

Hidden Markov Models I

Machine Learning 10-601B

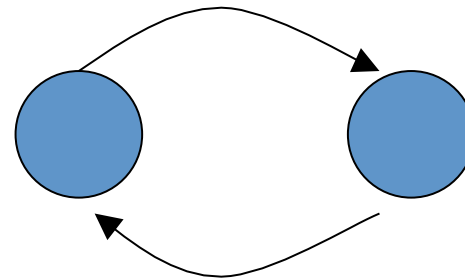
Seyoung Kim

Many of these slides are derived from Tom Mitchell, Ziv Bar-Joseph. Thanks!

What's wrong with Bayesian networks

- Bayesian networks are very useful for modeling joint distributions
- But they have their limitations:
 - Cannot account for temporal / sequence models
 - DAG's (no self or any other loops)

**This is not a valid
Bayesian network!**



Hidden Markov models

- Model a set of observation with a set of hidden states
 - Robot movement

Observations: range sensor, visual sensor

Hidden states: location (on a map)

Hidden Markov models

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Observations: range sensor, visual sensor

 **Hidden states:** location (on a map)



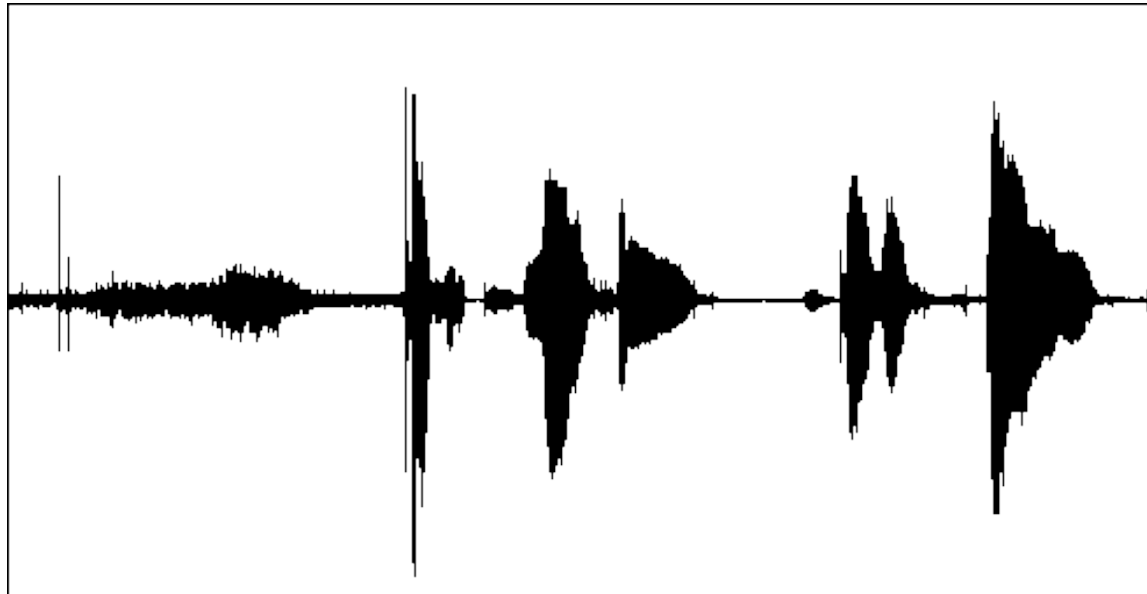
1. Hidden states generate observations
2. Hidden states transition to other hidden states

Examples: Speech processing

Speech processing

Observations: sound signals

Hidden states: parts of speech, words



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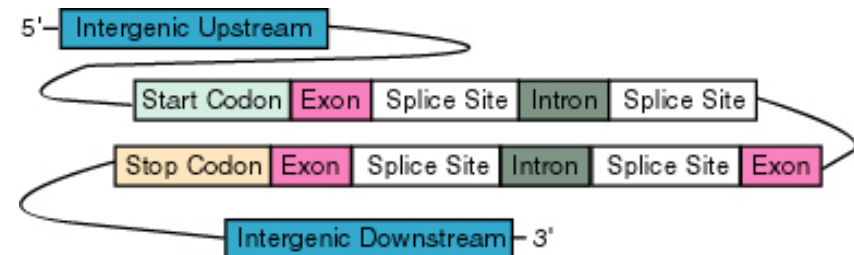
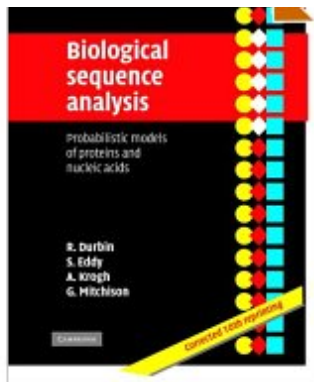
sil	spk	spk	sil	spk	spk
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Example: Biological data

Biology

Observations: DNA base pairs

Hidden states: Genes



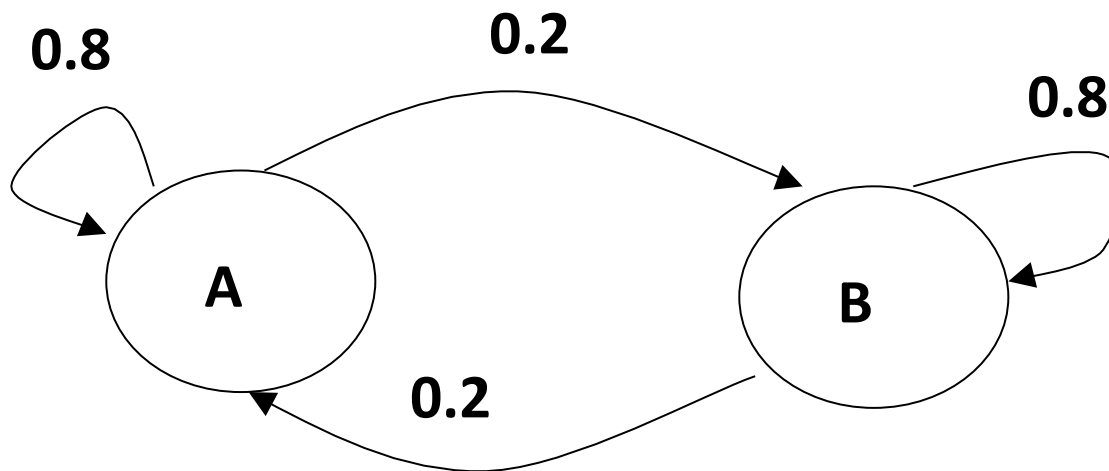
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TGCTCCAAAGAAAAACCGAAGTGCGCCAAGTGTCTGAAGAA
CAACTGGGAGTGTCTGCTAC
TCTCCCAAACCAAAGGTCTCCGCTGACTAGGGCACATCTG
ACAGAAGTGGAATCAAGG
CTAGAAAGACTGGAACAGCTATTTCTACTGATTTTTCCTCGAG
AAGACCTTGACATGATT
```

Example: Gambling on dice outcome

- Two dice A and B, both skewed (output model).
- Can either stay with the same die or switch to the second die (transition model).



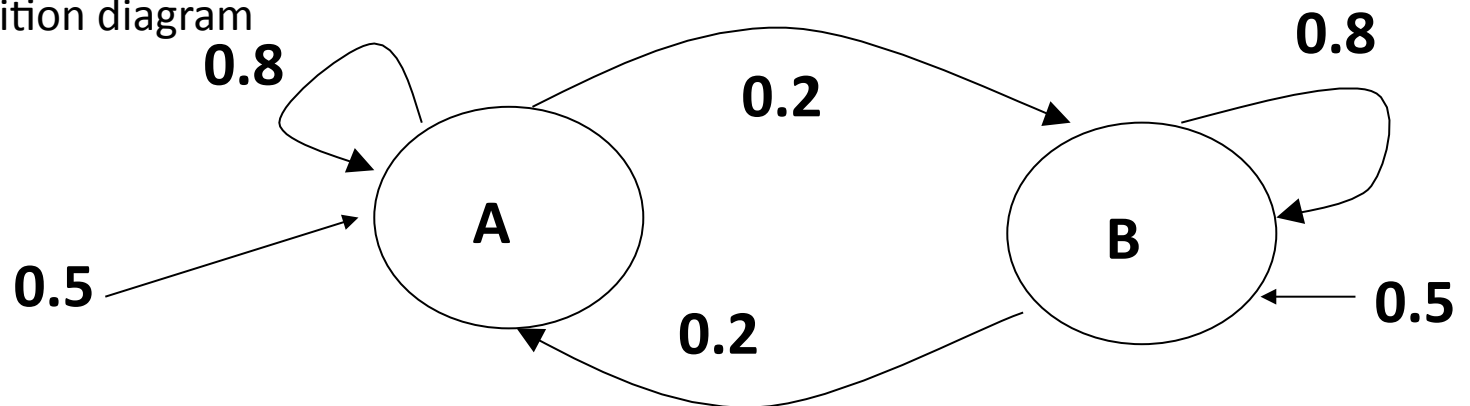
State transition diagram



A Hidden Markov Model

- A set of states $S = \{s_1 \dots s_n\}$
 - In each time point we are in exactly one of these states
- A set of possible outputs $\Sigma = \{\sigma_1, \dots, \sigma_m\}$
 - In each time point we emit a symbol $\sigma_j \in \Sigma$

State transition diagram



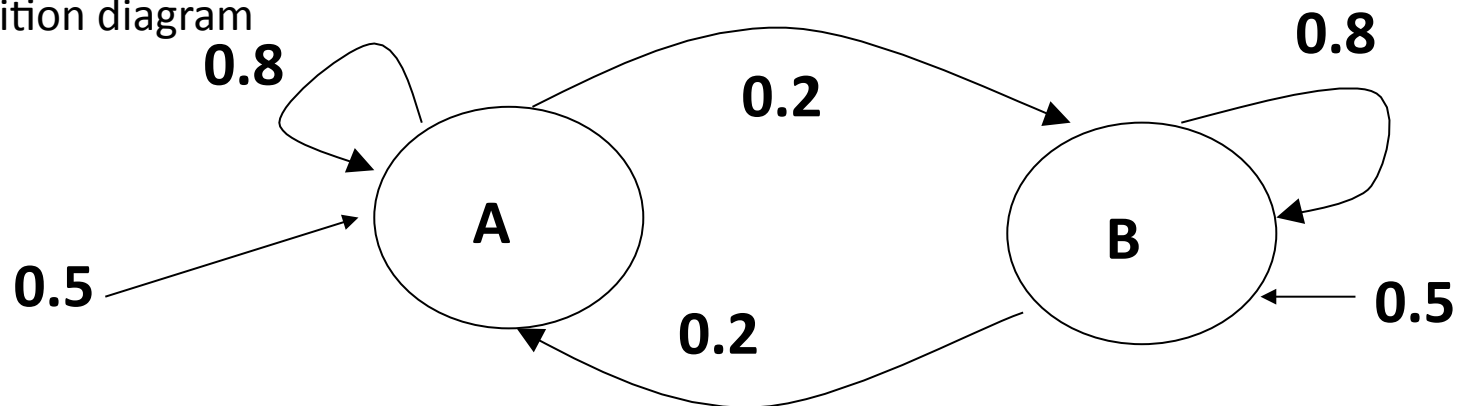
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States: A A A B B B B B A A

Observations: 1 2 2 1 1 2 1 1 1 2 2

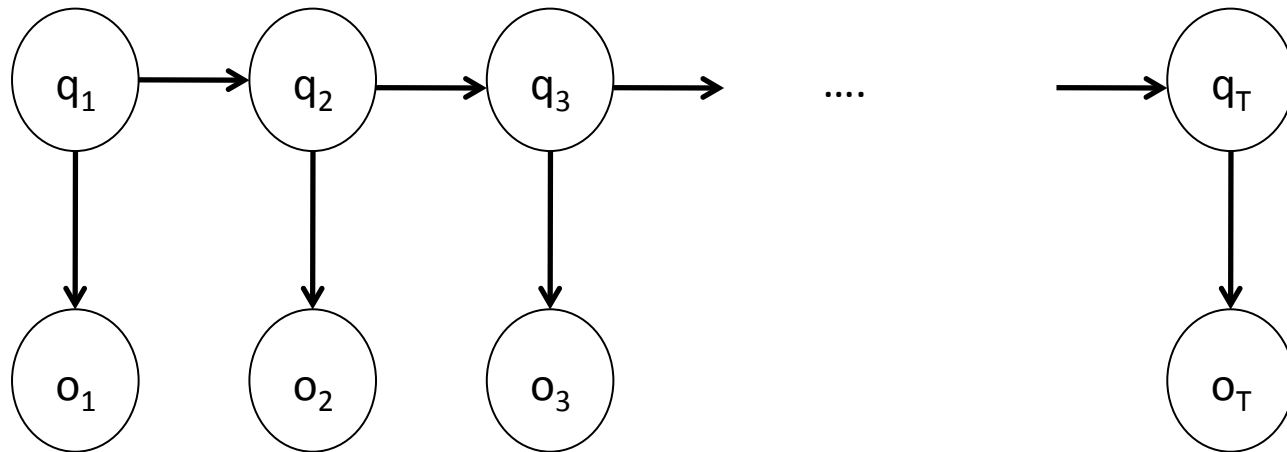
State transition diagram



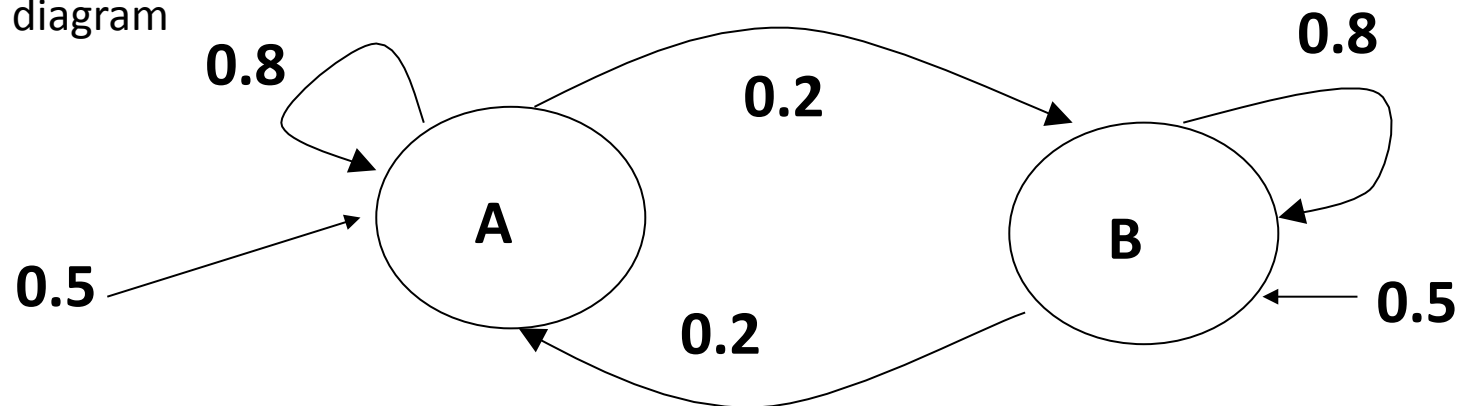
A Hidden Markov Model

- Probabilistic graphical models

States: A A A B B B B B B A A
Observations: 1 2 2 1 1 2 1 1 1 2 2



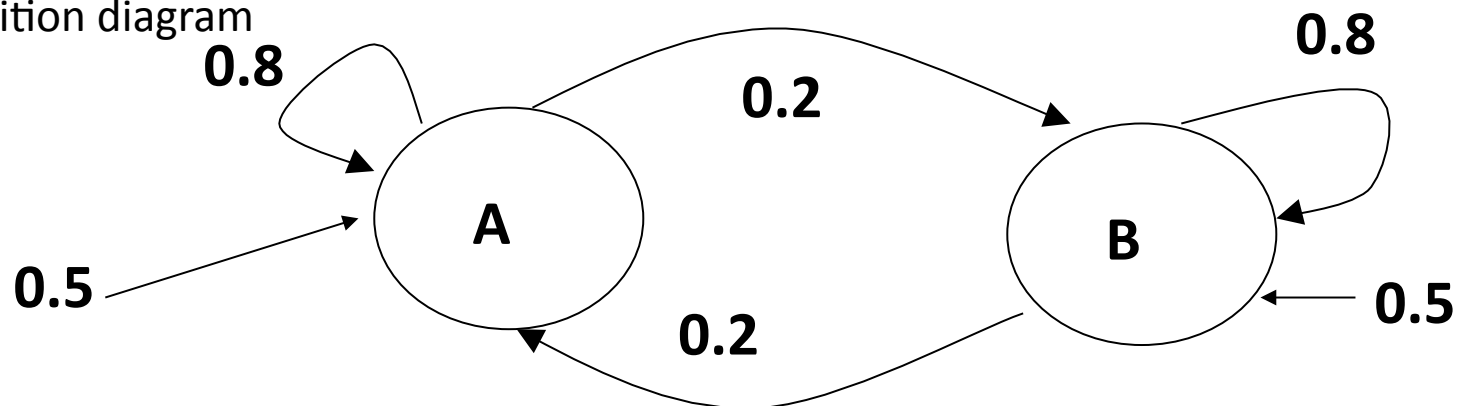
State transition diagram



A Hidden Markov Model

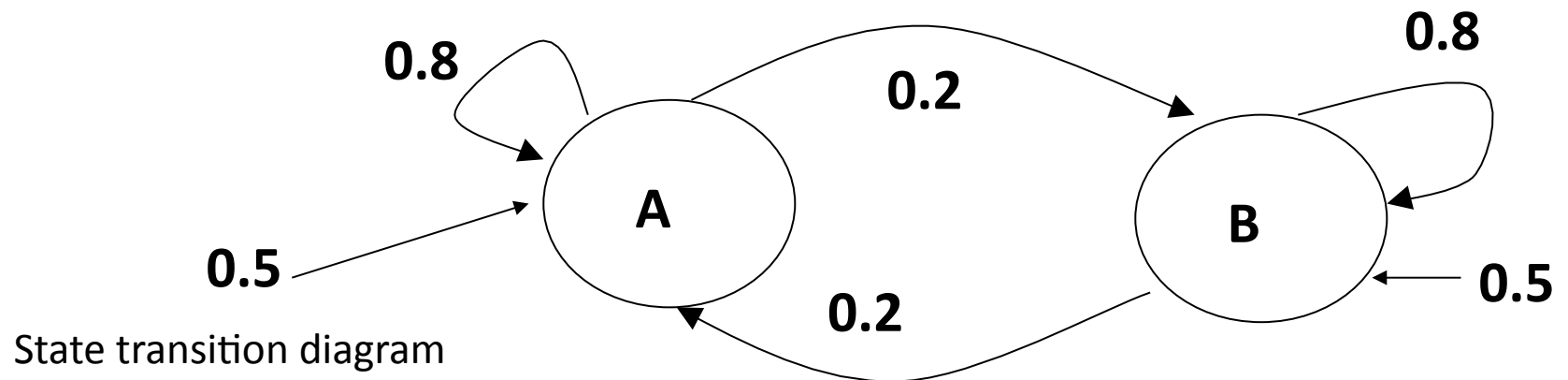
- A set of states $S=\{s_1 \dots s_n\}$
 - In each time point we are in exactly one of these states
- A set of possible outputs $\Sigma=\{\sigma_1, \dots, \sigma_m\}$
 - In each time point we emit a symbol $\sigma_j \in \Sigma$
- Random variables
 - States at each time point $Q = \{q_1, \dots, q_T\}$
 - Each q_t can take on values from $\{s_1 \dots s_n\}$
 - Outputs at each time point $O = \{o_1, \dots, o_T\}$
 - Each o_t can take on values from Σ

State transition diagram



A Hidden Markov Model

- Parameters of the model
 - $\Pi_i = \{\pi_1, \dots, \pi_n\}$: initial state probabilities $P(q_1=s_i)$
 - the probability that we *start* at state s_i , $i=1,\dots,n$
 - A transition probability model, $P(q_t = s_i \mid q_{t-1} = s_j)$
 - $n \times n$ matrix of transition probabilities
 - An emission probability model, $p(o_t = \sigma_j \mid q_t = s_i)$
 - $n \times m$ matrix of emission probabilities

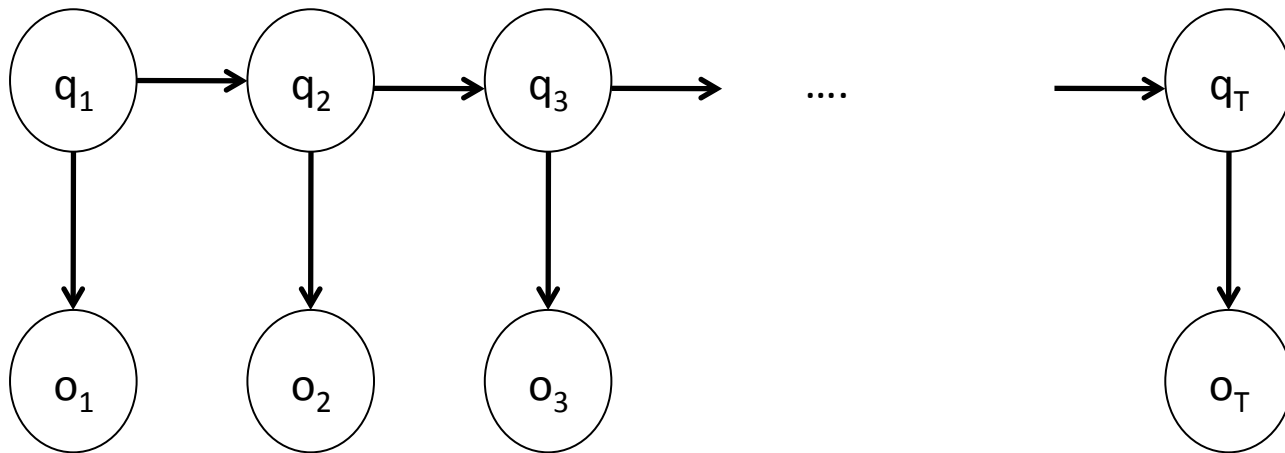


A Hidden Markov Model

- The joint probability of (Q,O) is defined as

$$P(Q,O) = p(q_1) \prod_{t=1}^T p(q_t | q_{t-1}) p(o_t | q_t)$$

Initial probability transition probability emission probability

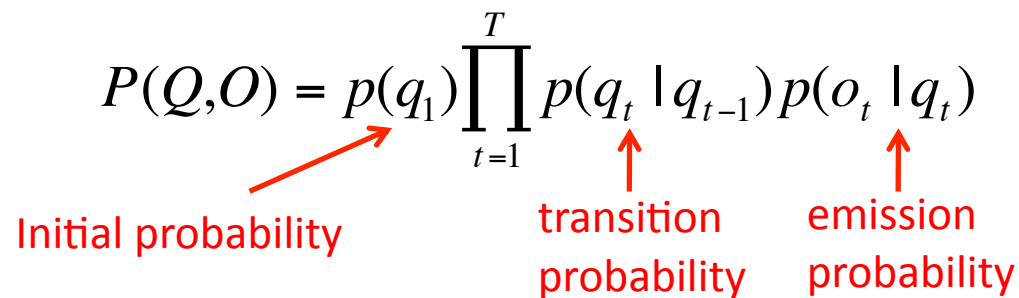


A Hidden Markov Model

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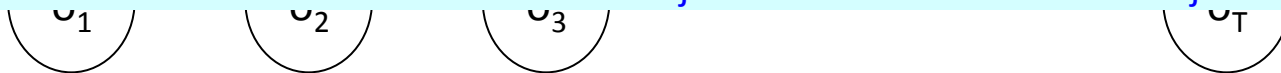
$$P(Q,O) = p(q_1) \prod_{t=1}^T p(q_t | q_{t-1}) p(o_t | q_t)$$

Initial probability transition probability emission probability



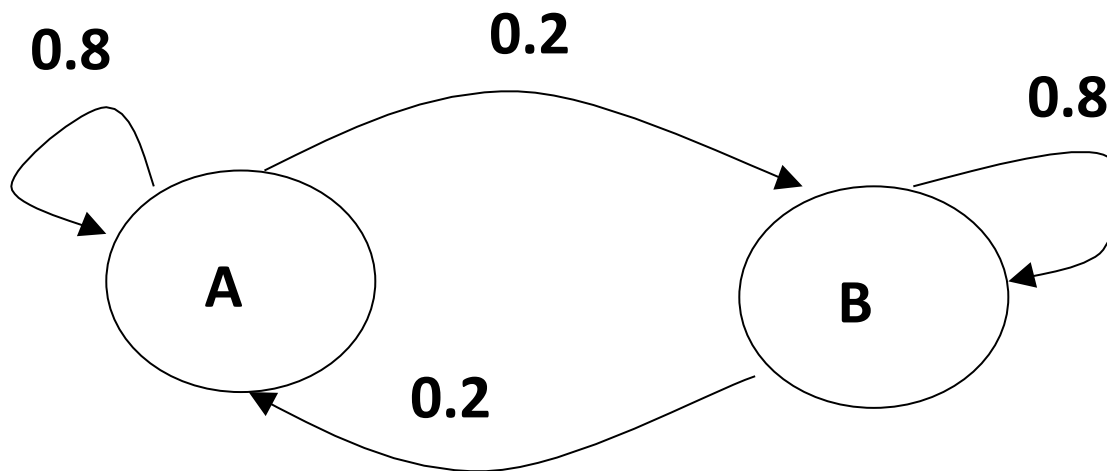
An important aspect of this definitions is the Markov property:
 q_{t+1} is conditionally independent of q_{t-1} (and any earlier time points)
given q_t

More formally $P(q_{t+1} = s_i | q_t = s_j) = P(q_{t+1} = s_i | q_t = s_j, q_{t-1} = s_j)$



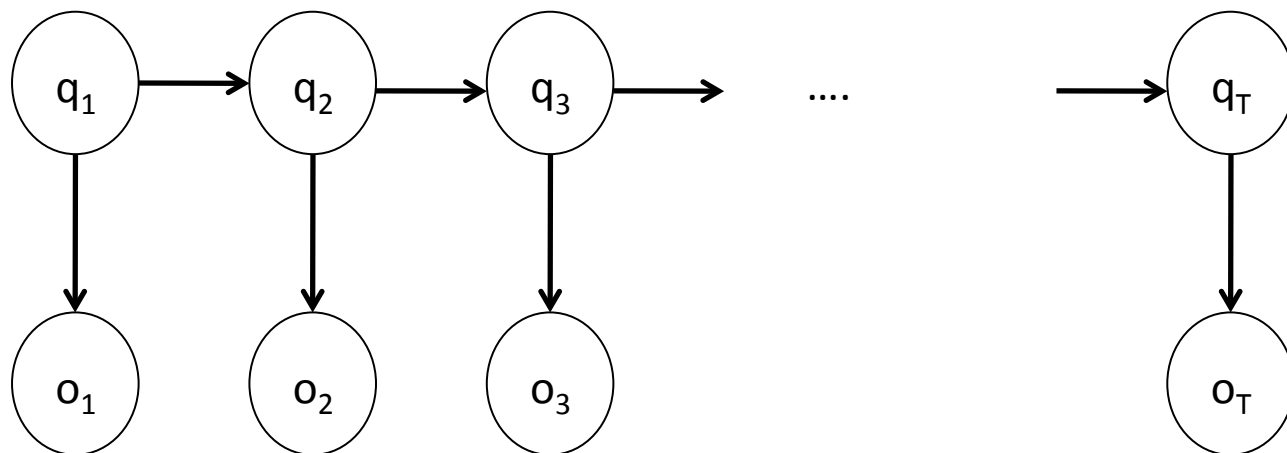
What can we ask when using a HMM?

- A few examples:
 - “Which die is currently being used?”
 - “What is the probability of a 6 in the next role?”
 - “What is the probability of 6 in any of the next 3 roles?”



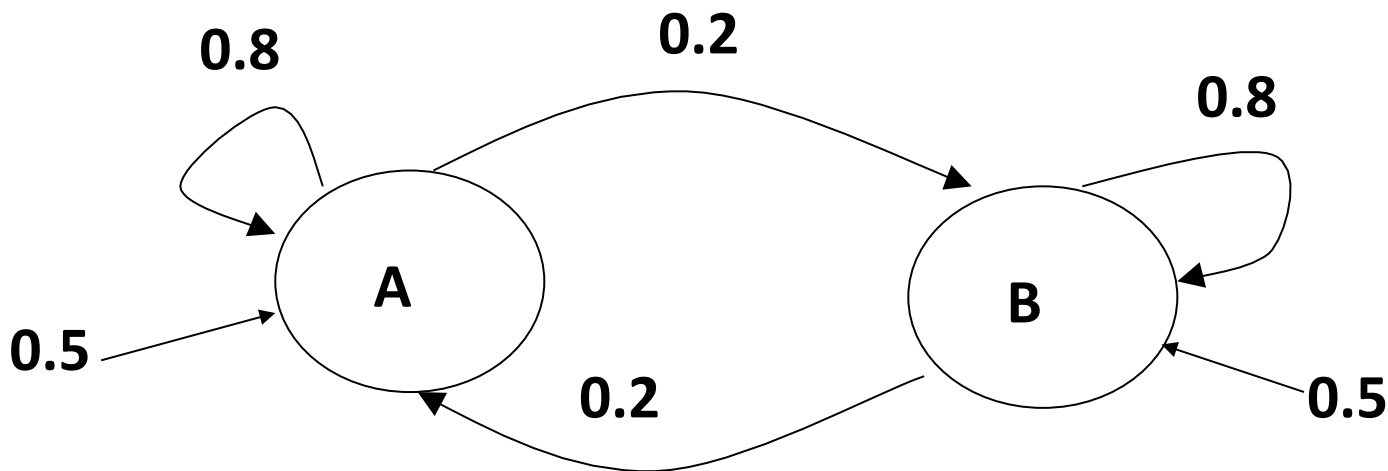
Inference in HMMs

- Computing $P(Q)$ and $P(q_t = s_i)$
 - If we cannot look at observations
- Computing $P(Q | O)$ and $P(q_t = s_i | O)$
 - When we have observation and care about the last state only
- Computing $\text{argmax}_Q P(Q | O)$
 - When we care about the entire path



Which die is currently being used?

- We played t rounds so far
- We want to determine $P(q_t = A)$
- Let's assume for now that we cannot observe any outputs (we are blind folded)
- How can we compute this?



$P(q_t = A)?$

- Simple answer: Consider “all” paths that end in A. For each such path Q, let's determine $P(Q)$

$$Q = q_1, \dots, q_{t-1}, A$$

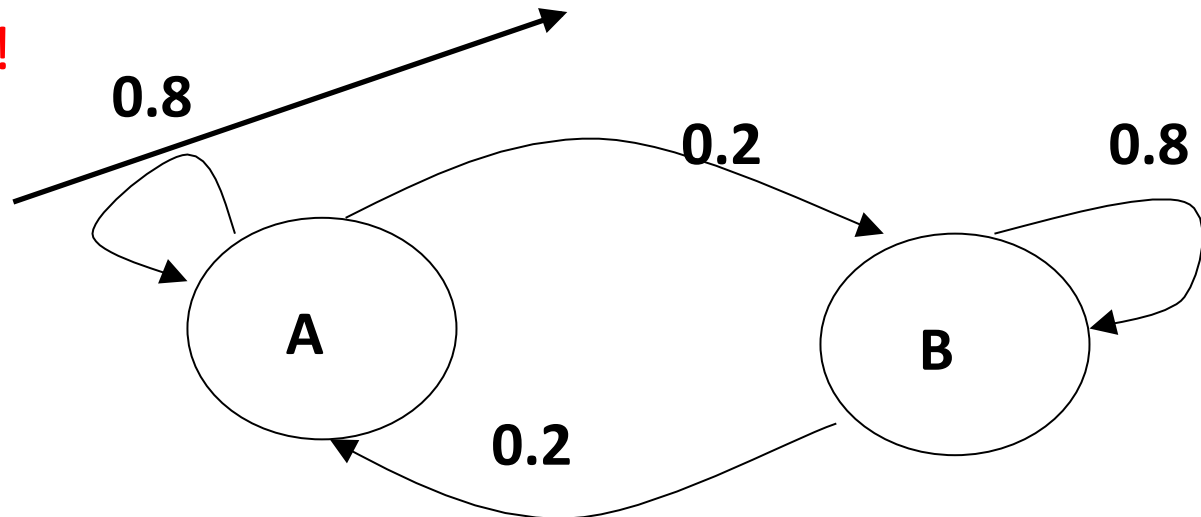
$$P(Q) = P(q_1, \dots, q_{t-1}, A) = P(A \mid q_1, \dots, q_{t-1}) P(q_1, \dots, q_{t-1})$$

$$= P(A \mid q_{t-1}) P(q_1, \dots, q_{t-1})$$

$$= P(A \mid q_{t-1}) \dots P(q_2 \mid q_1) P(q_1)$$

Markov property!

Initial probability



$P(q_t = A)?$

- Simple answer:

1. Let's determine $P(Q)$ where Q is any path that ends in A

$$Q = q_1, \dots, q_{t-1}, A$$

$$P(Q) = P(q_1, \dots, q_{t-1}, A)$$

$$= P(A \mid q_1, \dots, q_{t-1}) P(q_1, \dots, q_{t-1})$$

$$= P(A \mid q_{t-1}) P(q_1, \dots, q_{t-1})$$

$$= P(A \mid q_{t-1}) \dots P(q_2 \mid q_1) P(q_1)$$

2. $P(q_t = A) = \sum P(Q)$

where the sum is over all sets of t states
that end in A

$P(q_t = A)?$

- Simple answer:

1. Let's determine $P(Q)$ where Q is any path that ends in A

$$Q = q_1, \dots, q_{t-1}, A$$

$$P(Q) = P(q_1, \dots, q_{t-1}, A)$$

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2. $P(q_t = A) = \sum P(Q)$

where the sum is over all sets of t states
that end in A

Q: How many sets Q are
there?

A: A lot! (2^{t-1})

Not a feasible solution

$P(q_t = A)$, the smart way

- Let's define $p_t(i)$ as the probability of being in state i at time t : $p_t(i) = p(q_t = s_i)$
- We can determine $p_t(i)$ by induction
 1. $p_1(i) = \Pi_i$
 2. $p_t(i) = ?$

$P(q_t = A)$, the smart way

- Let's define $p_t(i)$ = probability state i at time t = $p(q_t = s_i)$
- We can determine $p_t(i)$ by induction
 1. $p_1(i) = \Pi_i$
 2. $p_t(i) = \sum_j p(q_t = s_i \mid q_{t-1} = s_j) p_{t-1}(j)$

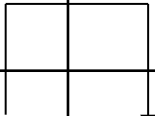
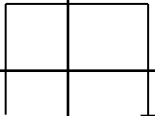

$P(q_t = A)$, the smart way

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This type of computation is called
dynamic programming

Complexity: $O(n^2 * t)$

Number of states in our HMM

Time / state	t1	t2	t3
s1	.3		
s2	.7		

Inference in HMMs

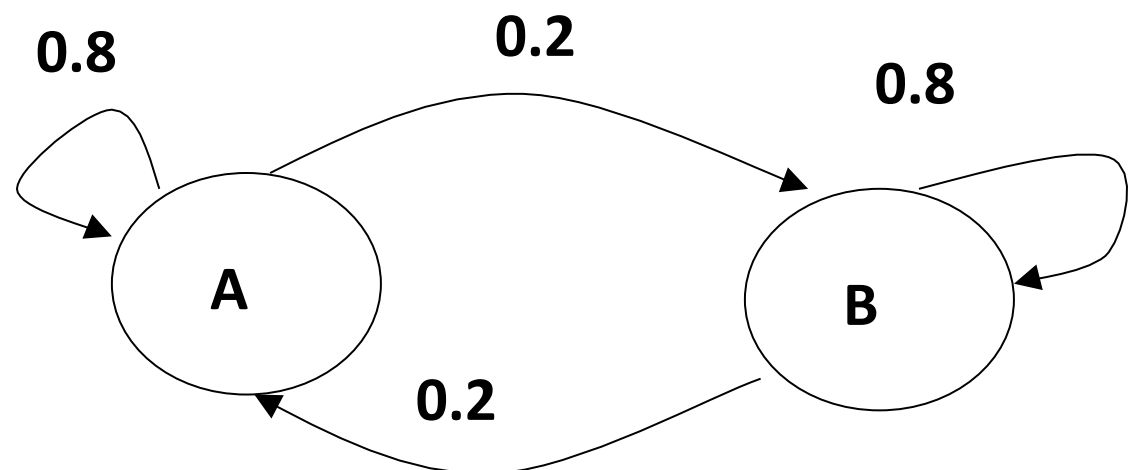
- Computing $P(Q)$ and $P(q_t = s_i)$ ✓
- Computing $P(Q \mid O)$ and $P(q_t = s_i \mid O)$
- Computing $\operatorname{argmax}_Q P(Q)$

But what if we observe outputs?

- So far, we assumed that we could not observe the outputs
- In reality, we almost always can.



v	$P(v A)$	$P(v B)$
1	.3	.1
2	.2	.1
3	.2	.1
4	.1	.2
5	.1	.2
6	.1	.3



But what if we observe outputs?

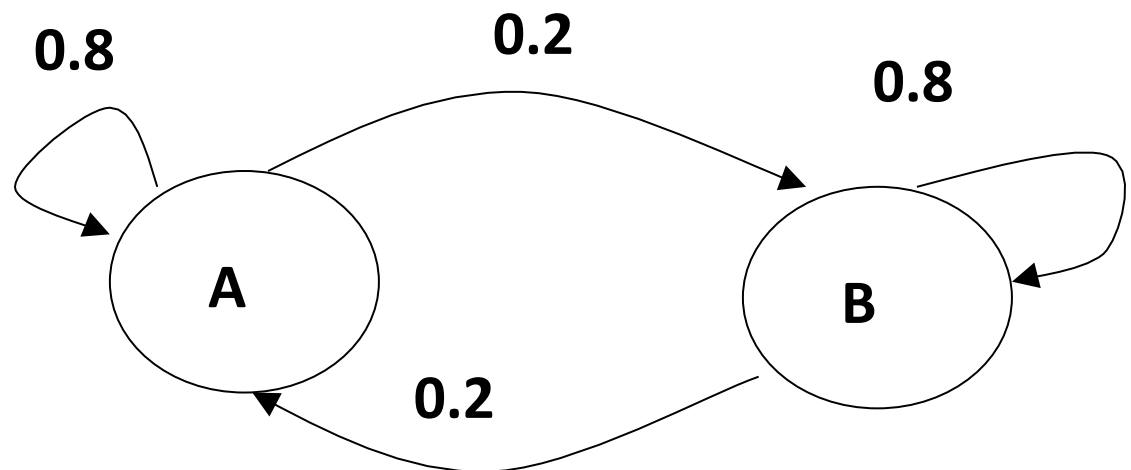
- So far, we assumed that we could not observe the outputs
- In reality, we almost always can

Does observing the sequence

5, 6, 4, 5, 6, 6

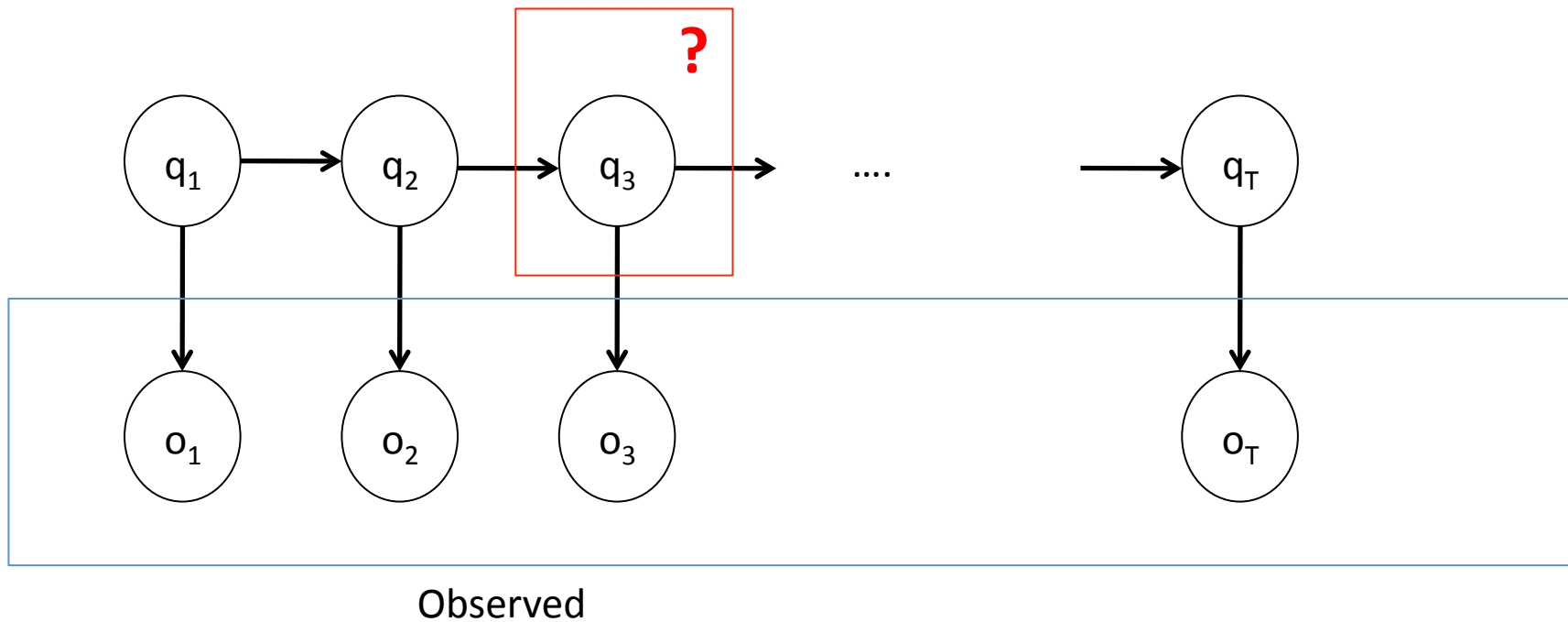
Change our belief about the state?

v	$P(v A)$	$P(v B)$
1	.3	.1
2	.2	.1
3	.2	.1
4	.1	.2
5	.1	.2
6	.1	.3



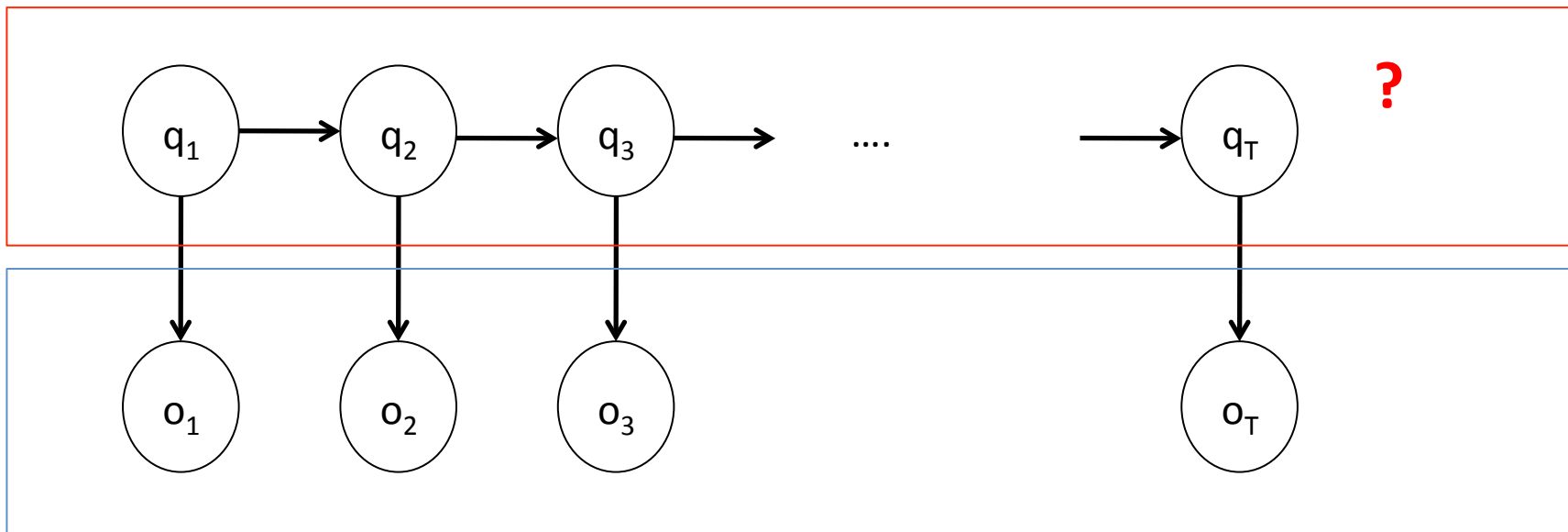
When outputs are observed

- We want to compute $P(q_t = A \mid O_1 \dots O_t)$



$P(q_t = A | O)$ when outputs are observed

- We want to compute $P(q_t = A | O_1 \dots O_t)$
- Let's start with a simpler question. Given a sequence of states Q , what is $P(Q | O_1 \dots O_t) = P(Q | O)$?
 - It is pretty simple to move from $P(Q)$ to $P(q_t = A)$



Observed

$P(q_t = A | O)$ when outputs are observed

- We want to compute $P(q_t = A | O_1 \dots O_t)$
- Let's start with a simpler question. Given a sequence of states Q , what is $P(Q | O_1 \dots O_t) = P(Q | O)$?
 - It is pretty simple to move from $P(Q)$ to $P(q_t = A)$
 - In some cases $P(Q)$ is the more important question
 - Speech processing
 - NLP

$$P(Q \mid O)$$

- We can use Bayes rule:

$$P(Q|O) = \frac{P(O|Q)P(Q)}{P(O)}$$

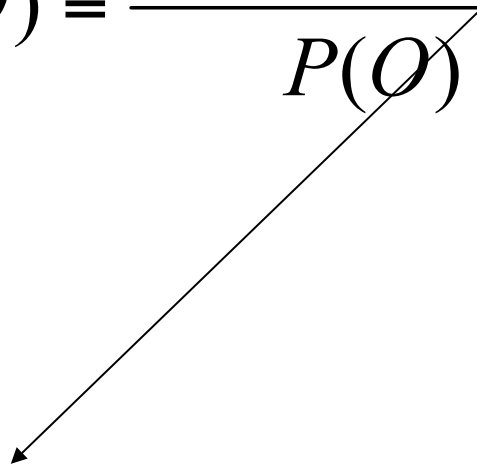


Easy, $P(O \mid Q) = P(o_1 \mid q_1) P(o_2 \mid q_2) \dots P(o_t \mid q_t)$

$$P(Q \mid O)$$

- We can use Bayes rule:

$$P(Q|O) = \frac{P(O|Q)P(Q)}{P(O)}$$

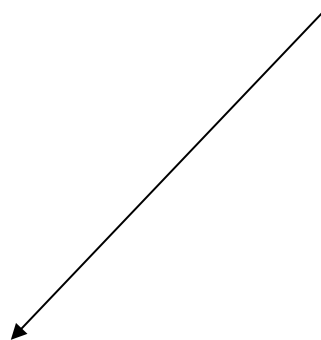


Easy, $P(Q) = P(q_1) P(q_2 \mid q_1) \dots P(q_t \mid q_{t-1})$

$$P(Q \mid O)$$

- We can use Bayes rule:

$$P(Q|O) = \frac{P(O|Q)P(Q)}{P(O)}$$



Hard!

$P(O)$

- What is the probability of seeing a set of observations:
 - An important question in it own rights, for example classification using two HMMs H_1 and H_2
 - Compute $P(O|H_1)$ and $P(O|H_2)$, classify to the model with higher probability

$P(O)$

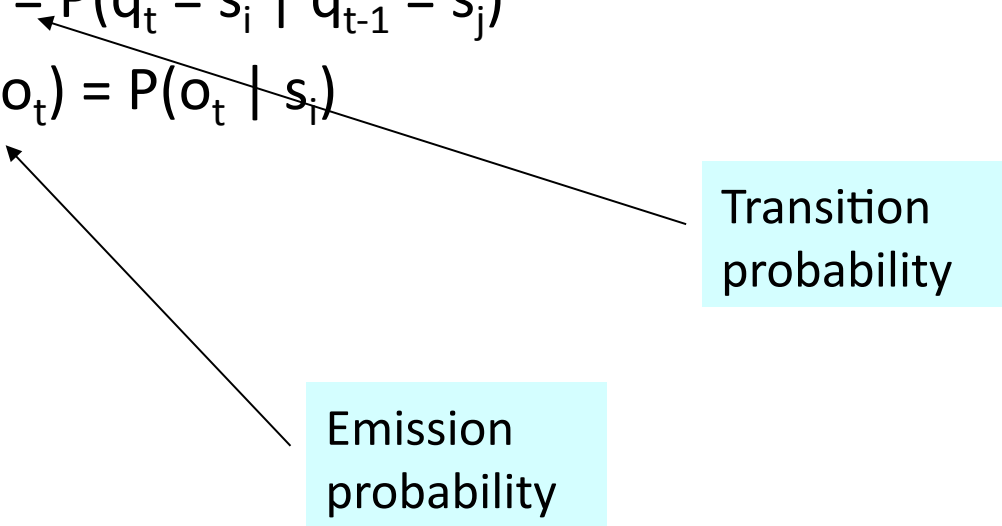
- Define $\alpha_t(i) = P(o_1, o_2, \dots, o_t \wedge q_t = s_i)$
- $\alpha_t(i)$ is the probability that we:
 1. Observe o_1, o_2, \dots, o_t
 2. End up at state i

How do we compute $\alpha_t(i)$?

When outputs are observed

- We want to compute $P(q_t = A \mid O_1 \dots O_t)$
- For ease of writing we will use the following notations (commonly used in the literature)
- $a_{j,i} = P(q_t = s_i \mid q_{t-1} = s_j)$
- $b_i(o_t) = P(o_t \mid s_i)$

Transition
probability



Emission
probability

Computing $\alpha_t(i)$

- $\alpha_1(i) = P(o_1 \wedge q_1 = i) = P(o_1 \mid q_1 = s_i)\Pi_i$

We must be at a state in time t

chain rule

Markov property

Computing $\alpha_t(i)$

- $\alpha_1(i) = P(o_1 \wedge q_1 = i) = P(o_1 \mid q_1 = s_i) \Pi_i$

$$\begin{aligned}
 \alpha_{t+1}(i) &= P(O_1 \dots O_{t+1} \wedge q_{t+1} = s_i) = && \text{We must be at a state in time } t \\
 \sum_j P(O_1 \dots O_t \wedge q_t = s_j \wedge O_{t+1} \wedge q_{t+1} = s_i) &= && \text{chain rule} \\
 \sum_j P(O_{t+1} \wedge q_{t+1} = s_i \mid O_1 \dots O_t \wedge q_t = s_j) P(O_1 \dots O_t \wedge q_t = s_j) &= \\
 \sum_j P(O_{t+1} \wedge q_{t+1} = s_i \mid O_1 \dots O_t \wedge q_t = s_j) \alpha_t(j) &= && \text{Markov property} \\
 \sum_j P(O_{t+1} \mid q_{t+1} = s_i) P(q_{t+1} = s_i \mid q_t = s_j) \alpha_t(j) &= \\
 \sum_j b_i(O_{t+1}) a_{j,i} \alpha_t(j)
 \end{aligned}$$

Example: Computing $\alpha_3(B)$

- We observed 2,3,6

$$\alpha_1(A) = P(2 \wedge q_1 = A) = P(2 \mid q_1 = A)\Pi_A = .2 * .7 = .14, \alpha_1(B) = .1 * .3 = .03$$

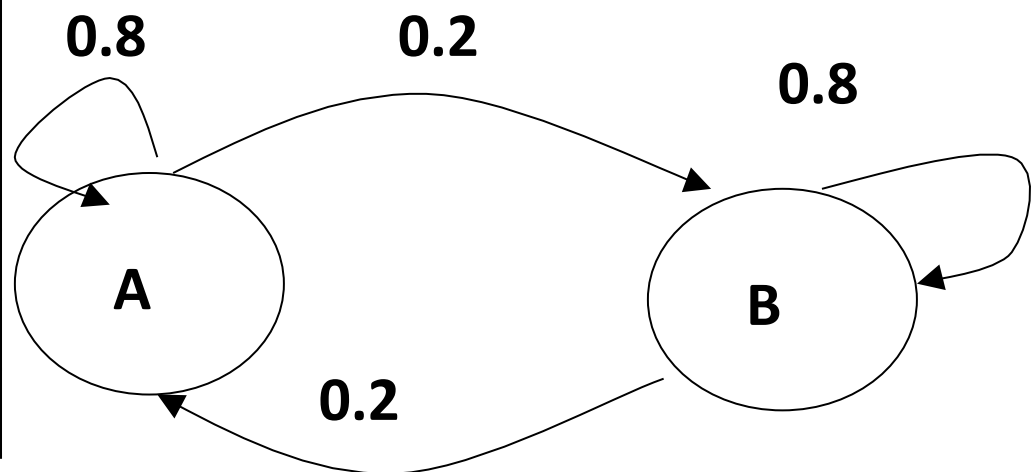
$$\alpha_2(A) = \sum_{j=A,B} b_A(3) a_{j,A} \alpha_1(j) = .2 * .8 * .14 + .2 * .2 * .03 = 0.0236, \alpha_2(B) = 0.0052$$

$$\alpha_3(B) = \sum_{j=A,B} b_B(6) a_{j,B} \alpha_2(j) = .3 * .2 * .0236 + .3 * .8 * .0052 = 0.00264$$

$$\Pi_A = 0.7$$

$$\Pi_B = 0.3$$

v	P(v A)	P(v B)
1	.3	.1
2	.2	.1
3	.2	.1
4	.1	.2
5	.1	.2
6	.1	.3



Where we are

- We want to compute $P(Q \mid O)$
- For this, we only need to compute $P(O)$
- We know how to compute $\alpha_t(i)$

From now its easy

$$\alpha_t(i) = P(o_1, o_2 \dots, o_t \wedge q_t = s_i)$$

so

$$P(O) = P(o_1, o_2 \dots, o_t) = \sum_i P(o_1, o_2 \dots, o_t \wedge q_t = s_i) = \sum_i \alpha_t(i)$$

note that

$$p(q_t=s_i \mid o_1, o_2 \dots, o_t) = \frac{\alpha_t(i)}{\sum_j \alpha_t(j)}$$

$$P(A \mid B) = P(A \wedge B) / P(B)$$

Complexity

- How long does it take to compute $P(Q \mid O)$?
 - $P(Q)$: $O(n)$
 - $P(O \mid Q)$: $O(n)$
 - $P(O)$: $O(n^2t)$

Inference in HMMs

- Computing $P(Q)$ and $P(q_t = s_i)$ ✓
- Computing $P(Q \mid O)$ and $P(q_t = s_i \mid O)$ ✓
- Computing $\operatorname{argmax}_Q P(Q)$

Most probable path

- We are almost done ...
- One final question remains

How do we find the most probable path, that is Q^* such that

$$P(Q^* | O) = \operatorname{argmax}_Q P(Q | O)?$$

- This is an important path
 - The words in speech processing
 - The set of genes in the genome
 - etc.

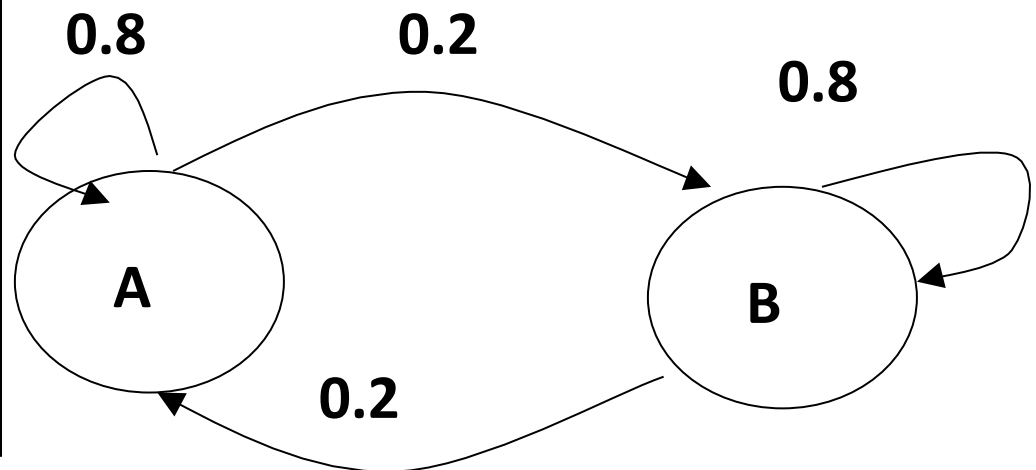
Example

- What is the most probable set of states leading to the sequence:

1,2,2,5,6,5,1,2,3 ?

$$\Pi_A = 0.7$$
$$\Pi_B = 0.3$$

v	P(v A)	P(v B)
1	.3	.1
2	.2	.1
3	.2	.1
4	.1	.2
5	.1	.2
6	.1	.3



Most probable path

$$\begin{aligned}\arg \max_Q P(Q | O) &= \arg \max_Q \frac{P(O | Q)P(Q)}{P(O)} \\ &= \arg \max_Q P(O | Q)P(Q)\end{aligned}$$

We will use the following definition:

$$\delta_t(i) = \max_{q_1 \dots q_{t-1}} p(q_1 \dots q_{t-1} \wedge q_t = s_i \wedge O_1 \dots O_t)$$

In other words we are interested in the most likely path from 1 to t that:

1. Ends in S_i
2. Produces outputs $O_1 \dots O_t$

Computing $\delta_t(i)$

$$\delta_t(i) = \max_{q_1 \dots q_{t-1}} p(q_1 \dots q_{t-1} \wedge q_t = s_i \wedge O_1 \dots O_t)$$

Initialization at $t=1$

$$\begin{aligned}\delta_1(i) &= p(q_1 = s_i \wedge O_1) \\ &= p(q_1 = s_i) p(O_1 | q_1 = s_i) \\ &= \pi_i b_i(O_1)\end{aligned}$$

Q: Given $\delta_t(i)$, how can we compute $\delta_{t+1}(i)$?

A: To get from $\delta_t(i)$ to $\delta_{t+1}(i)$ we need to

1. Add an emission for time $t+1$ (O_{t+1})
2. Transition to state s_i

$$\begin{aligned}\delta_{t+1}(i) &= \max_{q_1 \dots q_t} p(q_1 \dots q_t \wedge q_{t+1} = s_i \wedge O_1 \dots O_{t+1}) \\ &= \max_j \delta_t(j) p(q_{t+1} = s_i | q_t = s_j) p(O_{t+1} | q_{t+1} = s_i) \\ &= \max_j \delta_t(j) a_{j,i} b_i(O_{t+1})\end{aligned}$$

The Viterbi algorithm

$$\begin{aligned}\delta_{t+1}(i) &= \max_{q_1 \dots q_t} p(q_1 \dots q_t \wedge q_{t+1} = s_i \wedge O_1 \dots O_{t+1}) \\ &= \max_j \delta_t(j) p(q_{t+1} = s_i \mid q_t = s_j) p(O_{t+1} \mid q_{t+1} = s_i) \\ &= \max_j \delta_t(j) a_{j,i} b_i(O_{t+1})\end{aligned}$$

- Once again we use dynamic programming for solving $\delta_t(i)$
- Once we have $\delta_t(i)$, we can solve for our $P(Q^* \mid O)$ by:

$$P(Q^* \mid O) = \operatorname{argmax}_Q P(Q \mid O) =$$

path defined by $\operatorname{argmax}_j \delta_t(j)$,

Inference in HMMs

- Computing $P(Q)$ and $P(q_t = s_i)$ ✓
- Computing $P(Q \mid O)$ and $P(q_t = s_i \mid O)$ ✓
- Computing $\operatorname{argmax}_Q P(Q)$ ✓

What you should know

- Why HMMs? Which applications are suitable?
- Inference in HMMs
 - No observations
 - Probability of next state w. observations
 - Maximum scoring path (Viterbi)