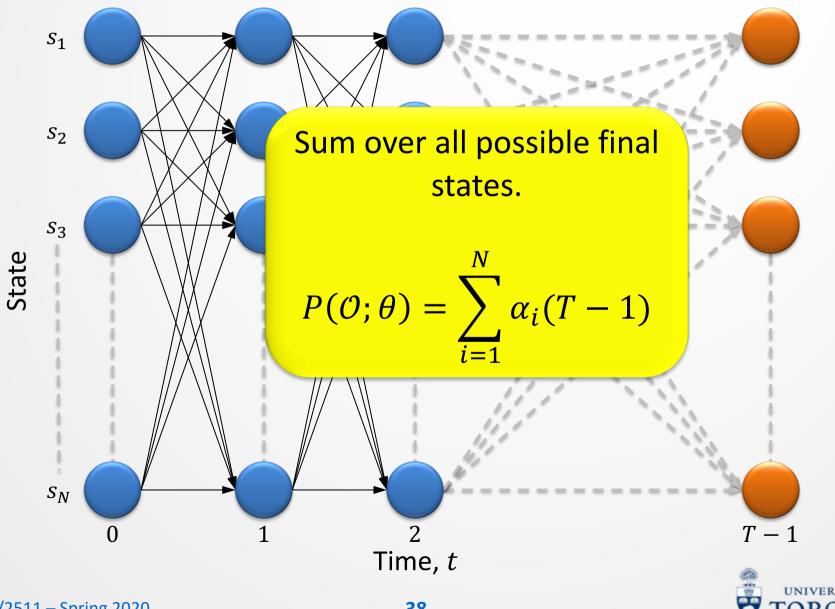


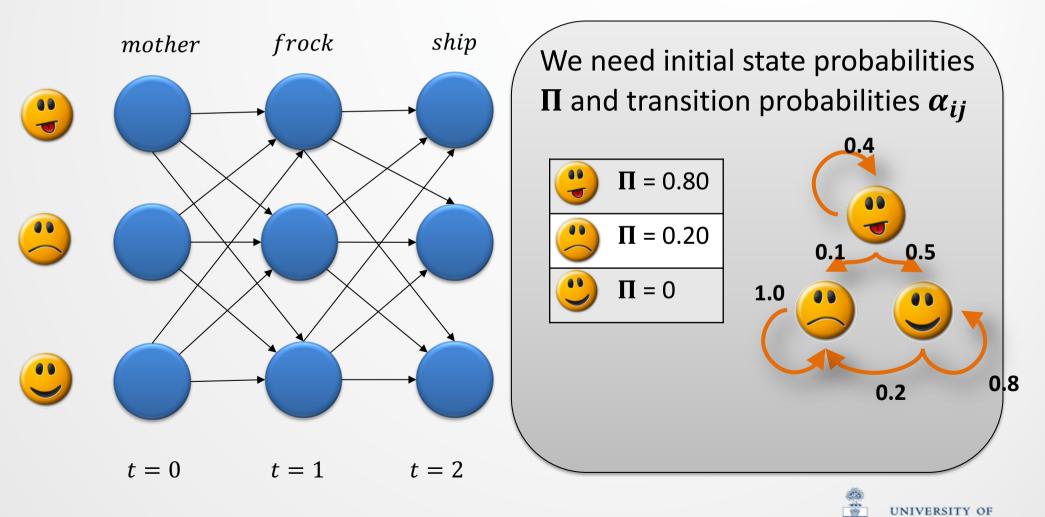
Induction of Forward procedure

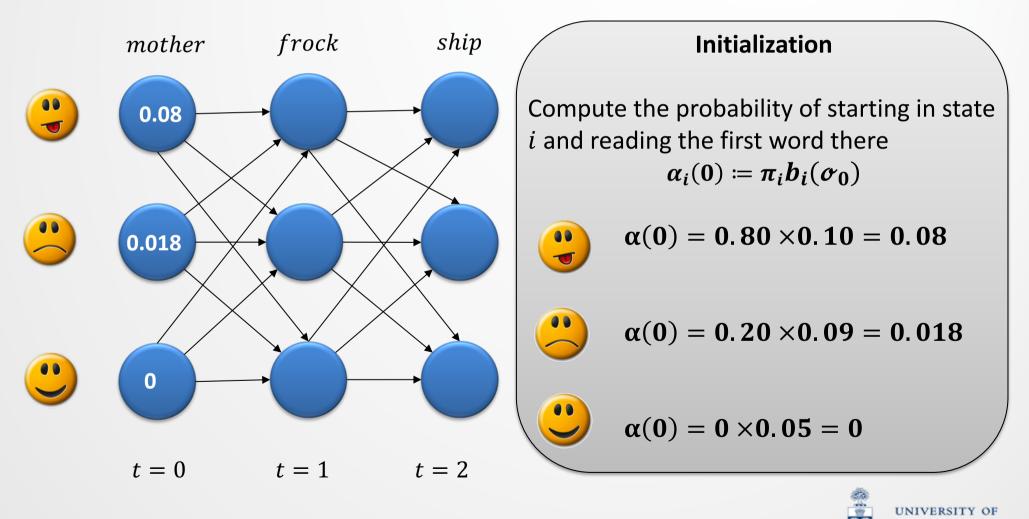


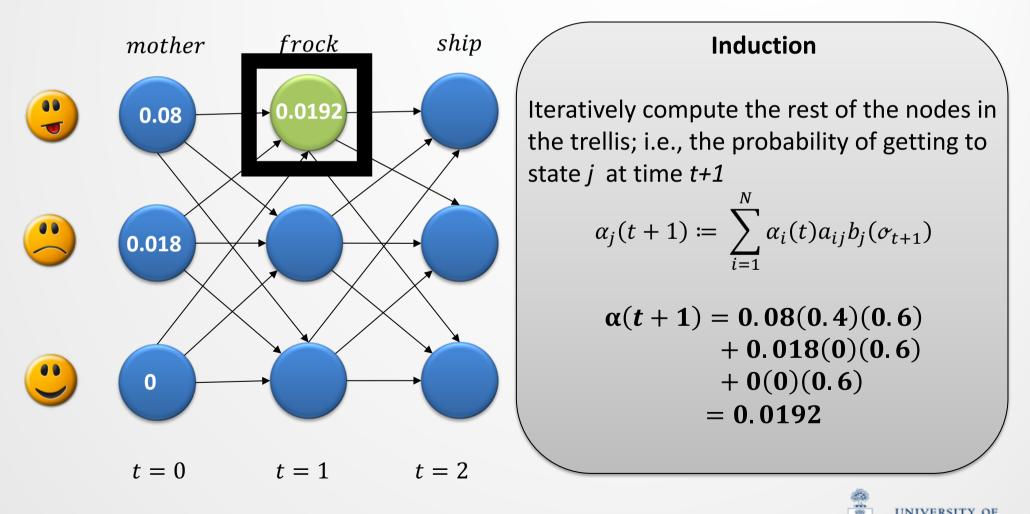


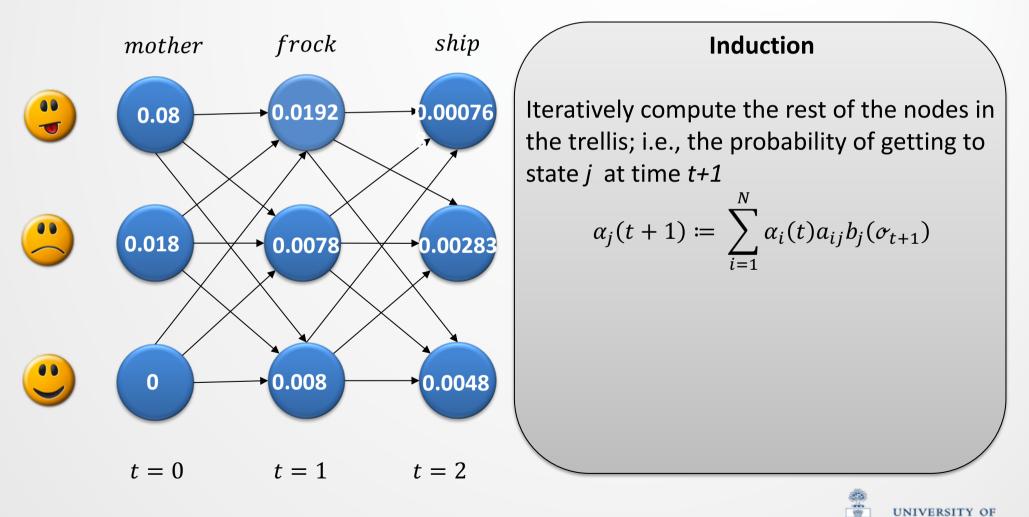
Conclusion of Forward procedure

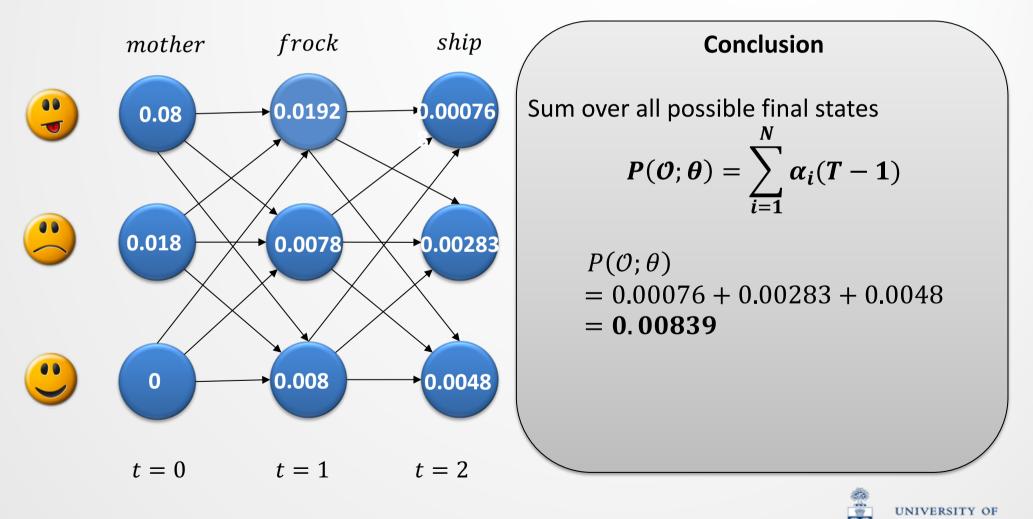












The Forward procedure

- The naïve approach needed $(2T) \cdot N^T$ multiplications.
- The Forward procedure (using dynamic programming) needs only 2N²T multiplications.
- The Forward procedure gives us $P(\mathcal{O}; \theta)$.
- Clearly, but less intuitively, we can also compute the trellis from back-to-front, i.e., backwards in time...



Remember the point

The point was to compute the equivalent of

$$P(\mathcal{O};\theta) = \sum_{Q} P(\mathcal{O},Q;\theta)$$

where $P(\mathcal{O}, Q; \theta) = P(\mathcal{O}|Q; \theta)P(Q; \theta)$ $= \pi_{q_0} b_{q_0}(\sigma_0) a_{q_0q_1} b_{q_1}(\sigma_1) a_{q_1q_2} b_{q_2}(\sigma_2) \dots$ $a_i(0)$ $a_i(1)$

The Forward algorithm stores all possible 1-state sequences (from the start), to store all possible 2-state sequences (from the start), to store all possible 3-state sequences (from the start)...

Remember the point

But, we can compute these factors in reverse
 P(O,Q;θ) = P(O|Q;θ)P(Q;θ)

$$= \pi_{q_0} \dots b_{q_{T-3}} (\sigma_{T-3}) a_{q_{T-3}q_{T-2}} b_{q_{T-2}} (\sigma_{T-2}) a_{q_{T-2}q_{T-1}} b_{q_{T-1}} (\sigma_{T-1})$$

$$\beta_i (T-2)$$

$$\beta_i (T-4)$$

We can still deal with sequences that evolve *forward* in time, but simply **store temporary results in reverse**...



The Backward procedure

• In the $(i, t)^{th}$ node of the **trellis**, we store $\beta_i(t) = P(\sigma_{t+1:T-1} | \sigma_{0:t}, q_t = s_i; \theta)$ $= P(\sigma_{t+1:T-1} | q_t = s_i; \theta)$

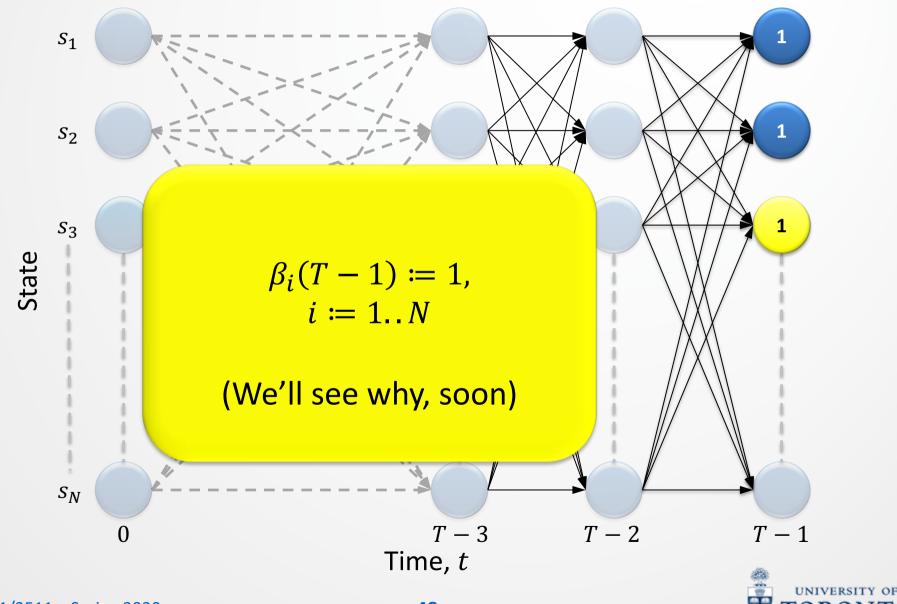
which is computed by summing probabilities on **outgoing** arcs **from** that node.

 $\beta_i(t)$ is the probability of starting in state *i* at time *t* then observing everything that comes thereafter.

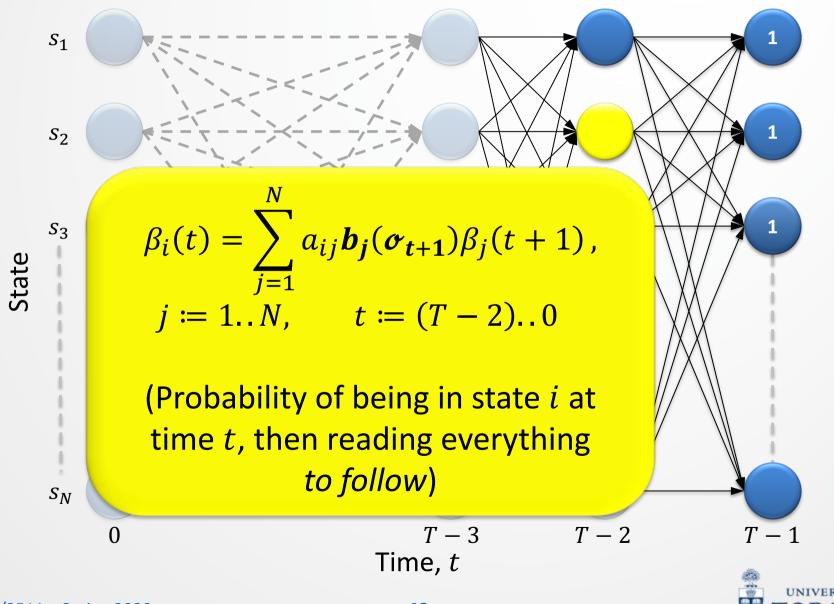
The trellis is computed right-to-left and top-to-bottom.



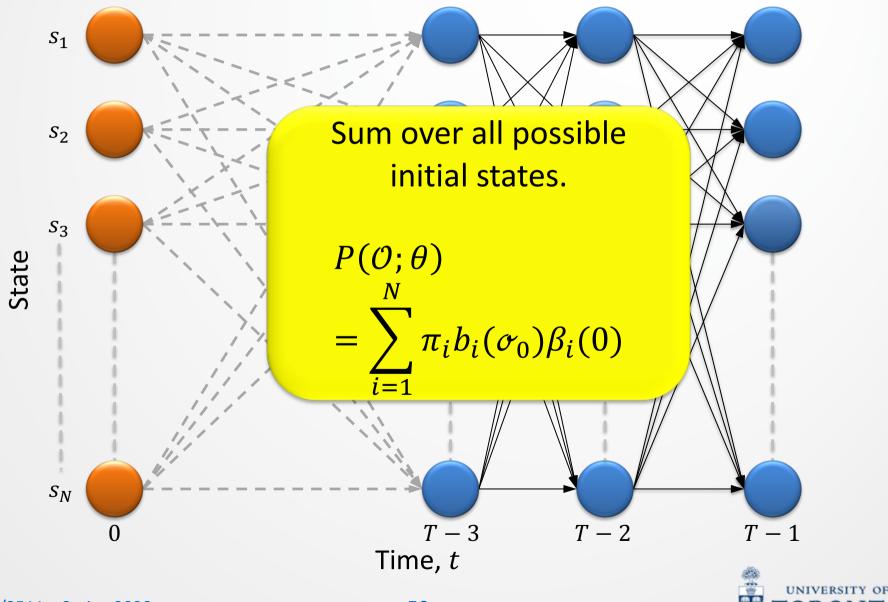
Step 1: Backward initialization



Step 2: Backward induction



Step 3: Backward conclusion



The Backward procedure

- Initialization
 - $\beta_i(T-1) = 1, \qquad i \coloneqq 1..N$
- Induction $\beta_i(t) = \sum_{j=1}^N a_{ij} b_j(\sigma_{t+1}) \beta_j(t+1), \qquad i \coloneqq 1..N$ $t \coloneqq T - 2..0$
- Conclusion $P(\mathcal{O};\theta) = \sum_{i=1}^{N} \pi_{i} b_{i}(\sigma_{0}) \beta_{i}(0)$



The Backward procedure – so what?

- The combination of Forward and Backward procedures will be vital for solving parameter re-estimation, i.e., training.
- Generally, we can combine α and β at any point in time to represent the probability of an entire observation sequence...



Combining α and β

$$P(\mathcal{O}, q_t = i; \theta) = \alpha_i(t)\beta_i(t)$$

$$\therefore P(\mathcal{O}; \theta) = \sum_{i=1}^{N} \alpha_i(t)\beta_i(t)$$

This requires the current word to be incorporated by $\alpha_i(t)$, but **not** $\beta_i(t)$.

> This isn't merely for fun – it will soon become useful...



Fundamental tasks for HMMs

2. Given an observation sequence O and a model θ , how do we choose a state sequence $Q^* = \{q_0, \dots, q_{T-1}\}$ that *best explains* the observations?

This is the task of **inference** – i.e., guessing at the best explanation of unknown ('latent') variables given our model.

This is often an important part of **classification**.



Task 2: Choosing $Q^* = \{q_0 ... q_{T-1}\}$

- The purpose of finding the best state sequence Q* out of all possible state sequences Q is that it tells us what is most likely to be going on 'under the hood'.
- With the Forward algorithm, we didn't care about specific state sequences – we were summing over all possible state sequences.



Task 2: Choosing $Q^* = \{q_0 ... q_{T-1}\}$

In other words,

$$Q^* = \operatorname*{argmax}_{Q} P(\mathcal{O}, Q; \theta)$$

where

$$P(\mathcal{O}, Q; \theta) = \pi_{q_0} b_{q_0}(\sigma_0) \prod_{t=1}^{T-1} a_{q_{t-1}q_t} b_{q_t}(\sigma_t)$$



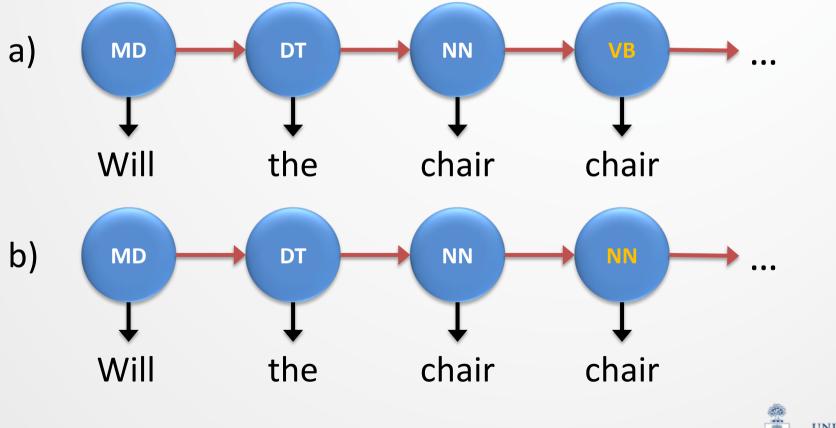
Why choose $Q^* = \{q_0 ... q_{T-1}\}$?

- Recall the purpose of HMMs:
 - To represent multivariate systems where some variable is unknown/hidden/latent.
- Finding the best hidden-state sequence Q^* allows us to:
 - Identify unseen parts-of-speech given words,
 - Identify equivalent English words given French words,
 - Identify unknown phonemes given speech sounds,
 - Decipher hidden messages from encrypted symbols,
 - Identify hidden relationships from gene sequences,
 - Identify hidden market conditions given stock prices,



Example – PoS state sequences

 Will/MD the/DT chair/NN chair/?? the/DT meeting/NN from/IN that/DT chair/NN?



Recall

- Observation likelihoods depend on the state, which changes over time
- We cannot simply choose the state that maximizes the probability of o_t without considering the state

| word | P(word) | |
|--------|---------|--|
| ship | 0.1 | |
| pass | 0.05 | |
| camp | 0.05 | |
| frock | 0.6 | |
| soccer | 0.05 | |
| mother | 0.1 | |
| tops | 0.05 | |

sequence.

| word | P(word) |
|--------|---------|
| ship | 0.25 |
| pass | 0.25 |
| camp | 0.05 |
| frock | 0.3 |
| soccer | 0.05 |
| mother | 0.09 |
| tops | 0.01 |

| word | P(word) | |
|--------|-------------|--|
| ship | 0.3 | |
| pass | 0 | |
| camp | 0 | |
| frock | 0.2 | |
| soccer | 0.05 | |
| mother | 0.05 | |
| tops | 0.4 | |
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The Viterbi algorithm

- The Viterbi algorithm is an inductive dynamicprogramming algorithm that uses a *new kind* of trellis.
- We define **the probability of the most probable path** leading to the trellis node at (state *i*, time *t*) as

$$\boldsymbol{\delta_i(t)} = \max_{q_0 \dots q_{t-1}} P(q_0 \dots q_{t-1}, \boldsymbol{\sigma_0} \dots \boldsymbol{\sigma_t}, \boldsymbol{q_t} = \boldsymbol{s_i}; \boldsymbol{\theta})$$

• $\psi_i(t)$: The best possible previous state, if If I'm in state *i* at time *t*.



Viterbi example

For illustration, we assume a simpler state-transition topology:

0.01

| ler sta | ate-tra | nsition | camp |
|---------|---------|-------------|--------|
| | | | frock |
| logy: | | | soccer |
| | | 0.4 S_d | mother |
| | | Su Su | tops |
| | | | |
| | | 0.1 0.5 | |
| word | P(word) | ↓ ↓ | |
| ship | 0.25 | S_s S_h | |
| pass | 0.25 | 5 10 | |
| camp | 0.05 | | |
| frock | 0.3 | | |
| soccer | 0.05 | 1.0 0.2 | 0.8 |
| mother | 0.09 | | |
| | | | |

| word | P(word) | |
|--------|---------|--|
| ship | 0.1 | |
| pass | 0.05 | |
| camp | 0.05 | |
| frock | 0.6 | |
| soccer | 0.05 | |
| mother | 0.1 | |
| tops | 0.05 | |
| | | |

| word | P(word) | |
|--------|---------|----------|
| ship | 0.3 | |
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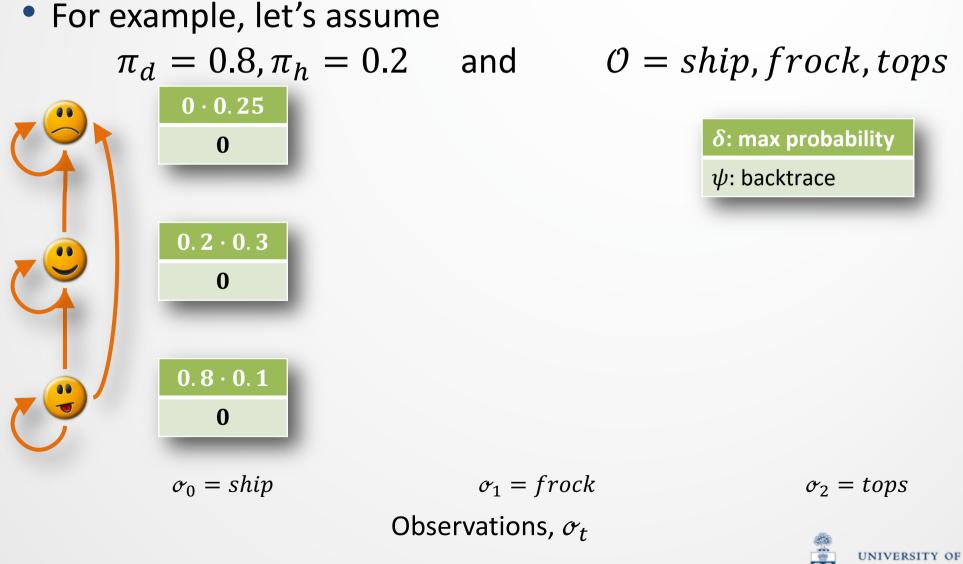
tops

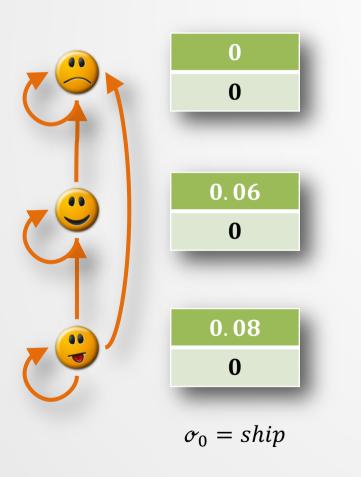
Step 1: Initialization of Viterbi

• Initialize with $\delta_i(0) = \pi_i b_i(\sigma_0)$ and $\psi_i(0) = 0$ for all states.



Step 1: Initialization of Viterbi





The best path to state s_j at time t, $\delta_j(t)$, depends on the best path to each possible previous state, $\delta_i(t-1)$, and their transitions to j, a_{ij}

$$\delta_j(t) = \max_i \left[\delta_i(t-1)a_{ij} \right] b_j(\sigma_t)$$

$$\psi_j(t) = \operatorname*{argmax}_i \left[\delta_i(t-1)a_{ij} \right]$$

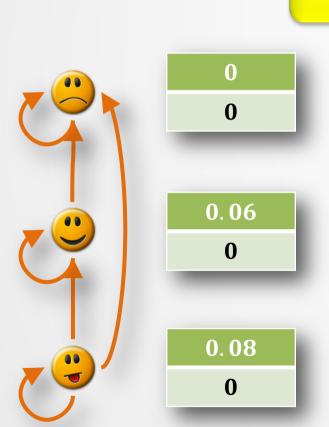
 $\sigma_1 = frock$

Observations, σ_t

 $\sigma_2 = tops$



Specifically...



$$\delta_s(1) = \max_i \left[\delta_i(0) a_{is} \right] b_s(\sigma_1)$$
$$\psi_s(1) = \operatorname*{argmax}_i \left[\delta_i(0) a_{is} \right]$$

$$\delta_h(1) = \max_i \left[\delta_i(0) a_{ih} \right] b_h(\sigma_1)$$
$$\psi_h(1) = \operatorname*{argmax}_i \left[\delta_i(0) a_{ih} \right]$$

| $\delta_d(1) = \max_i \left[\delta_i(0) a_{id} \right] b_d(\sigma_1)$ |
|---|
| $\psi_d(1) = \underset{i}{\operatorname{argmax}} \left[\delta_i(0) a_{id} \right]$ |

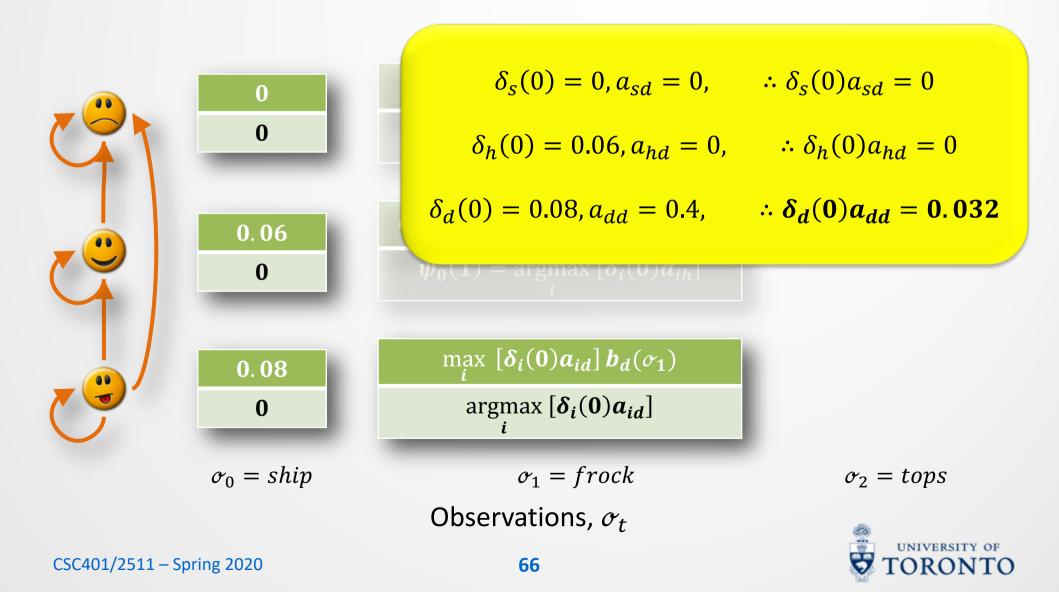
 $\sigma_1 = frock$

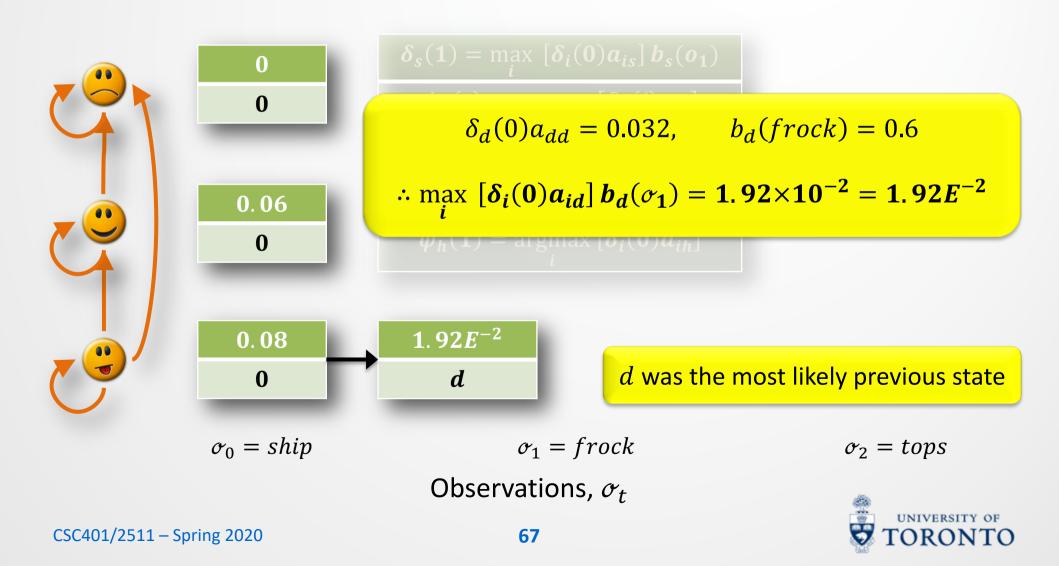
 $\sigma_0 = ship$

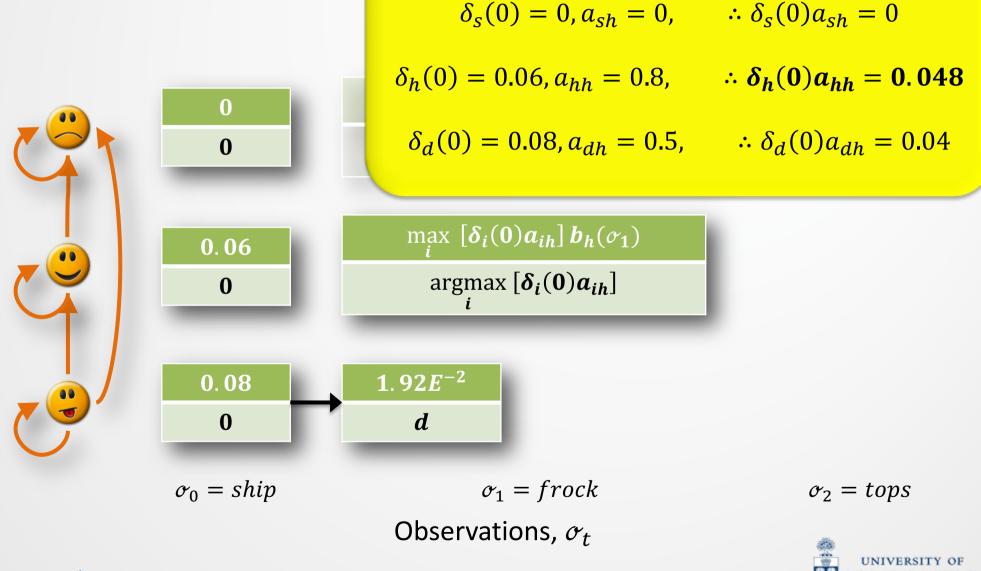
 $\sigma_2 = tops$

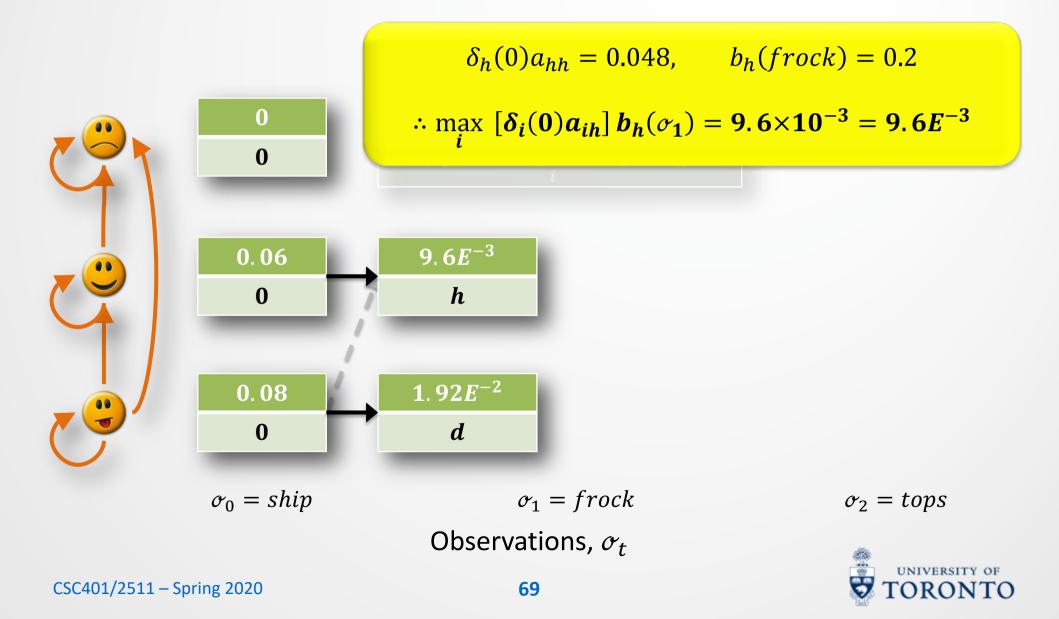


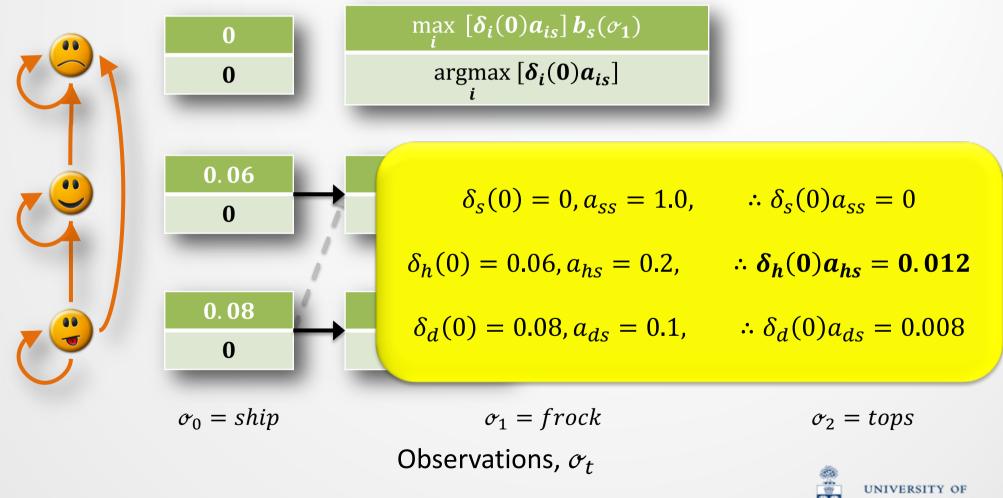
Observations, σ_t



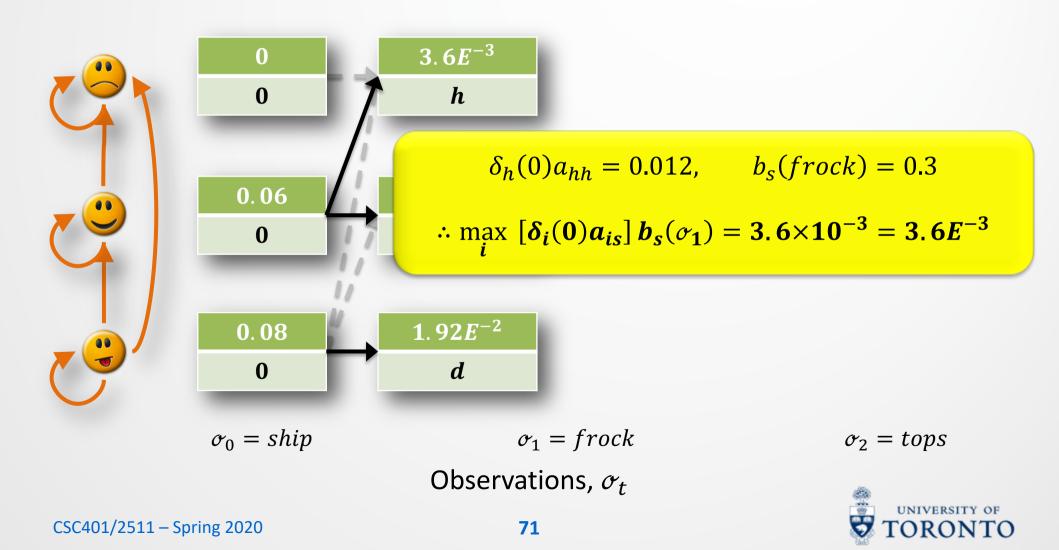


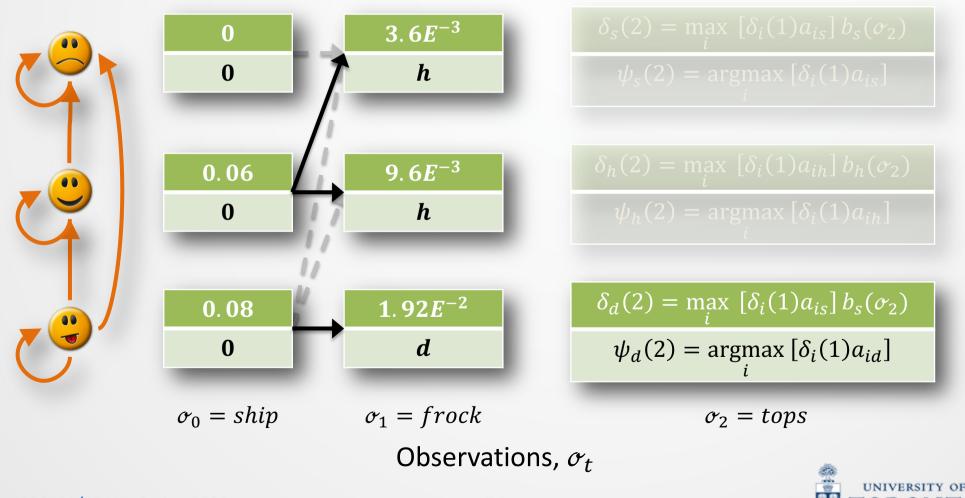


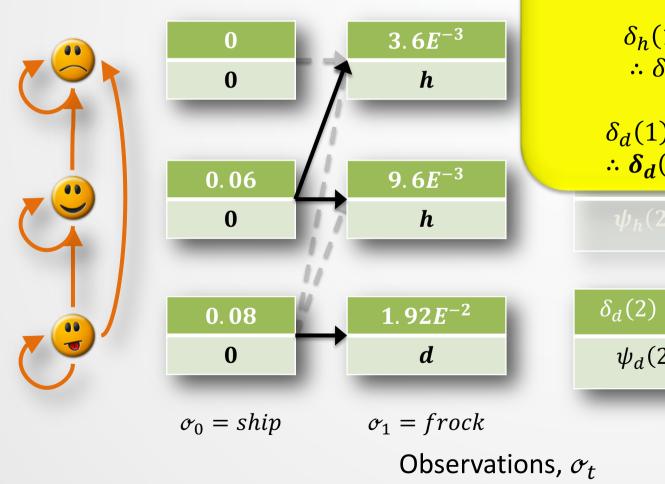




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 $\delta_s(1) = 3.6E^{-3}, a_{sd} = 0,$ $\therefore \delta_s(1)a_{sd} = 0$

$$\delta_h(1) = 9.6E^{-3}, a_{hd} = 0,$$

$$\therefore \delta_h(1)a_{hd} = 0$$

$$\delta_d(1) = 1.92E^{-2}, a_{dd} = 0.4,$$

 $\therefore \delta_d(1)a_{dd} = 0.00768$

 $\boldsymbol{\psi}_{h}(2) = \operatorname{argmax} \left[\boldsymbol{\delta}_{i}(1) \boldsymbol{a}_{ih} \right]$

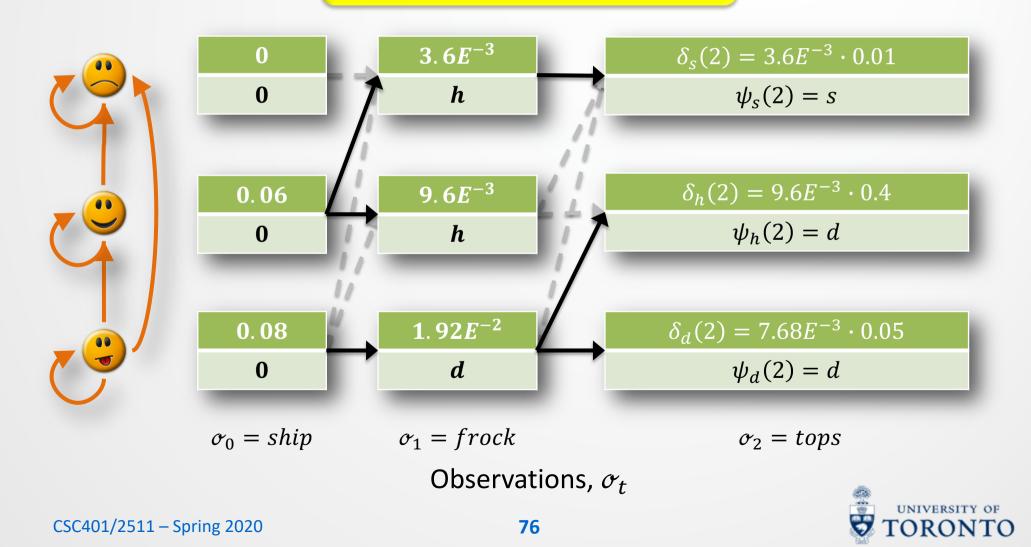
$$\delta_d(2) = \max_i \left[\delta_i(1) a_{is} \right] b_s(\sigma_2)$$

$$\psi_d(2) = \operatorname*{argmax}_i \left[\delta_i(1) a_{id} \right]$$

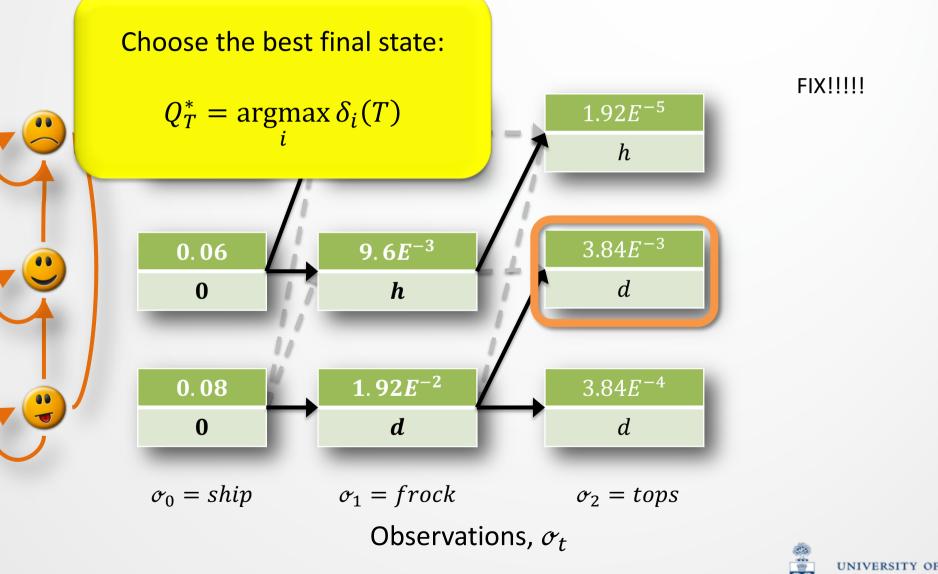
 $\sigma_2 = tops$



Continuing...

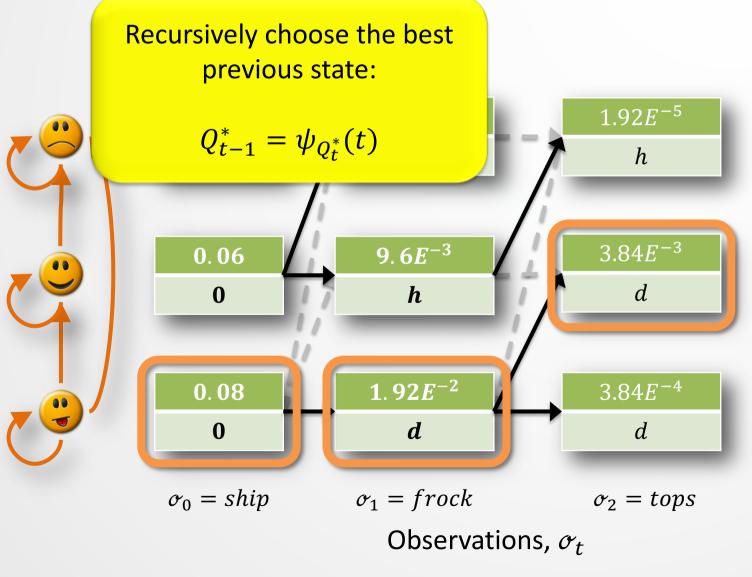


Step 3: Conclusion of Viterbi



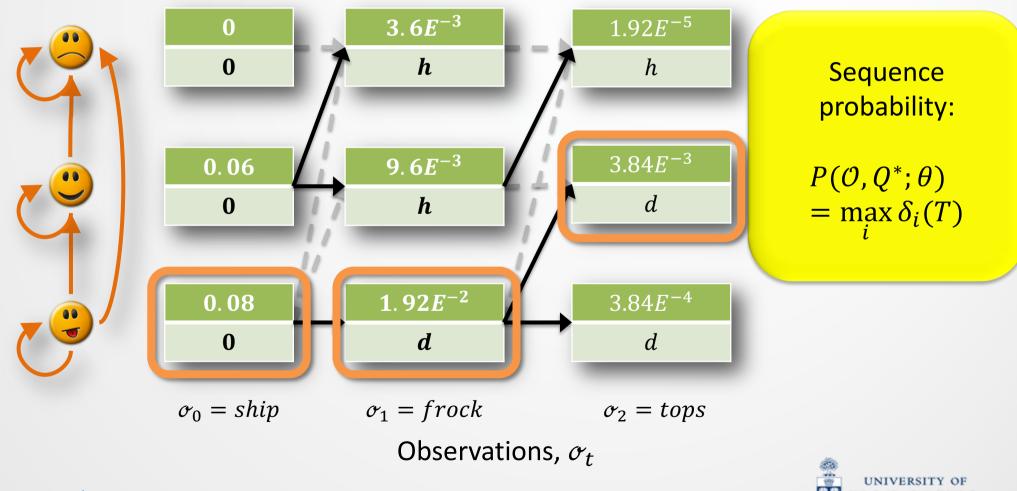
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Step 3: Conclusion of Viterbi





Step 3: Conclusion of Viterbi



ORON

Aside - Working in the log domain

Our formulation was

$$Q^* = \operatorname{argmax}_Q P(\mathcal{O}, Q; \theta)$$

his is equivalent to
$$Q^* = \operatorname{argmin}_Q - \log_2 P(\mathcal{O}, Q; \theta)$$

where

t

$$-\log_2 P(\mathcal{O}, Q; \theta) = -\log_2 \left(\pi_{q_0} b_{q_0}(\sigma_0) \right) - \sum_{t=1}^{T-1} \log_2 \left(a_{q_{t-1}q_t} b_{q_t}(\sigma_t) \right)$$



Fundamental tasks for HMMs

3. Given a large **observation sequence** \mathcal{O} for **training**, but *not* the state sequence, how do we choose the 'best' parameters $\theta = \langle \Pi, A, B \rangle$ that explain the data \mathcal{O} ?

This is the task of **training**.

As with observable Markov models and **MLE**, we want our parameters to be set so that **the available training data is maximally likely**, But doing so will involve **guessing unseen information**...



Task 3: Choosing $\theta = \langle \Pi, A, B \rangle$

We want to modify the parameters of our model
 θ = (Π, A, B) so that P(O; θ) is maximized for some
 training data O:

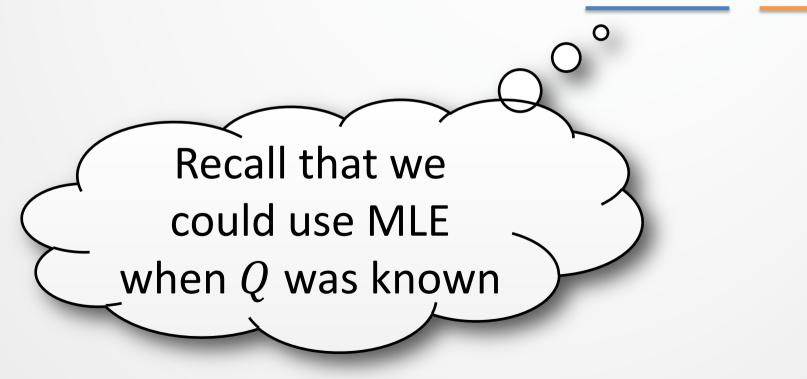
$$\hat{\theta} = \operatorname*{argmax}_{\theta} \frac{P(\mathcal{O}; \theta)}{P(\mathcal{O}; \theta)}$$

 Why? E.g., if we later want to choose the best state sequence Q* for previously unseen test data, the parameters of the HMM should be tuned to similar training data.



Task 3: Choosing $\theta = \langle \Pi, A, B \rangle$

- $\hat{\theta} = \underset{\theta}{\operatorname{argmax}} P(\mathcal{O}; \theta) = \underset{\theta}{\operatorname{argmax}} \sum_{Q} P(\mathcal{O}, Q; \theta)$
- $P(\mathcal{O}, Q; \theta) = P(q_{0:T-1})P(w_{0:t}|q_{0:t}) \approx \prod_{i=0}^{t} P(q_i|q_{i-1})P(w_i|q_i)$





Can we do

this?

Task 3: Choosing $\theta = \langle \Pi, A, B \rangle$

- $P(\mathcal{O}, Q; \theta) = P(q_{0:t})P(w_{0:t}|q_{0:t}) \approx \prod_{i=0}^{t} P(q_i|q_{i-1})P(w_i|q_i)$
- If the training data contained state sequences, we could simply do maximum likelihood estimation, as before:

•
$$P(q_i|q_{i-1}) = \frac{Count(q_{i-1}q_i)}{Count(q_{i-1})}$$
 $P(w_i|q_i) = \frac{Count(w_i \land q_i)}{Count(q_i)}$

- But we <u>don't</u> know the states; we <u>can't</u> count them.
- However, we can use an iterative hill-climbing approach if we can guess the counts using a "good" pre-existing model



Expecting and maximizing

• If we knew θ , we could make **expectations** such as

- Expected number of times in state s_i,
- Expected number of transitions $s_i \rightarrow s_j$
- If we knew:
 - Expected number of times in state s_i,
 - Expected number of transitions $s_i \rightarrow s_j$

then we could compute the maximum likelihood estimate of

 $\theta = \left\langle \pi_i, \left\{ a_{ij} \right\}, \left\{ b_i(w) \right\} \right\rangle$



Expectation-maximization

 Expectation-maximization (EM) is an iterative training algorithm that alternates between two steps:



guesses the **expected** counts for the hidden sequence using the current model θ_k .

• Maximization (M): computes a new θ that maximizes the likelihood of the data, given the guesses of the E-step. This θ_{k+1} is then used in the next E-step.

Continue until convergence or stopping condition...



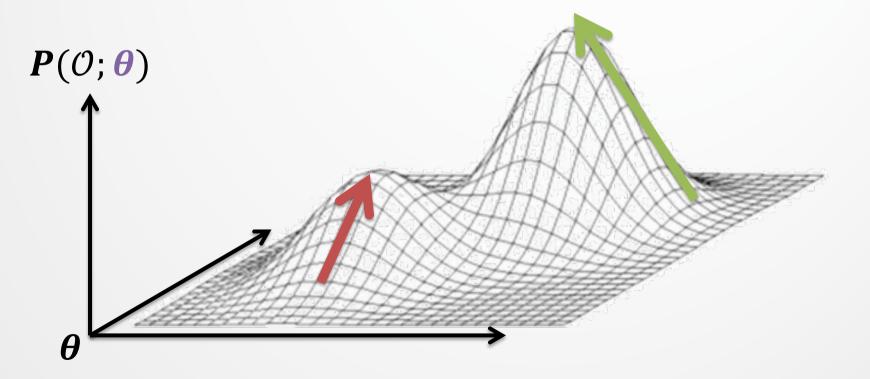
Baum-Welch re-estimation

- Baum-Welch (BW): *n.* a specific version of EM for HMMs. a.k.a. 'forward-backward' algorithm.
 - **1.** Initialize the model.
 - 2. Compute **expectations** for $Count(q_{t-1}q_t)$ and $Count(q_t \land w_t)$ given model, training data \mathcal{O} .
 - 3. Adjust our start, transition, and observation probabilities to maximize the likelihood of O.
 - 4. Go to 2. and repeat until convergence or stopping condition...



Local maxima

- Baum-Welch changes θ to climb a `hill' in $P(\mathcal{O}; \theta)$.
 - How we initialize θ can have a big effect.



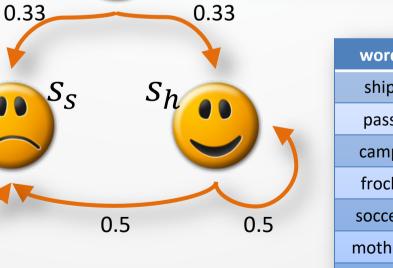


Step 1: BW initialization

- Our **initial guess** for the parameters, θ_0 , can be:
 - a) All probabilities are **uniform** (e.g., $b_i(w_a) = b_i(w_b)$ for all states *i* and words *w*) 0.33

| word | P(word) |
|--------|---------|
| ship | 0.143 |
| pass | 0.143 |
| camp | 0.143 |
| frock | 0.143 |
| soccer | 0.143 |
| mother | 0.143 |
| tops | 0.143 |

| word | P(word) | |
|--------|---------|--|
| ship | 0.143 | |
| pass | 0.143 | |
| camp | 0.143 | |
| frock | 0.143 | |
| soccer | 0.143 | |
| mother | 0.143 | |
| tops | 0.143 | |



 S_d

| word | P(word) | |
|--------------------------|---------|--|
| ship | 0.143 | |
| pass | 0.143 | |
| camp | 0.143 | |
| frock | 0.143 | |
| soccer | 0.143 | |
| mother | 0.143 | |
| tops | 0.143 | |
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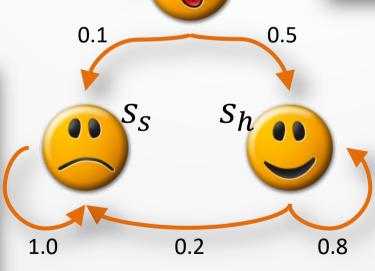
1.0

Step 1: BW initialization

• Our initial guess for the parameters, θ_0 , can be: b) All probabilities are drawn randomly (subject to the condition that $\sum_i P(i) = 1$) $0.4 \bigcirc S_d$

| word | P(word) |
|--------|---------|
| ship | 0.1 |
| pass | 0.05 |
| camp | 0.05 |
| frock | 0.6 |
| soccer | 0.05 |
| mother | 0.1 |
| tops | 0.05 |

| word | P(word) |
|--------|---------|
| ship | 0.25 |
| pass | 0.25 |
| camp | 0.05 |
| frock | 0.3 |
| soccer | 0.05 |
| mother | 0.09 |
| tops | 0.01 |

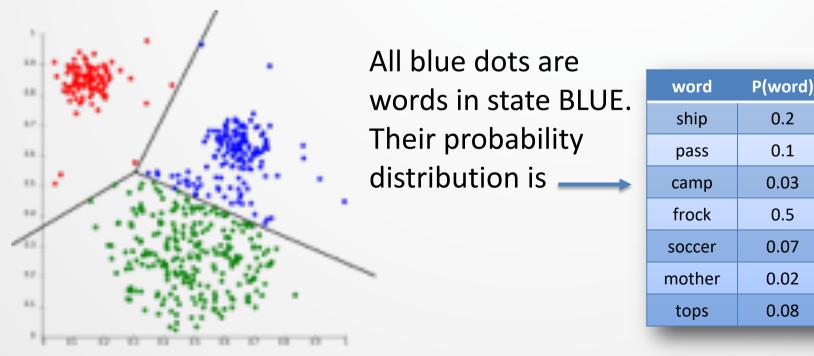


| word | P(word) | | |
|---------------|---------|--|--|
| ship | 0.3 | | |
| pass | 0 | | |
| camp | 0 | | |
| frock | 0.2 | | |
| soccer | 0.05 | | |
| mother | 0.05 | | |
| tops | 0.4 | | |
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Step 1: BW initialization

• Our **initial guess** for the parameters, θ_0 , can be:

c) Observation distributions are drawn from prior distributions: e.g., $b_i(w_a) = P(w_a)$ for all states *i*. *sometimes* this involves pre-clustering, e.g. *k*-means





What to expect when you're expecting

- If we knew θ , we could estimate **expectations** such as
 - Expected number of times in state s_i,
 - Expected number of transitions $s_i \rightarrow s_j$
- If we knew:
 - Expected number of times in state s_i,
 - Expected number of transitions $s_i \rightarrow s_j$

then we could compute the **maximum likelihood estimate** of $\theta = \langle \{a_{ij}\}, \{b_i(w)\}, \pi_i \rangle$



BW E-step (occupation)

• We define

$$\gamma_i(t) = P(q_t = i | \mathcal{O}; \theta_k)$$

as the probability of **being** in state *i* at time *t*, based on our current model, θ_k , **given** the <u>entire</u> observation, O.

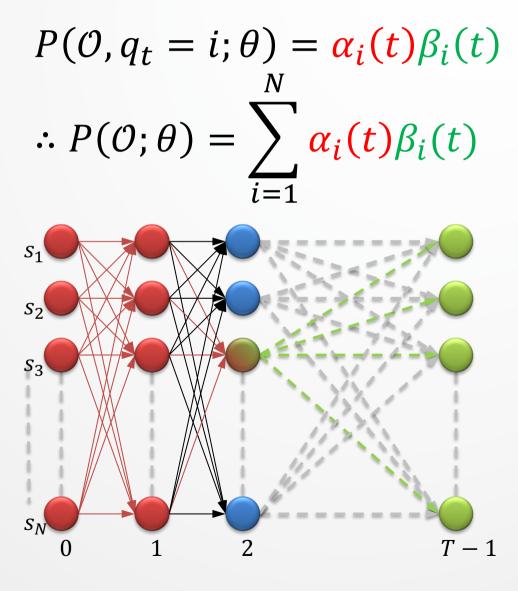
and rewrite as:

$$\gamma_{i}(t) = \frac{P(q_{t} = i, \mathcal{O}; \theta_{k})}{P(\mathcal{O}; \theta_{k})}$$
$$= \frac{\alpha_{i}(t)\beta_{i}(t)}{P(\mathcal{O}; \theta_{k})}$$

Remember, $\alpha_i(t)$ and $\beta_i(t)$ depend on values from $\theta = \langle \pi_i, a_{ij}, b_i(w) \rangle$



Combining α and β





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BW E-step (transition)

• We define

$$\xi_{ij}(t) = P(q_t = i, q_{t+1} = j | \mathcal{O}; \theta_k)$$

as the probability of **transitioning** from state *i* at time *t* to state *j* at time t + 1 **based on** our current model, θ_k , and **given** the <u>entire</u> observation, O. This is:

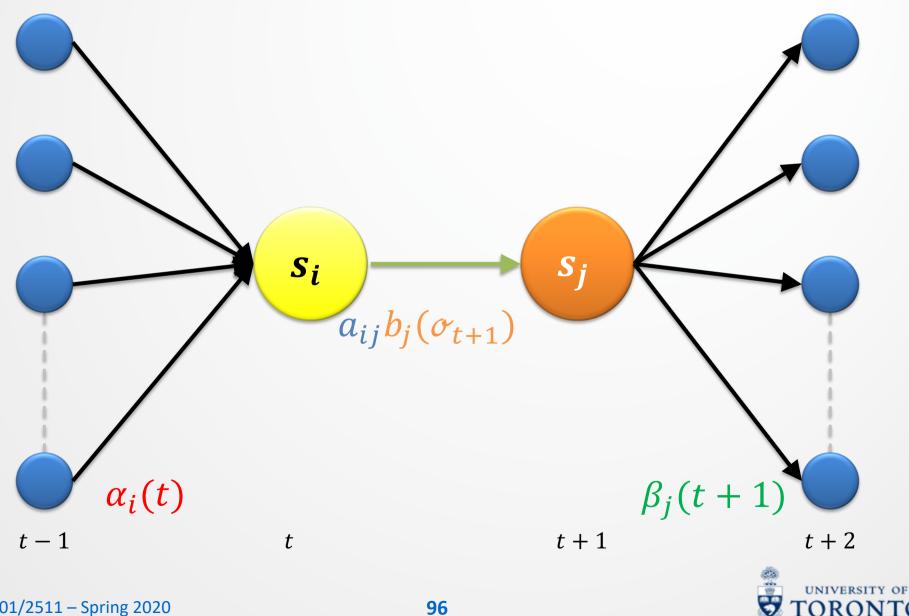
$$\xi_{ij}(t) = \frac{P(q_t = i, q_{t+1} = j, 0; \theta_k)}{P(0; \theta_k)}$$

=
$$\frac{\alpha_i(t)a_{ij}b_j(\sigma_{t+1})\beta_j(t+1)}{P(0; \theta_k)}$$
Again, these estimates come from our model at

iteration k, θ_k .



BW E-step (transition)



Expecting and maximizing

• If we knew θ , we could estimate **expectations** such as

- Expected number of times in state s_i,
- Expected number of transitions $s_i \rightarrow s_j$
- If we knew:
 - Expected number of times in state s_i,
 - Expected number of transitions $s_i \rightarrow s_j$

then we could compute the **maximum likelihood estimate** of $\theta = \langle \{a_{ij}\}, \{b_i(w)\}, \pi_i \rangle$



BW M-step

We update our parameters as if we were doing MLE:

Initial-state probabilities: for $i \coloneqq 1..N$ $\hat{\pi}_i = \gamma_i(0)$ $P(q_j|q_i)$ $Count(q_i q_j)$ $Count(q_i)$ II. State-transition probabilities: \circ O C $\hat{a}_{ij} = \frac{\sum_{t=0}^{T-2} \xi_{ij}(t)}{\sum_{t=0}^{T-2} \gamma_i(t)}$ for $i, j \coloneqq 1..N$ $P(w_i|q_i)$ $Count(w_i \wedge q_i)$ III. Discrete observation probabilities: ° O $\widehat{b}_{j}(w) = \frac{\sum_{t=0}^{T-1} \gamma_{j}(t)|_{\sigma_{t}=w}}{\sum_{t=0}^{T-1} \gamma_{i}(t)} \text{ for } j \coloneqq 1..N \text{ and } w \in \mathcal{V}$



Baum-Welch iteration

• We update our parameters after **each iteration** $\theta_{k+1} = \left\langle \hat{\pi}_i, \hat{a}_{ij}, \hat{b}_j(w) \right\rangle$

rinse, and repeat until $\theta_k \approx \theta_{k+1}$ (until change almost stops).

• Baum proved that $P(\mathcal{O}; \theta_{k+1}) \ge P(\mathcal{O}; \theta_k)$

although this method does *not* guarantee a *global maximum*.



Features of Baum-Welch

- Although we're not guaranteed to achieve a global optimum, the local optima are often 'good enough'.
- BW does not estimate the number of states, which must be 'known' beforehand.
 - Moreover, some constraints on topology are often imposed beforehand to assist training.





Discrete vs. continuous

- If our observations are drawn from a continuous space (e.g., speech acoustics), the probabilities *b_i(X)* must also be continuous.
- HMMs generalize to continuous distributions, or multivariate observations,

e.g., $b_i([-14.28, 0.85, 0.21])$.



Adaptation

- It can take a <u>LOT</u> of data to train HMMs.
- Imagine that we're given a **trained** HMM but not the data.
 - Also imagine that this HMM has been trained with data from many sources (e.g., many speakers).
- We want to use this HMM with a particular new source for whom we have some data (but not enough to fully train the HMM properly from scratch).
 - To be more accurate for that source, we want to change the original HMM parameters *slightly* given the new data.



HMM interpolation

For added robustness, we can combine estimates of a generic HMM, *G*, trained with lots of data from many sources with a specific HMM, *S*, trained with a little data from a single source.

$$P_{Interp}(\sigma) = \lambda P(\sigma; \theta_G) + (1 - \lambda) P(\sigma; \theta_S)$$

- This gives us a model tuned to our target source (S), but with some general 'knowledge' (G) built in.
 - How do we pick $\lambda \in [0..1]$?



EM for interpolated models

- Strategy can be used for any $P(\mathcal{O}; \lambda) = \sum_i \lambda_i P_i(\mathcal{O})$
- Introduce latent states s such that $P(s = i; \lambda) = \lambda_i$
- Once in state i, $P(\mathcal{O}|s = i; \lambda) = P_i(\mathcal{O})$
- Like with HMMs, we estimate Count(s = i) using EM:

$$\lambda_{i}^{new} = \frac{P(s = i, \mathcal{O}; \lambda^{old})}{P(\mathcal{O}; \lambda^{old})}$$

 This is a (simplified) version of what is done for Jelinek-Mercer interpolation, as well as Gaussian Mixture Models (covered in ASR lecture)



Held-out data

- Let $T_{\lambda} = \{O\}$ be the data used to learn λ , T_i for $P_i(\cdot)$
- If for most $\mathcal{O} \in T_{\lambda}$, $j \cdot P_i(\mathcal{O}) \ge P_j(\mathcal{O})$, then $\lambda_i \to 1$
- This can easily occur when $T_i = T_{\lambda}$, e.g.:
 - If $P_i(\cdot)$ is an MLE *i*-gram model trained on T_{λ} , it will outperform $P_{\langle i}(\cdot)$ (even if also trained on T_{λ})
 - If $P(\sigma; \theta_S)$ was trained on T_{λ} but not $P(\sigma; \theta_G)$
- Less likely to happen when $T_i \cap T_\lambda = \emptyset$
- A disjoint T_{λ} is often called **held-out** or **development data**



Aside – Maximum a Posteriori (MAP)

- Given adaptation data \mathcal{O}_a , the MAP estimate is $\hat{\theta} = \operatorname{argmax}_{\theta} P(\mathcal{O}_a | \theta) P(\theta)$
- If we can guess some structure for $P(\theta)$, we can use EM to estimate new parameters (or Monte Carlo).
- For continuous $b_i(\sigma)$, we use **Dirichlet distribution** that defines the hyper-parameters of the model and the **Lagrange method** to describe the change in parameters $\theta \Rightarrow \hat{\theta}$.



Summary

• Important ideas to know:

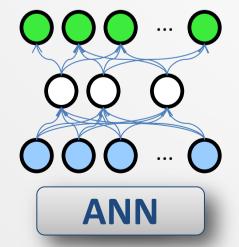
- The definition of an HMM (e.g., its parameters).
- The purpose of the Forward algorithm.
 - How to compute $\alpha_i(t)$ and $\beta_i(t)$
- The purpose of the Viterbi algorithm.
 - How to compute $\delta_i(t)$ and $\psi_i(t)$.
- The purpose of the **Baum-Welch algorithm**.
 - Some understanding of EM.
 - Some understanding of the equations.

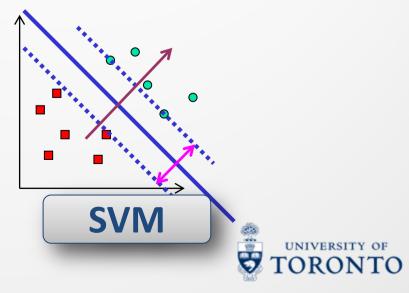


Generative vs. discriminative

- HMMs are generative classifiers. You can generate synthetic samples from because they model the phenomenon itself.
 (e.g. P(O,Q; θ) or P(O; θ))
- Other classifiers (e.g., artificial neural networks and support vector machines) are discriminative in that their probabilities are trained specifically to reduce the error in classification.
 (e.g. P(Q|O; θ))

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Reading

- (optional) Manning & Schütze: Section 9.2—9.4.1
 - Note that they use another formulation...
- Rabiner, L. (1990) A Tutorial on Hidden Markov Models and Selected Applications in Speech Recognition. In: *Readings in speech recognition*. Morgan Kaufmann. (posted on course website)
- Optional software:
 - Hidden Markov Model Toolkit (<u>http://htk.eng.cam.ac.uk/</u>)
 - Sci-kit's HMM (<u>http://scikit-learn.sourceforge.net/stable/modules/hmm.html</u>)

