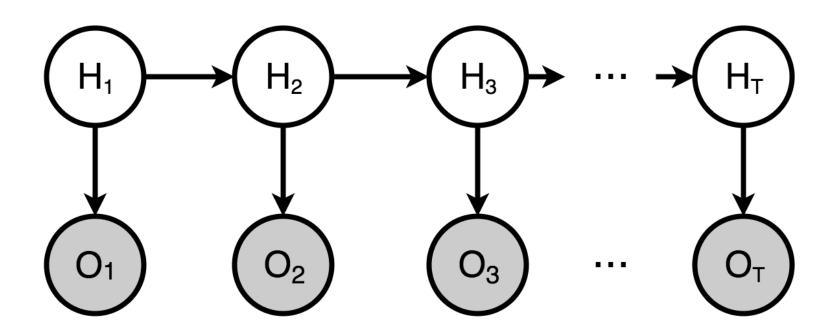
# DSCI 552, Machine Learning for Data Science

University of Southern California

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# Lesson 11 Hidden Markov Models



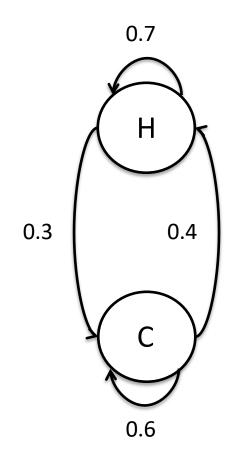
#### Hidden Markov Models

What is a hidden Markov model (HMM)? A machine learning technique and... ...a discrete hill climb technique Two for the price of one! Where are HMMs used? Speech recognition, information security, and too many other things to list Q: Why are HMMs so useful? A: Widely applicable and *efficient* algorithms

```
Markov chain
  "Memoryless random
  process"
  Transitions depend only on
  current state (Markov chain
  of order 1)...
  ...and transition probability
  matrix
```

Suppose we're interested in average annual temperature Only consider Hot and Cold From recorded history, obtain probabilities for...

Year-to-year transitions
Based on thermometer
readings for "recent" years



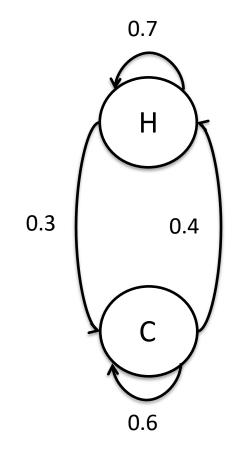
Transition probability matrix

Matrix is denoted as A

[ 0.4 0.0 ]

Note, A is "row stochastic"

$$A = \left[ \begin{array}{cc} 0.7 & 0.3 \\ 0.4 & 0.6 \end{array} \right]$$

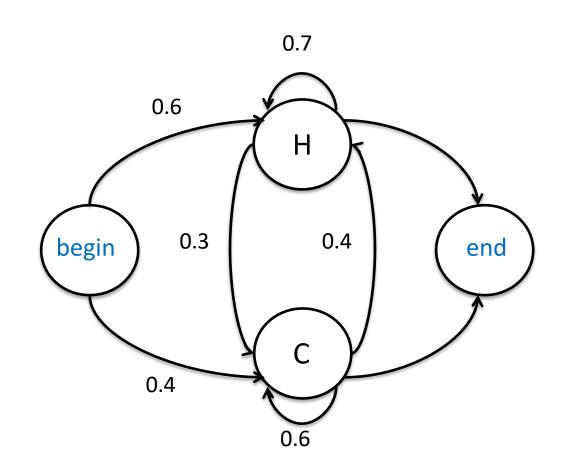


So, each element of A is between 0 and 1 and each row satisfies the definition of a discrete probability distribution, thus the elements of any given row sum to 1.

Can also include begin, end states Matrix for begin state denoted π In this example,

$$\pi = \left[ \begin{array}{cc} 0.6 & 0.4 \end{array} \right]$$

Note that π is also row stochastic



#### Hidden Markov Model

#### HMM includes a Markov chain/ process

- But the Markov process is "hidden", i.e., we can't directly observe the Markov process
- Instead, observe things that are probabilistically related to hidden states
- It's as if there is a "curtain" between Markov chain and the observations

Consider H/C annual temp example Suppose we want to know H or C annual temperature in distant past

- Before thermometers were invented
- Note, we only distinguish between H and C

We assume transition between Hot and Cold years is same as today

Then the A matrix is known

Temps in past follow a Markov process

But, we cannot observe temperature in past We find evidence that tree ring size is related to temperature

Looking at historical data, we find this holds We only consider 3 tree ring sizes

Small, Medium, Large (S, M, L, respectively)
Measure tree ring sizes and recorded temperatures
to determine relationship

We find that tree ring sizes and temperature related by S = M = L

$$\begin{array}{c|cccc}
H & \begin{bmatrix} 0.1 & 0.4 & 0.5 \\ 0.7 & 0.2 & 0.1 \end{bmatrix}
\end{array}$$

This is known as the B matrix

The matrix B is also row stochastic

$$B = \left[ \begin{array}{ccc} 0.1 & 0.4 & 0.5 \\ 0.7 & 0.2 & 0.1 \end{array} \right]$$

Can we now find H/C temps in past?
We cannot measure (observe) temps
But we can measure tree ring sizes...
...and tree ring sizes related to temps
By probabilities in the B matrix
Can we say something intelligent about temps over some interval in the past?

#### **HMM Notation**

# A lot of notation is required Notation may be the most difficult part...

```
T = the length of the observation sequence N = the number of states in the model M = the number of observation symbols Q = \{q_0, q_1, \dots, q_{N-1}\} = the states of the Markov process V = \{0, 1, \dots, M-1\} = set of possible observations A = the state transition probabilities B = the observation probability matrix \pi = the initial state distribution \mathcal{O} = (\mathcal{O}_0, \mathcal{O}_1, \dots, \mathcal{O}_{T-1}) = observation sequence.
```

#### **HMM Notation**

Note that for simplicity, observations taken from  $V = \{0, 1, ..., M-1\}$ That is,  $\mathcal{O}_i \in V$  for i = 0, 1, ..., T-1The matrix  $A = \{a_{ij}\}$  is N x N, where  $a_{ij} = P(\text{state } q_i \text{ at } t+1 | \text{state } q_i \text{ at } t)$ 

The matrix  $B = \{b_i(k)\}\$  is  $N \times M$ , where

 $b_j(k) = P(\text{observation } k \text{ at } t \mid \text{state } q_j \text{ at } t).$ 

Consider our temperature example...

What are the possible observations?

 $V = \{0,1,2\}$ , corresponding to S, M, L

What are states of Markov process?

$$Q = \{H,C\}$$

What are A,B,  $\pi$ , and T?

A,B, π on previous slides

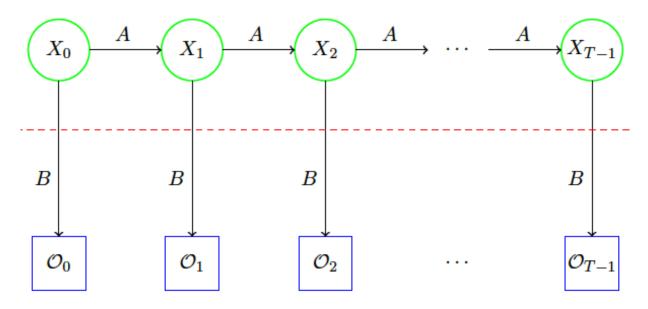
T is number of tree rings measured

What are N and M?

$$N = 2$$
 and  $M = 3$ 

#### Generic HMM

#### Generic view of HMM



HMM defined by A,B, and  $\pi$ We denote HMM "model" as  $\lambda = (A,B,\pi)$ 

Suppose that we observe tree ring sizes For a 4 year period of interest: S,M,S,L Then  $\mathcal{O} = (\mathcal{O}_0, \mathcal{O}_1, \mathcal{O}_2, \mathcal{O}_3) = (0, 1, 0, 2)$ Most likely (hidden) state sequence? That is, most likely  $X = (X_0, X_1, X_2, X_3)$ Let  $\pi_{X0}$  be prob. of starting in state  $x_0$ Note  $b_{x_0}(\mathcal{O}_0)$  prob. of initial observation And  $a_{X_0,X_1}$  is prob. of transition  $X_0$  to  $X_1$ And so on...

$$\pi = \begin{bmatrix} 0.6 & 0.4 \end{bmatrix}$$

Bottom line?

We can compute P(X,O) for any XFor  $X = (X_0, X_1, X_2, X_3)$  we have P(X,O)

$$= \pi_{x_0} b_{x_0}(\mathcal{O}_0) a_{x_0,x_1} b_{x_1}(\mathcal{O}_1) a_{x_1,x_2} b_{x_2}(\mathcal{O}_2) a_{x_2,x_3} b_{x_3}(\mathcal{O}_3)$$

Suppose we observe (0,1,0,2), then what is probability of, say, HHCC?

S,M,S,L

Plug into formula above to find *P*(*HHCC*,0,1,0,2)

$$= 0.6(0.1)(0.7)(0.4)(0.3)(0.7)(0.6)(0.1) = 0.000212.$$

Do same for all 4state sequences We find that the winner is... CCCH Not so fast!

		normalized
$\operatorname{state}$	probability	probability
HHHH	.000412	.042787
HHHC	.000035	.003635
HHCH	.000706	.073320
HHCC	.000212	.022017
HCHH	.000050	.005193
HCHC	.000004	.000415
HCCH	.000302	.031364
HCCC	.000091	.009451
CHHH	.001098	.114031
CHHC	.000094	.009762
CHCH	.001882	.195451
CHCC	.000564	.058573
CCHH	.000470	.048811
CCHC	.000040	.004154
CCCH	.002822	.293073
CCCC	.000847	.087963

Sum = P(O) P(X|O) = P(X,O)/P(O)

The *path* CCCH scores the highest In dynamic programming (DP), we find highest scoring path But, in HMM we maximize *expected number of correct states*Sometimes called "EM algorithm" For "Expectation Maximization" How does HMM work in this example?

For first position...

Sum probabilities for all paths that have H in 1<sup>st</sup> position, compare to sum of probabilities for paths with C in 1<sup>st</sup> position: biggest wins

Repeat for each position and we find

	element			
	0	1	2	3
P(H O)	0.188182	0.519576	0.228788	0.804029
P(C O)	0.811818	0.480424	0.771212	0.195971

Note: We could use both P(X,O) and P(X|O) to decide the best. Especially in paper and pencil problems, it is easier to use the unnormalized P(X,O)

	element			
	0	1	2	3
P(H O)	0.188182	0.519576	0.228788	0.804029
P(C O)	0.811818	0.480424	0.771212	0.195971

So, HMM solution gives us CHCH
While DP solution is CCCH
Which solution is better?
Neither solution is better!
Just using different definitions of "best"

	element			
	0	1	2	3
P(H O)	0.188182	0.519576	0.228788	0.804029
P(C O)	0.811818	0.480424	0.771212	0.195971

	element			
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So, HMM solution gives us CHCH
While DP solution is CCCH
Which solution is better?
Neither solution is better!
Just using different definitions of "best"

#### **HMM Paradox?**

HMM maximizes expected number of correct states

Whereas DP chooses "best" overall path Possible for HMM to choose a "path" that is impossible

Could be a transition probability of 0 Cannot get impossible path with DP Is this a flaw with HMM?

No, it's a feature

# Probability of Observations

Table computed for

$$\mathcal{O} = (\mathcal{O}_0, \mathcal{O}_1, \mathcal{O}_2, \mathcal{O}_3)$$
  
= (0,1,0,2)

For this sequence,

$$P(O) = .000412 + .000035 + .000706 + ... +$$

.000847

= left to the reader

Similarly for other observations  $\mathcal{O}$ 

		$ onume{normalized} $
$\operatorname{state}$	probability	probability
HHHH	.000412	.042787
HHHC	.000035	.003635
HHCH	.000706	.073320
HHCC	.000212	.022017
HCHH	.000050	.005193
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CHCH	.001882	.195451
CHCC	.000564	.058573
CCHH	.000470	.048811
CCHC	.000040	.004154
CCCH	.002822	.293073
CCCC	.000847	.087963

# Probability of Observations

If this calculation is made for all possible 4-observation sequences then the sum of the resulting probabilities (not the normalized probabilities) will be 1.

		normalized
	1 1 1114	
state	probability	probability
HHHH	.000412	.042787
HHHC	.000035	.003635
HHCH	.000706	.073320
HHCC	.000212	.022017
HCHH	.000050	.005193
HCHC	.000004	.000415
HCCH	.000302	.031364
HCCC	.000091	.009451
CHHH	.001098	.114031
CHHC	.000094	.009762
CHCH	.001882	.195451
CHCC	.000564	.058573
CCHH	.000470	.048811
CCHC	.000040	.004154
CCCH	.002822	.293073
CCCC	.000847	.087963

#### **HMM Model**

An HMM is defined by the three matrices, A, B, and  $\pi$ Note that M and N are implied, since they are the dimensions of matrices So, we denote an HMM "model" as  $\lambda = (A,B,\pi)$ 

# The Three Problems HMMs used to solve 3 problems:

Problem 1: Given a model  $\lambda = (A,B,\pi)$  and observation sequence O, find  $P(O|\lambda)$ 

That is, we can **score** an observation sequence to see how well it fits a given model

#### The Three Problems

HMMs used to solve 3 problems Problem 2: Given  $\lambda = (A,B,\pi)$  and O, find an optimal state sequence (in HMM sense)

Uncover hidden part (like previous example)

In many applications in NLP, the solution to Problem 2 is crucial. Example: Finding the grammatical roles of words in a sentence.

#### The Three Problems

HMMs used to solve 3 problems

Problem 3: Given O, N, and M, find the model λ that maximizes probability of O

That is, *train* a model to fit observations

#### **HMMs** in Practice

Often, HMMs used as follows:

Given an observation sequence...

Assume that (hidden) Markov process exists

Train a model based on observations

That is, solve Problem 3

"Best" N can be found by trial and error

Then given a sequence of observations, score it versus the model we trained

This is Problem 1: high score implies similar to training data, low score says it's not

#### **HMMs** in Practice

In this sense, HMM is a "machine learning" technique

To train a model, we do not need to specify anything except the parameter N "Best" N often found by trial and error So, we don't need to "think" too much Just train HMM and then use it Fortunately, there are efficient algorithms for HMMs

### The Three Solutions

We give detailed solutions to 3 problems

Note: We must find *efficient* solutions

The three problems:

Problem 1: Score an observation sequence versus a given model

Problem 2: Given a model, "uncover" hidden part

Problem 3: Given an observation sequence, train a model

Recall that we considered example for 2 and 1, but direct solutions are very inefficient

# The Three Solutions Appendix: Algorithmic Details

#### Solution 1

Score observations versus a given model Given model  $\lambda = (A,B,\pi)$  and observation sequence  $O=(O_0,O_1,...,O_{T-1})$ , find  $P(O|\lambda)$ 

Denote hidden states as

$$X = (X_0, X_1, ..., X_{T-1})$$

Then from definition of B,

$$P(O|X,\lambda)=b_{X0}(O_0) b_{X1}(O_1) \dots b_{XT-1}(O_{T-1})$$

And from definition of A and  $\pi$ ,

$$P(X|\lambda) = \pi_{X0} a_{X0,X1} a_{X1,X2} \dots a_{XT-2,XT-1}$$

Elementary conditional probability fact:

$$P(O,X|\lambda) = P(O|X,\lambda) P(X|\lambda)$$

Sum over all possible state sequences X,

$$P(O|\lambda) = \sum P(O,X|\lambda) = \sum P(O|X,\lambda) P(X|\lambda)$$
$$= \sum \pi_{X_0} b_{X_0}(O_0) a_{X_0,X_1} b_{X_1}(O_1) ... a_{X_{T-2},X_{T-1}} b_{X_{T-1}}(O_{T-1})$$

This "works" but way too costly

Requires about 2TN<sup>T</sup> multiplications

Why?

There should be a better way...

```
Instead, use forward algorithm
Or "alpha pass"
For t = 0,1,...,T-1 and i=0,1,...,N-1, let \alpha_t(i) = P(O_0,O_1,...,O_t,X_t=q_i|\lambda)
Probability of "partial observation" to t, and Markov process is in state q_i at step t Can be computed recursively, efficiently
```

Let  $\alpha_0(i) = \pi_i b_i(O_0)$  for i = 0,1,...,N-1For t = 1,2,...,T-1 and i=0,1,...,N-1, let

$$\alpha_t(i) = \left(\sum \alpha_{t-1}(j)a_{ji}\right)b_i(O_t)$$

Where the sum is from j = 0 to N-1 From definition of  $\alpha_t(i)$  we see

$$P(O|\lambda) = \Sigma \alpha_{T-1}(i)$$

Where the sum is from i = 0 to N-1 This requires only N<sup>2</sup>T multiplications

Note: The score  $P(O|\lambda)$  is dependent on the length of the observation sequence. Consequently, to compare scores for sequences of different length, we can normalize to a per observation score, that is, score =  $\Sigma \alpha_{T-1}(i) / T$ .

In a nutshell,  $\alpha_T(i)$  is the probability that the sequence of observations up to time T correspond to the state being the i<sup>th</sup> state at time T, given the model  $\lambda$ .

We compute  $\alpha_t(i)$  recursively using  $\alpha_{t-1}(j)$ 's.

Or "beta pass"

Given a model, find hidden states
Given λ = (A,B,π) and O, find an optimal state sequence
Recall that optimal means "maximize expected number of correct states"
In contrast, DP finds best scoring path
For temp/tree ring example, solved this
But hopelessly inefficient approach
A better way: *backward algorithm* 

### **Backward Algorithm**

For t = 0,1,...,T-1 and i = 0,1,...,N-1, let  $\beta_t(i) = P(O_{t+1},O_{t+2},...,O_{T-1}|X_t=q_i,\lambda)$ Probability of partial observation from t to end and Markov process in state  $q_i$  at step t Analogous to the forward algorithm As with forward algorithm, this can be computed recursively and efficiently

## **Backward Algorithm**

Let 
$$\beta_{T-1}(i) = 1$$
 for  $i = 0,1,...,N-1$   
For  $t = T-2,T-3,...,1$  and  $i = 0,1,...,N-1$ , let

$$\beta_t(i) = \sum a_{ij}b_j(O_{t+1})\beta_{t+1}(j)$$

Where the sum is from j = 0 to N-1

Note that 
$$\gamma_t(i) = \alpha_t(i)\beta_t(i)/P(O|\lambda)$$
  
And recall  $P(O|\lambda) = \Sigma\alpha_{T-1}(i)$ 

The bottom line?

Forward algorithm solves Problem 1 Forward/backward algorithms solve Problem 2 by computing  $\gamma_t(i)$  for each state at each time step (using both forward and backward paths) and choosing the state that maximizes it.

$$\widetilde{X}_t = \max_i \gamma_t(i).$$

Why is it necessary to normalize gamma by dividing by P(O | lambda)? Because these probabilities are computed assuming the observation sequence is known (i.e., given O), as opposed to being computed relative to the larger probability space.

Train a model: Given O, N, and M, find  $\lambda$  that maximizes probability of O We'll iteratively adjust  $\lambda = (A,B,\pi)$  to better fit the given observations O

The size of matrices are fixed (N and M)

But elements of matrices can change
It is nice that this works...

...and amazing that it's efficient!

```
For t=0,1,...,T-2 and i,j in {0,1,...,N-1}, define "di-gammas" as  \gamma_t(i,j) = P(X_t = q_i, \ X_{t+1} = q_j | O, \lambda)  Note \gamma_t(i,j) is prob of being in state q_i at time t and transiting to state q_j at t+1 Then \gamma_t(i,j) = \alpha_t(i)a_{ij}b_j(O_{t+1})\beta_{t+1}(j)/P(O|\lambda)  And \gamma_t(i) = \Sigma\gamma_t(i,j)  Where sum is from j=0 to N-1
```

### Baum-Welch Model Re-estimation

```
Given di-gammas and gammas...
For i = 0, 1, ..., N-1 let \pi_i = \gamma_0(i)
For i = 0,1,...,N-1 and j = 0,1,...,N-1
      a_{ii} = \sum \gamma_t(i,j)/\sum \gamma_t(i)
   Where both sums are from t = 0 to T-2
For j = 0,1,...,N-1 and k = 0,1,...,M-1
      b_i(k) = \sum \gamma_t(j) / \sum \gamma_t(j)
   Both sums from from t = 0 to T-1 but only t for
   which O₁ = k are counted in numerator
```

### Baum-Welch Model Re-estimation

#### Why does this work?

For the given observation sequence O, the sum  $\Sigma\gamma_t(i)$  gives us the current best estimate for the total probability of being in state i, while  $\Sigma\gamma_t(i,j)$  gives us the total probability of transitioning from state i to state j. Hence, the ratio  $\Sigma\gamma_t(i,j)/\Sigma\gamma_t(i)$  enables us to re-estimate  $a_{ij}$  based on the current model parameters and observation sequence.

#### To summarize...

- 1. Initialize  $\lambda = (A,B,\pi)$
- 2. Compute  $\alpha_t(i)$ ,  $\beta_t(i)$ ,  $\gamma_t(i,j)$ ,  $\gamma_t(i)$
- 3. Re-estimate the model  $\lambda = (A,B,\pi)$
- 4. If P(O|λ) increases by more than ε, goto 2 (where ε is small)

Some fine points... Model initialization If we have a good guess for  $\lambda = (A,B,\pi)$  then we can use it for initialization If not, let  $\pi_i \approx 1/N$ ,  $a_{i,i} \approx 1/N$ ,  $b_i(k) \approx 1/M$ Subject to row stochastic conditions But, do **not** initialize to exactly uniform values Stopping conditions

Stop after some number of iterations and/or... Stop if increase in  $P(O|\lambda)$  is too small

### HMM as Discrete Hill Climb

Algorithm on previous slides shows that HMM is a "discrete hill climb"

HMM consists of discrete parameters

Specifically, the elements of the matrices And re-estimation process improves model by modifying parameters

So, process "climbs" toward improved model

This happens in a high-dimensional space

Brief detour...

For  $\lambda = (A,B,\pi)$  as above, it's easy to define a dynamic program (DP)

**Executive summary:** 

DP is forward algorithm, with "sum" replaced by "max"

Let  $\delta_0(i) = \pi_i b_i(O_0)$  for i=0,1,...,N-1For t=1,2,...,T-1 and i=0,1,...,N-1 compute  $\delta_{t}(i) = \max \left( \delta_{t-1}(j) a_{ii} \right) b_{i}(O_{t})$ Where the max is over j in {0,1,...,N-1} Note that at each t, the DP computes best path for each state, up to that point So, probability of best path is max  $\delta_{T_{-1}}(j)$ This max gives the highest probability Not the best path, for that, see next slide

To determine optimal path

While computing deltas, keep track of pointers to previous state

When finished, construct optimal path by tracing back points

For example, consider temp example: recall that we observe (0,1,0,2)

$$\pi = \begin{bmatrix} 0.6 & 0.4 \end{bmatrix}$$
 $S \quad M \quad L$ 
 $H \quad C$ 
 $C \begin{bmatrix} 0.1 & 0.4 & 0.5 \\ 0.7 & 0.2 & 0.1 \end{bmatrix}$ 
 $C \begin{bmatrix} 0.7 & 0.3 \\ 0.4 & 0.6 \end{bmatrix}$ 

Probabilities for path of length 1:

$$P(H) = \pi_0 b_0(0) = 0.6(0.1) = 0.06$$
 and  $P(C) = \pi_1 b_1(0) = 0.4(0.7) = 0.28$ .

These are the only "paths" of length 1

Probabilities for each path of length 2

```
P(HH) = 0.06(0.7)(0.4) = 0.0168

P(HC) = 0.06(0.3)(0.2) = 0.0036

P(CH) = 0.28(0.4)(0.4) = 0.0448

P(CC) = 0.28(0.6)(0.2) = 0.0336
```

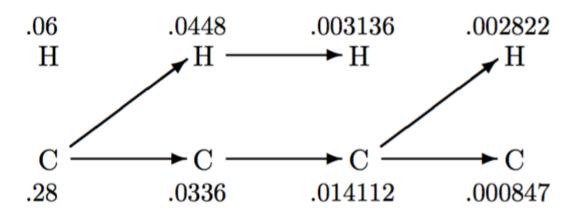
Best path of length 2 ending with H is CH

Best path of length 2 ending with C is CC

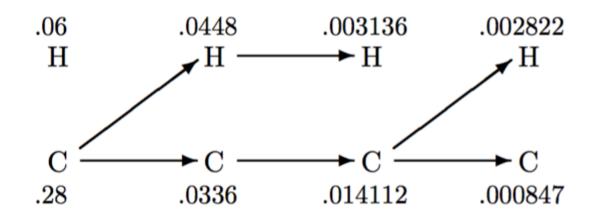
## Dynamic Program $\delta_{t}(i) = \max \left( \delta_{t-1}(j) a_{ji} \right) b_{i}(O_{t})$

Continuing, we compute best path ending at H and C at each step

And save pointers: why?



### Dynamic Program



Best final score is .002822

And thanks to pointers, best path is CCCH

But what about underflow?

A serious problem in bigger cases

## Dynamic Program

https://www.youtube.com/watch?v=6JVq utwtzmo&t=4s

### **Underflow Resistant DP**

```
Common trick to prevent underflow:
  Instead of multiplying probabilities...
  ...add logarithms of probabilities
Why does this work?
  Because log(xy) = log x + log y
  Adding logs does not tend to 0
Note that these logs are negative...
...and we must avoid 0 probabilities
```

#### **Underflow Resistant DP**

Underflow resistant DP algorithm:

Let 
$$\delta_0(i) = log(\pi_i \ b_i(O_0))$$
 for  $i=0,1,...,N-1$   
For  $t=1,2,...,T-1$  and  $i=0,1,...,N-1$  compute 
$$\delta_t(i) = max \left( \delta_{t-1}(j) + log(a_{ji}) + log(b_i(O_t)) \right)$$
Where the max is over  $j$  in  $\{0,1,...,N-1\}$   
And **score** of best path is max  $\delta_{T-1}(j)$   
As before, must also keep track of paths

Trickier to prevent underflow in HMM We consider solution 3

Since it includes solutions 1 and 2 Recall for t = 1,2,...,T-1, i=0,1,...,N-1,

$$\alpha_{t}(i) = \left( \sum \alpha_{t-1}(j) a_{j,i} \right) b_{i}(O_{t})$$

The idea is to normalize alphas so that they sum to 1

Algorithm on next slide

```
Given \alpha_t(i) = \left(\Sigma \alpha_{t-1}(j) a_{j,i}\right) b_i(O_t)

Let a_0(i) = \alpha_0(i) for i=0,1,...,N-1

Let c_0 = 1/\Sigma a_0(j)

For i=0,1,...,N-1, let \alpha_0(i) = c_0 a_0(i)

This takes care of t=0 case

Algorithm continued on next slide...
```

```
For t = 1,2,...,T-1 do the following:

For i = 0,1,...,N-1,
a_t(i) = \left(\Sigma \alpha_{t-1}(j)a_{j,i}\right)b_i(O_t)
Let c_t = 1/\Sigma a_t(j)

For i = 0,1,...,N-1 let \alpha_t(i) = c_t a_t(i)
```

Easy to show 
$$\alpha_t(i) = c_0c_1...c_t \alpha_t(i)$$
 (#)  
Simple proof by induction  
So,  $c_0c_1...c_t$  is scaling factor at step t  
Also, easy to show that  $\alpha_t(i) = \alpha_t(i)/\Sigma\alpha_t(j)$   
Which implies  $\Sigma a_{T-1}(i) = 1$  (##)

```
By combining (#) and (##), we have 1 = \Sigma \alpha_{f-1}(i) = c_0 c_1 ... c_{T-1} \Sigma \alpha_{T-1}(i) = c_0 c_1 ... c_{T-1} P(O|\lambda) Therefore, P(O|\lambda) = 1 / c_0 c_1 ... c_{T-1} To avoid underflow, we compute \log P(O|\lambda) = -\Sigma \log(c_j) Where sum is from j = 0 to T-1
```

Similarly, scale betas as  $c_t\beta_t(i)$ For re-estimation,

Compute  $\gamma_t(i,j)$  and  $\gamma_t(i)$  using original formulas, but with scaled alphas, betas

This gives us new values for  $\lambda = (A,B,\pi)$  "Easy exercise" to show re-estimate is exact when scaled alphas and betas used

Also,  $P(O|\lambda)$  cancels from formula Use log  $P(O|\lambda) = -\Sigma \log(c_i)$  to decide if iterate improves

## All Together Now

```
Complete pseudo code for Solution 3
Given: (O_0, O_1, ..., O_{T-1}) and N and M
Initialize: \lambda = (A,B,\pi)
   A is NxN, B is NxM and \pi is 1xN
   \pi_i \approx 1/N, a_{ii} \approx 1/N, b_i(k) \approx 1/M, each matrix row
   stochastic, but not uniform
Initialize:
   maxIters = max number of re-estimation steps
   iters = 0
   oldLogProb = -\infty
```

## Forward algorithm With scaling

```
// compute \alpha_0(i)
c_0 = 0
for i = 0 to N - 1
     \alpha_0(i) = \pi(i)b_i(\mathcal{O}_0)
     c_0 = c_0 + \alpha_0(i)
next i
// scale the \alpha_0(i)
c_0 = 1/c_0
for i = 0 to N - 1
     \alpha_0(i) = c_0\alpha_0(i)
next i
// compute \alpha_t(i)
for t = 1 to T - 1
     c_t = 0
     for i = 0 to N - 1
          \alpha_t(i) = 0
          for j = 0 to N - 1
               \alpha_t(i) = \alpha_t(i) + \alpha_{t-1}(j)a_{ii}
          next i
          \alpha_t(i) = \alpha_t(i)b_i(\mathcal{O}_t)
          c_t = c_t + \alpha_t(i)
     next i
     // scale \alpha_t(i)
     c_t = 1/c_t
     for i = 0 to N - 1
          \alpha_t(i) = c_t \alpha_t(i)
     next i
next t
```

## Backward Algorithm

Backward algorithm or "beta pass"

With scaling Note: same scaling factor as alphas

```
// Let \beta_{T-1}(i)=1 scaled by c_{T-1} for i=0 to N-1 \beta_{T-1}(i)=c_{T-1} next i

// \beta-pass for t=T-2 to 0 by -1 for i=0 to N-1 \beta_t(i)=0 for j=0 to N-1 \beta_t(i)=\beta_t(i)+a_{ij}b_j(\mathcal{O}_{t+1})\beta_{T+1}(j) next j // scale \beta_t(i) with same scale factor as \alpha_t(i) \beta_t(i)=c_t\beta_t(i) next i next t
```

### Gammas

# Using scaled alphas and betas

```
for t=0 to T-2
    denom = 0
    for i = 0 to N - 1
         for j = 0 to N - 1
              denom = denom + \alpha_t(i)a_{ij}b_j(\mathcal{O}_{t+1})\beta_{t+1}(j)
         next j
    next i
    for i = 0 to N - 1
         \gamma_t(i) = 0
         for j = 0 to N - 1
              \gamma_t(i,j) = (lpha_t(i)a_{ij}b_j(\mathcal{O}_{t+1})eta_{t+1}(j))/	ext{denom}
              \gamma_t(i) = \gamma_t(i) + \gamma_t(i, j)
         next j
    next i
next t
// Special case for \gamma_{T-1}(i)
denom = 0
for i = 0 to N - 1
    \mathtt{denom} = \mathtt{denom} + \alpha_{T-1}(i)
next i
for i = 0 to N - 1
    \gamma_{T-1}(i) = \alpha_{T-1}(i)/\mathtt{denom}
next i
```

#### Re-Estimation

Again, using scaled gammas So formulas unchanged

```
// re-estimate \pi
for i = 0 to N - 1
   \pi_i = \gamma_0(i)
next i
// re-estimate A
for i = 0 to N - 1
   for j = 0 to N - 1
       numer = 0
       denom = 0
       for t = 0 to T - 2
           numer = numer + \gamma_t(i, j)
           denom = denom + \gamma_t(i)
       next t
       a_{ij} = numer/denom
   next i
next i
// re-estimate B
for i = 0 to N - 1
   for j = 0 to M - 1
       numer = 0
        denom = 0
       for t = 0 to T - 2
           if(\mathcal{O}_t == j) then
               numer = numer + \gamma_t(i)
           end if
           denom = denom + \gamma_t(i)
       next t
       b_i(j) = \text{numer/denom}
   next j
next i
```

## Stopping Criteria

Check that
probability increases
In practice, want
logProb >
oldLogProb + ε
And don't exceed
max iterations

```
// Compute \log[P(\mathcal{O} \mid \lambda)]
\log \operatorname{Prob} = 0
\text{for } i = 0 \text{ to } T - 1
\log \operatorname{Prob} = \log \operatorname{Prob} + \log(c_i)
\text{next } i
\log \operatorname{Prob} = -\log \operatorname{Prob}
// To iterate or not to iterate, that is the question...
\text{iters} = \text{iters} + 1
\text{if (iters} < \max \operatorname{Iters} \text{ and logProb} > \operatorname{oldLogProb}) \text{ then }
\operatorname{oldLogProb} = \operatorname{logProb}
\operatorname{goto} \alpha \text{-pass}
\text{else}
\operatorname{output} \lambda = (\pi, A, B)
\text{end if}
```

#### References

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