# Precept 4: HMMs, MEMMs, CRFs

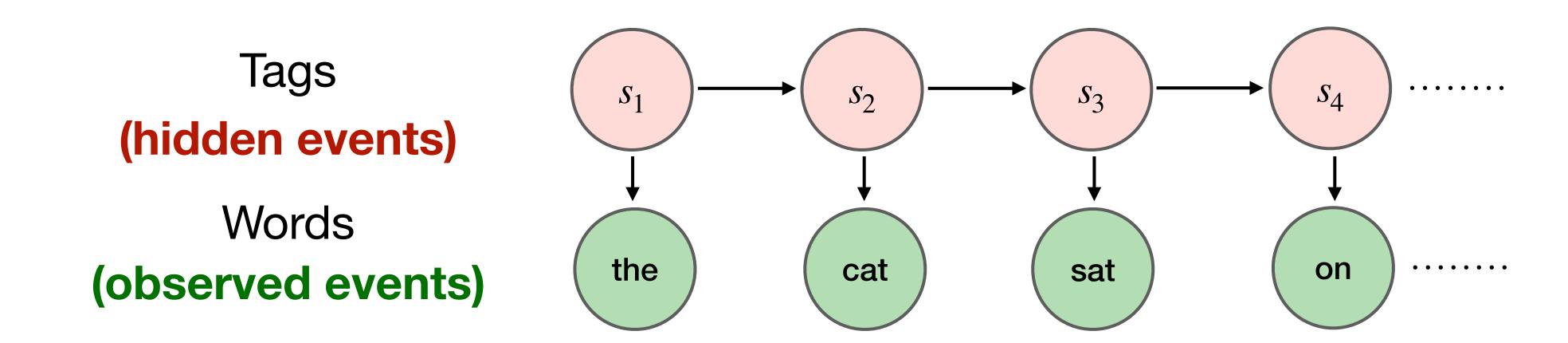
COS 484

Tyler Zhu (Viterbi figures adapted from Howard Chen, lecture)

### Today's Plan

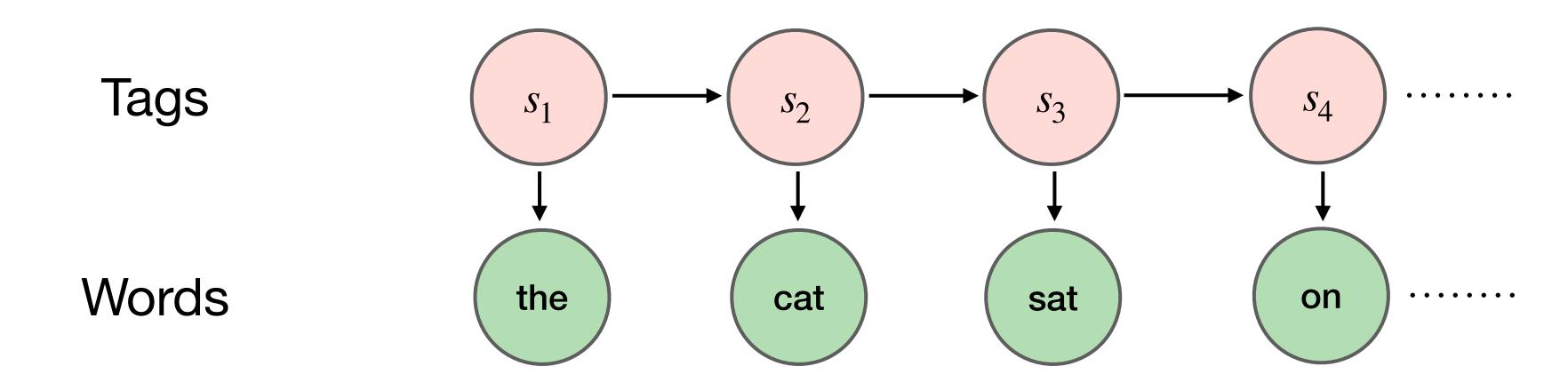
- 1. Hidden Markov Models (20 min)
- 2. Maximum Entropy Markov Models (20 min)
- 3. Conditional Random Fields (5 min)

### Hidden Markov Model (HMM)



- We don't normally see sequences of POS tags in text
- However, we do observe the words!
- The HMM allows us to jointly reason over both hidden and observed events.
  - Assume that each position has a tag that generates a word

### HMMs: Assumptions



What are the key assumptions?

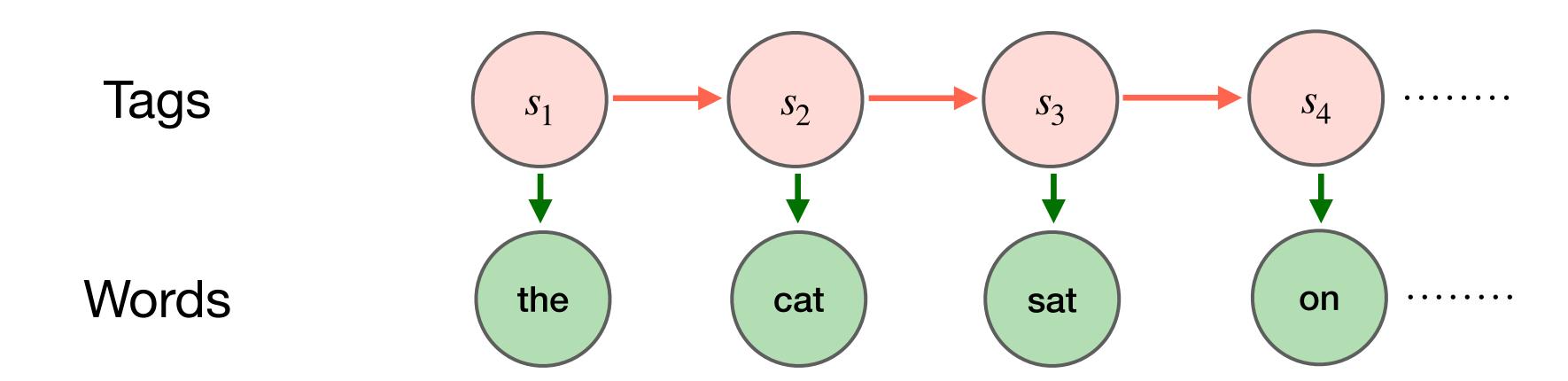
1. Markov assumption:

$$P(s_t | s_1, \dots, s_{t-1}) \approx P(s_t | s_{t-1})$$

2. Output independence:

$$P(o_t | s_1, \dots, s_t) \approx P(o_t | s_t)$$

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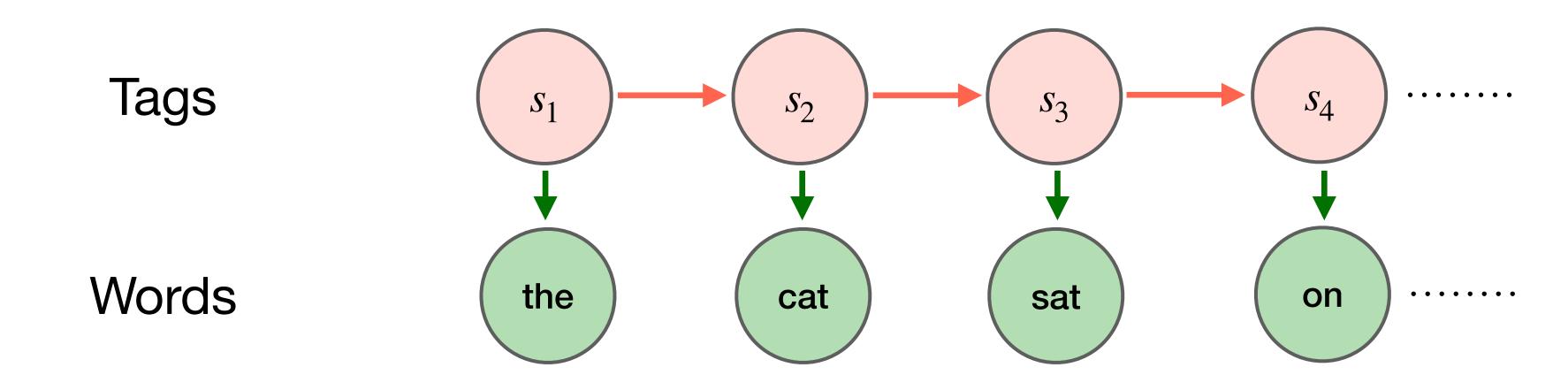
**Transition probabilities** 

2. Output independence:

$$P(o_t | s_1, \dots, s_t) \approx P(o_t | s_t)$$

**Emission probabilities** 

### HMMs: Training



What do we train and how?

#### **Transition probabilities**

$$P(s_t | s_1, \dots, s_{t-1}) \approx P(s_t | s_{t-1})$$

#### **Emission probabilities**

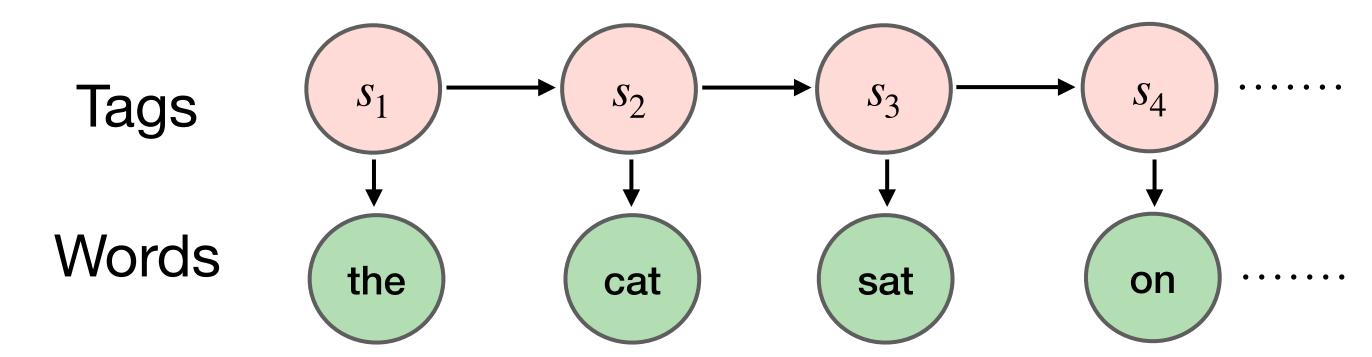
$$P(o_t | s_1, \dots, s_t) \approx P(o_t | s_t)$$

	DT	NN	VB
Ø	0.5	0.3	0.2
DT	0.1	0.5	0.4
NN	0.2	0.3	0.5
VB	0.4	0.3	0.3

From training data!

	the	cat	runs
DT	0.4	0.5	0.1
NN	0.5	0.4	0.1
VB	0.2	0.3	0.5

#### HMMs: Inference



**Task**: Find the most probable sequence of states  $S = s_1, s_2, \ldots, s_n$  given the observations  $O = o_1, o_2, \dots, o_n$ 

$$\hat{S} = \arg \max_{S} P(S \mid O) = \arg \max_{S} \frac{P(O \mid S)P(S)}{P(O)}$$
 [Bayes' rule]

= 
$$\underset{S}{\operatorname{arg}} \max_{S} P(O | S)P(S)$$
 [ $P(O)$  doesn't depend on  $S$ !]

How can we maximize this? Search over all state sequences?

= arg 
$$\max_{s_1, s_2, \dots s_n} \prod_{i=1}^n P(o_i | s_i) P(s_i | s_{i-1})$$
 [Markov assumption]

You are building a lie detector taking as input a stream of recorded behaviors:

•  $x_t \in \{a, b\}$ , i.e., face touching (a) or blinking (b)

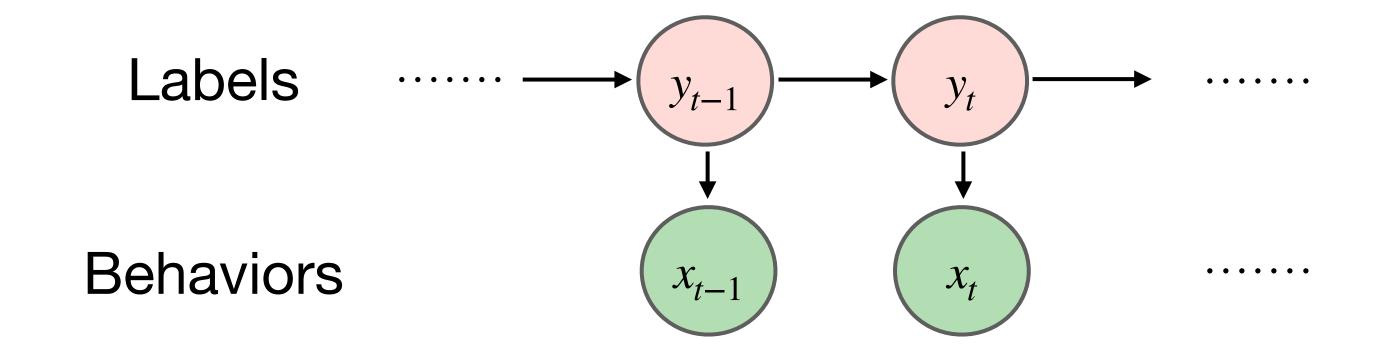
Detector at every moment then predicts one of four labels:

•  $y_t \in \{N, U, L, H\}$ , i.e., Neutral (N), Unclear (U), Lying (L), and Honest (H)

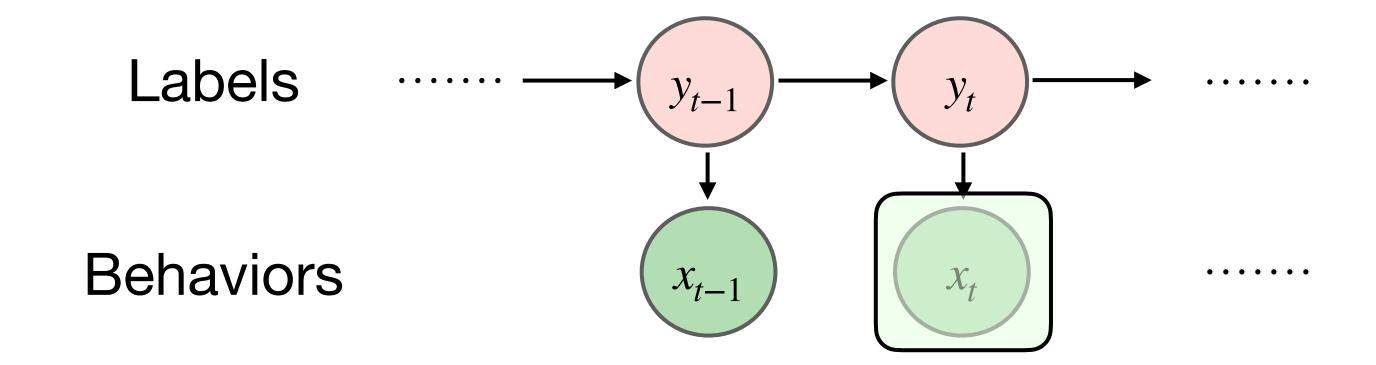
Dataset is triplets  $(y_{t-1}, y_t, x_t)$ : 9x of (N, L, a), 9x of (U, L, b), 1x of (N, H, b)

• Labels  $y_t$  come from body language experts' annotation

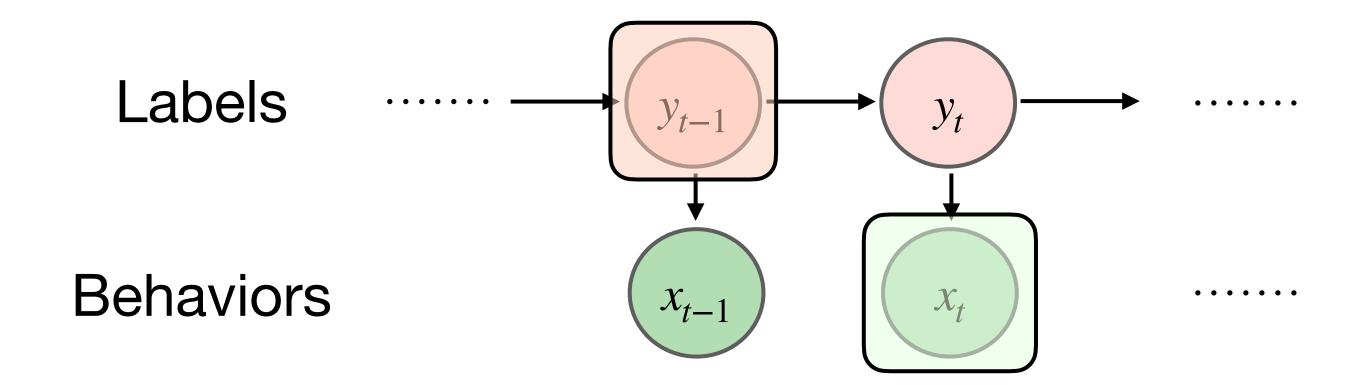
**Q1:** You use the above data to build an HMM. What is  $P(x_t = b \mid y_{t-1} = N)$ ?



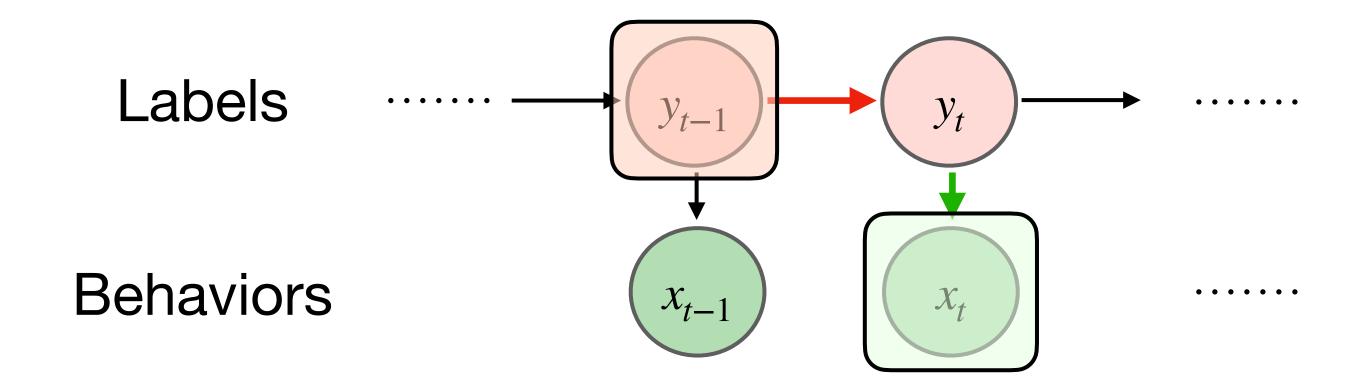
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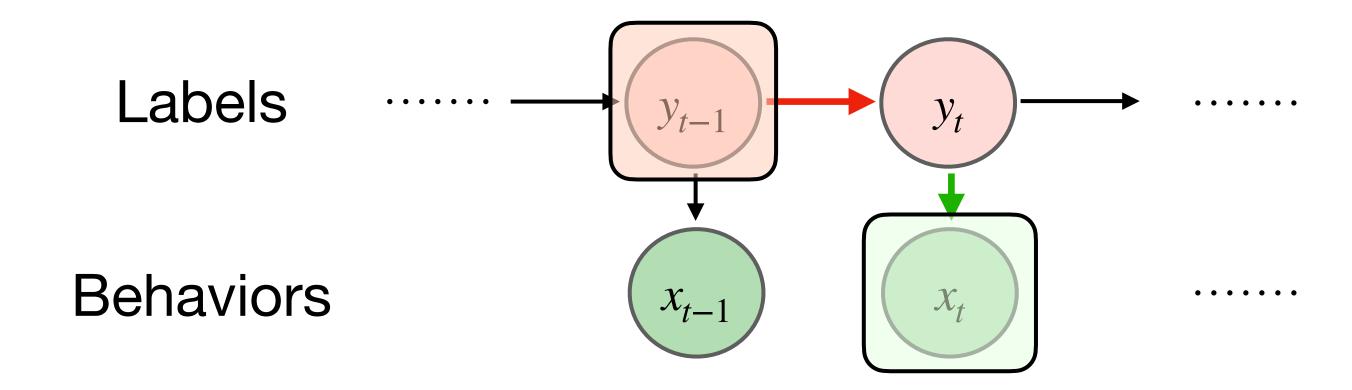
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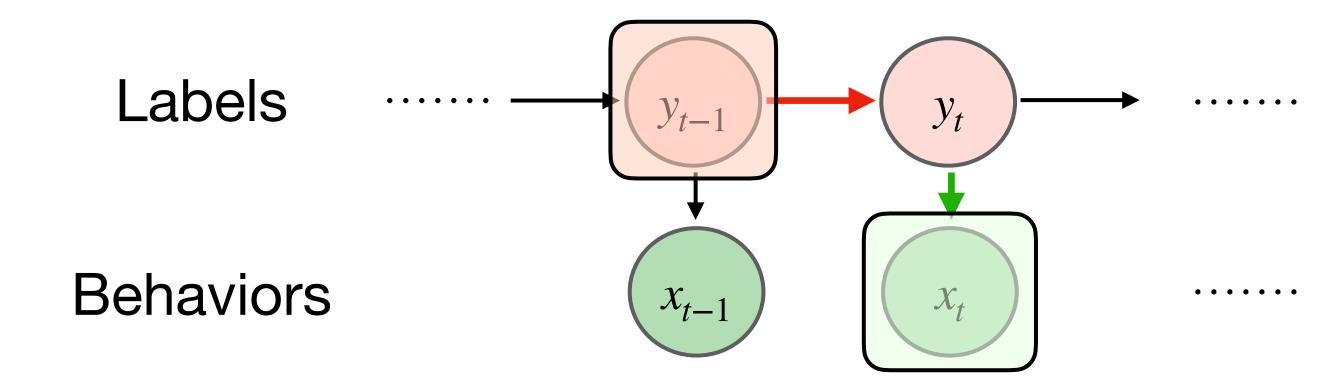


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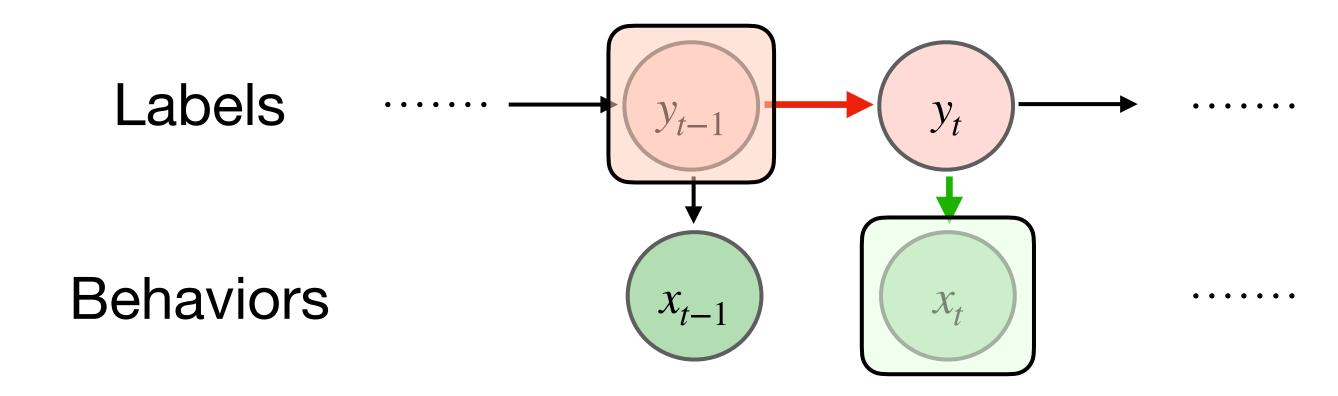


Simplified model of the HMM we care about.

We need to condition over  $y_t$  using our emission and transition probabilities.

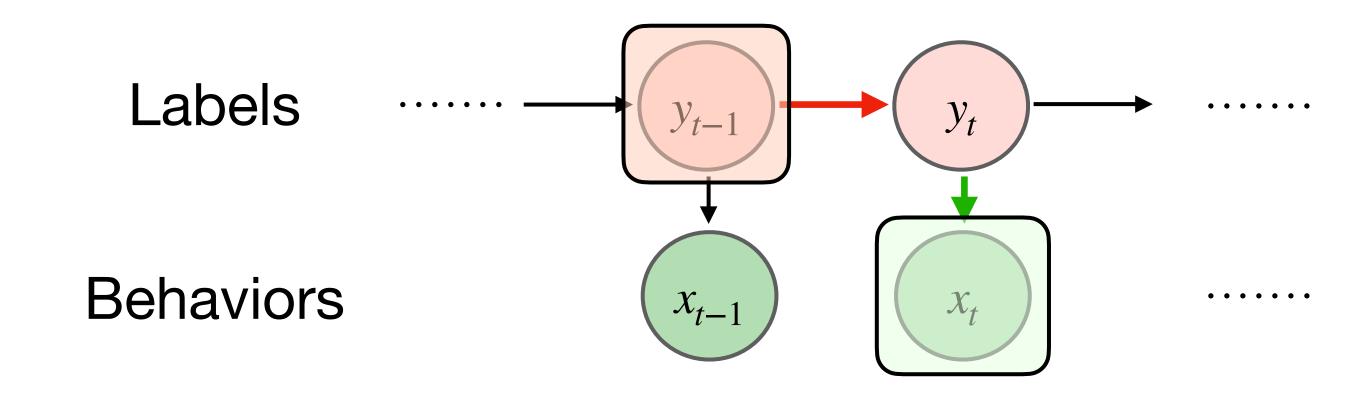


$$P(x_t = b | y_{t-1} = N) = \sum_{y_t} P(x_t = b, y_t | y_{t-1} = N)$$



$$P(x_{t} = b | y_{t-1} = N) = \sum_{y_{t}} P(x_{t} = b, y_{t} | y_{t-1} = N)$$

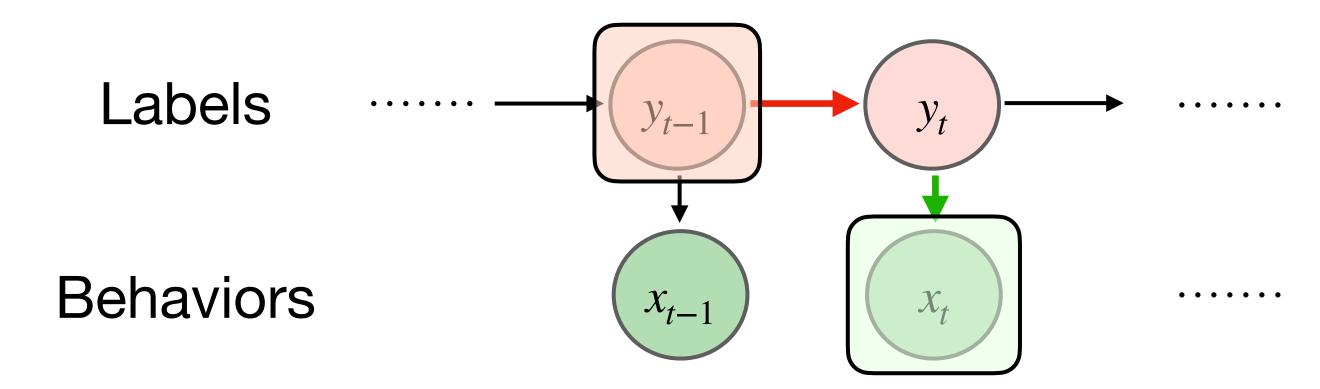
$$= \sum_{y_{t}} P(x_{t} = b | y_{t}) P(y_{t} | y_{t-1} = N)$$
emission transition



$$P(x_{t} = b | y_{t-1} = N) = \sum_{y_{t}} P(x_{t} = b, y_{t} | y_{t-1} = N)$$

$$= \sum_{y_{t}} P(x_{t} = b | y_{t}) P(y_{t} | y_{t-1} = N)$$

$$= P(b | L) P(L | N) + P(b | H) P(H | N)$$



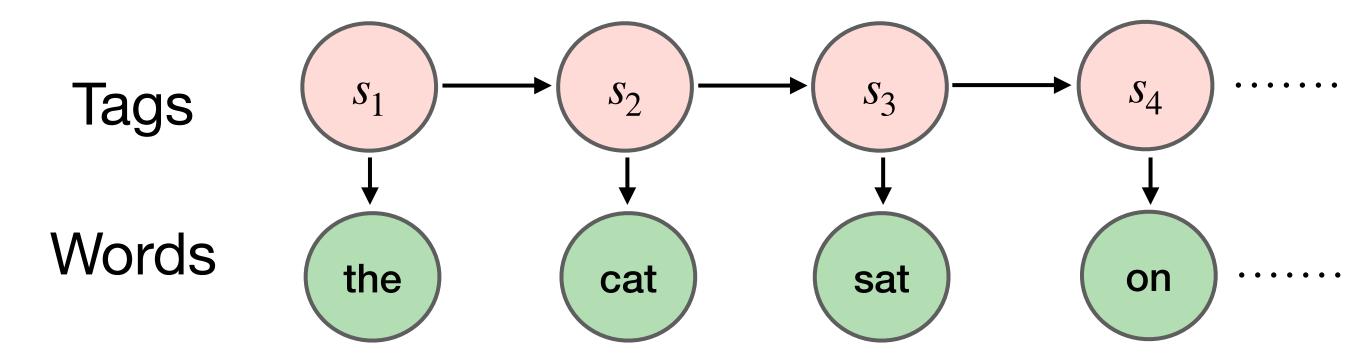
$$P(x_t = b | y_{t-1} = N) = \sum_{y_t} P(x_t = b, y_t | y_{t-1} = N)$$

$$= \sum_{y_t} P(x_t = b | y_t) P(y_t | y_{t-1} = N)$$

$$= P(b | L) P(L | N) + P(b | H) P(H | N)$$

$$= \frac{1}{2} \times \frac{9}{10} + 1 \times \frac{1}{10} = \frac{11}{20}$$

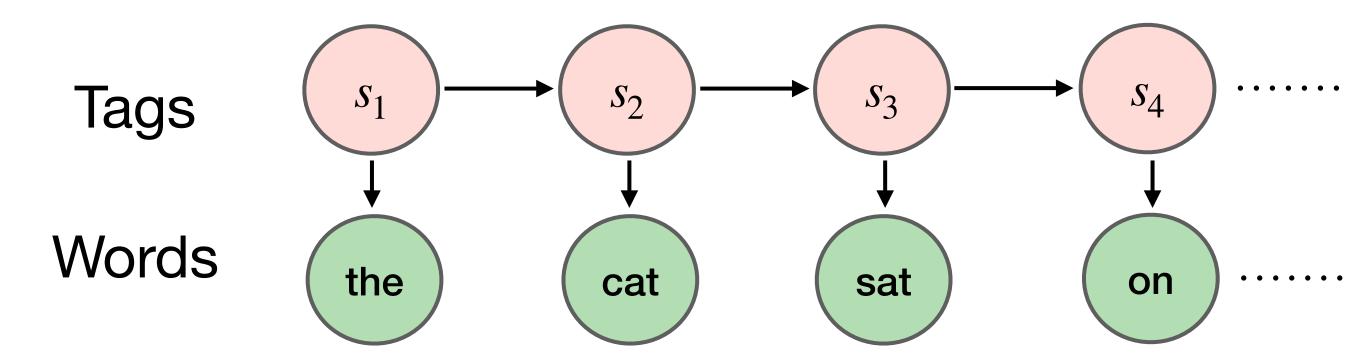
#### HMMs: Efficient Inference



**Task**: Find the most probable sequence of states  $S=s_1,s_2,\ldots,s_n$  given the observations  $O=o_1,o_2,\ldots,o_n$ 

$$\hat{S} = \arg \max_{S} P(S \mid O) = \arg \max_{s_1, s_2, \dots s_n} \prod_{i=1}^{n} P(o_i \mid s_i) P(s_i \mid s_{i-1})$$

#### HMMs: Efficient Inference



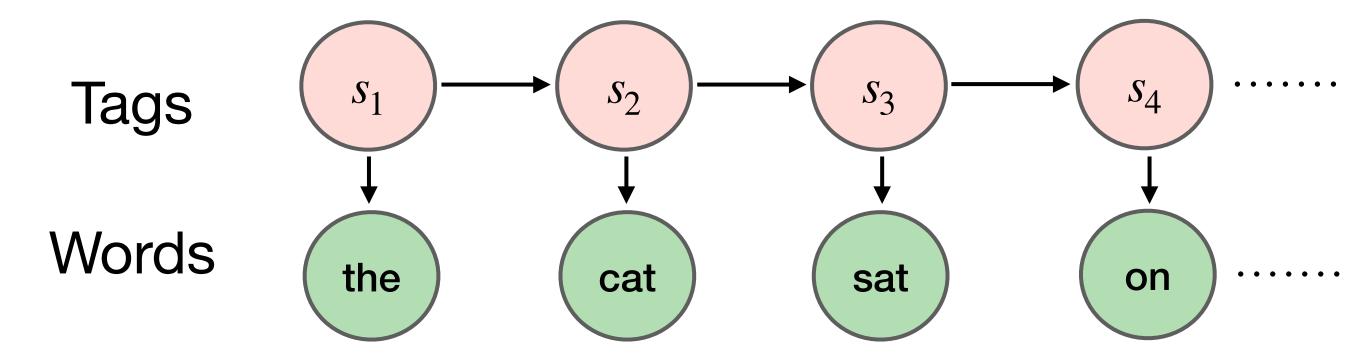
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Do what we just did for the example, but for every possible state!

- Viterbi is simply realizing that we only need to do this one-step calculation
- Try all possible explanations  $s_i$  for  $o_i$  to find the most likely one
- Multiply by (highest)  $P(s_{i-1})$ , i.e. score, so "most likely" is over all observations

#### HMMs: Efficient Inference



**Task**: Find the most probable sequence of states  $S = s_1, s_2, \ldots, s_n$  given the observations  $O = o_1, o_2, \ldots, o_n$ 

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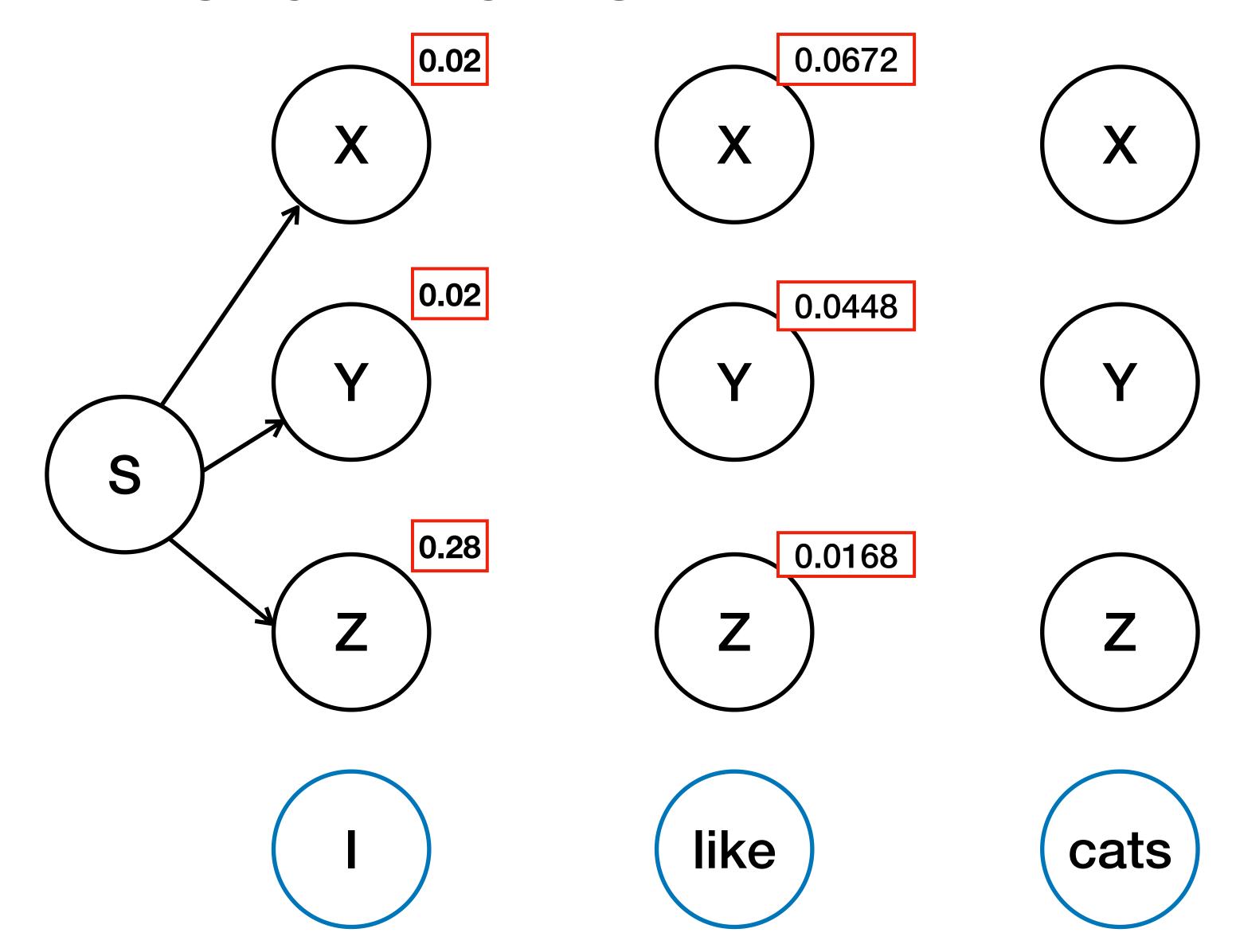
Thanks to Markov, we only need to "look-back" one time step to decode!

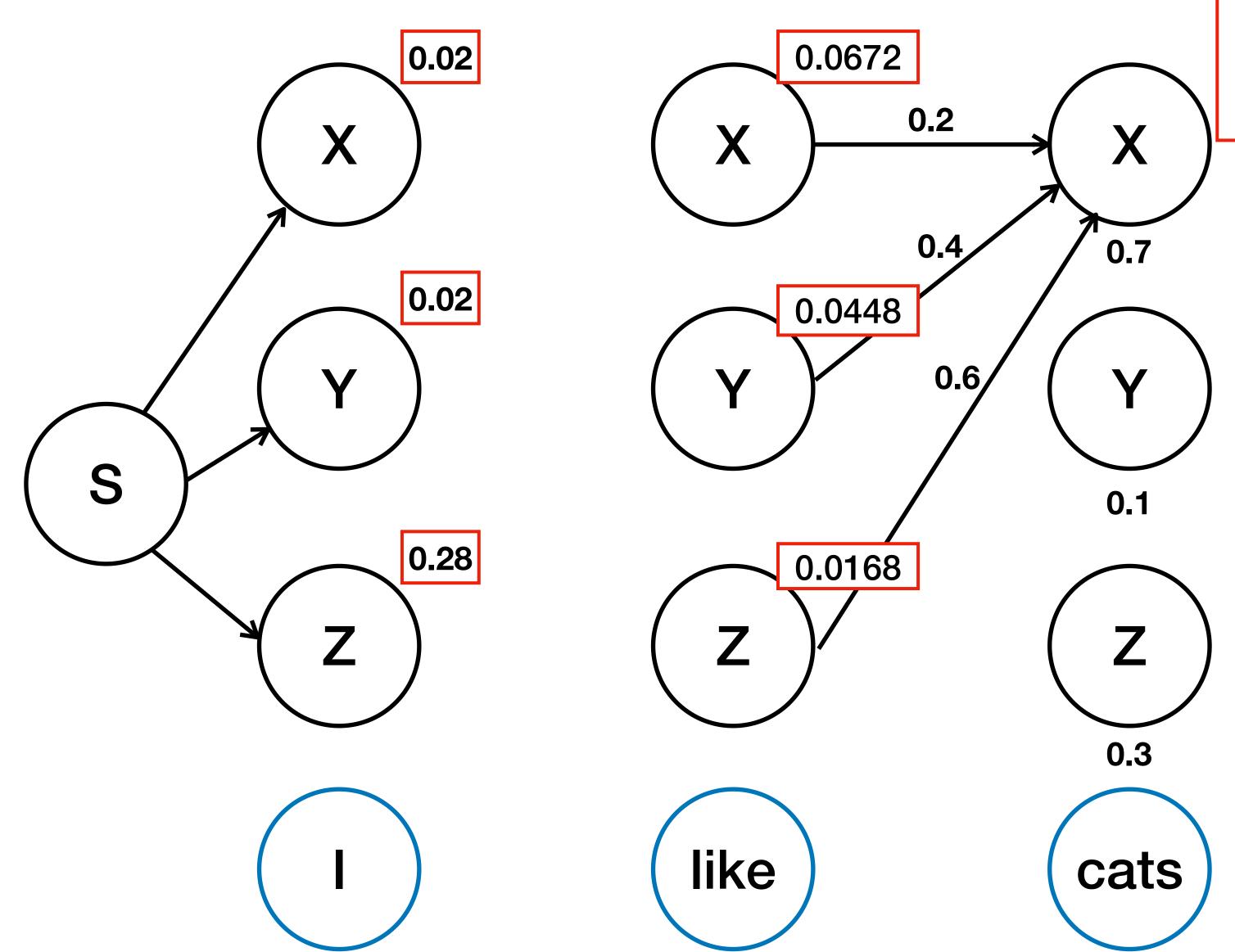
Makes dynamic programming possible, so we only need to keep the "best" for each previous time step to decode the next (instead of more history).



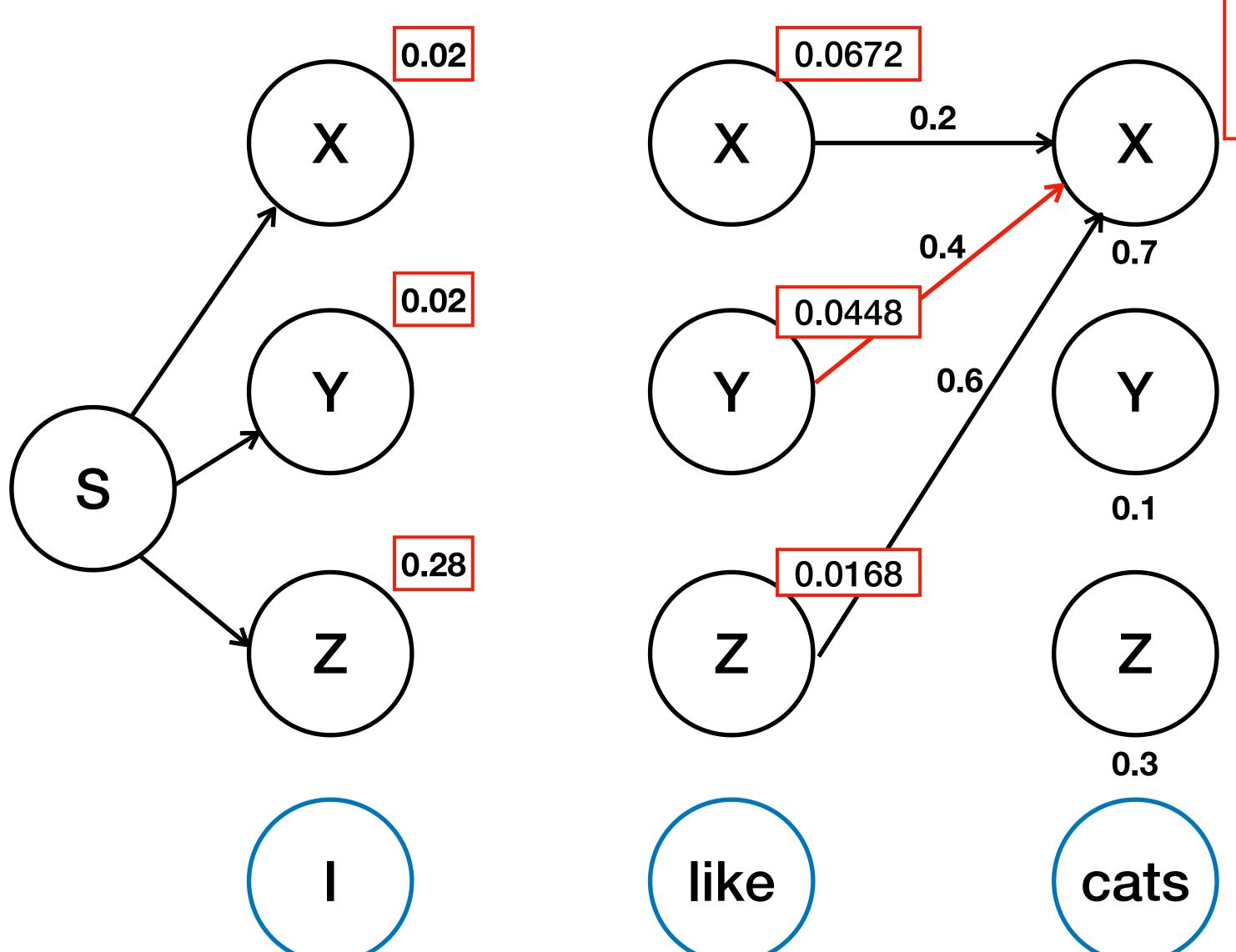
	X	Υ	Z
S	0.1	0.2	0.7
X	0.2	0.5	0.3
Υ	0.4	0.4	0.2
Z	0.6	0.2	0.2

		like	cats
X	0.2	0.1	0.7
Υ	0.1	0.8	0.1
Z	0.4	0.3	0.3

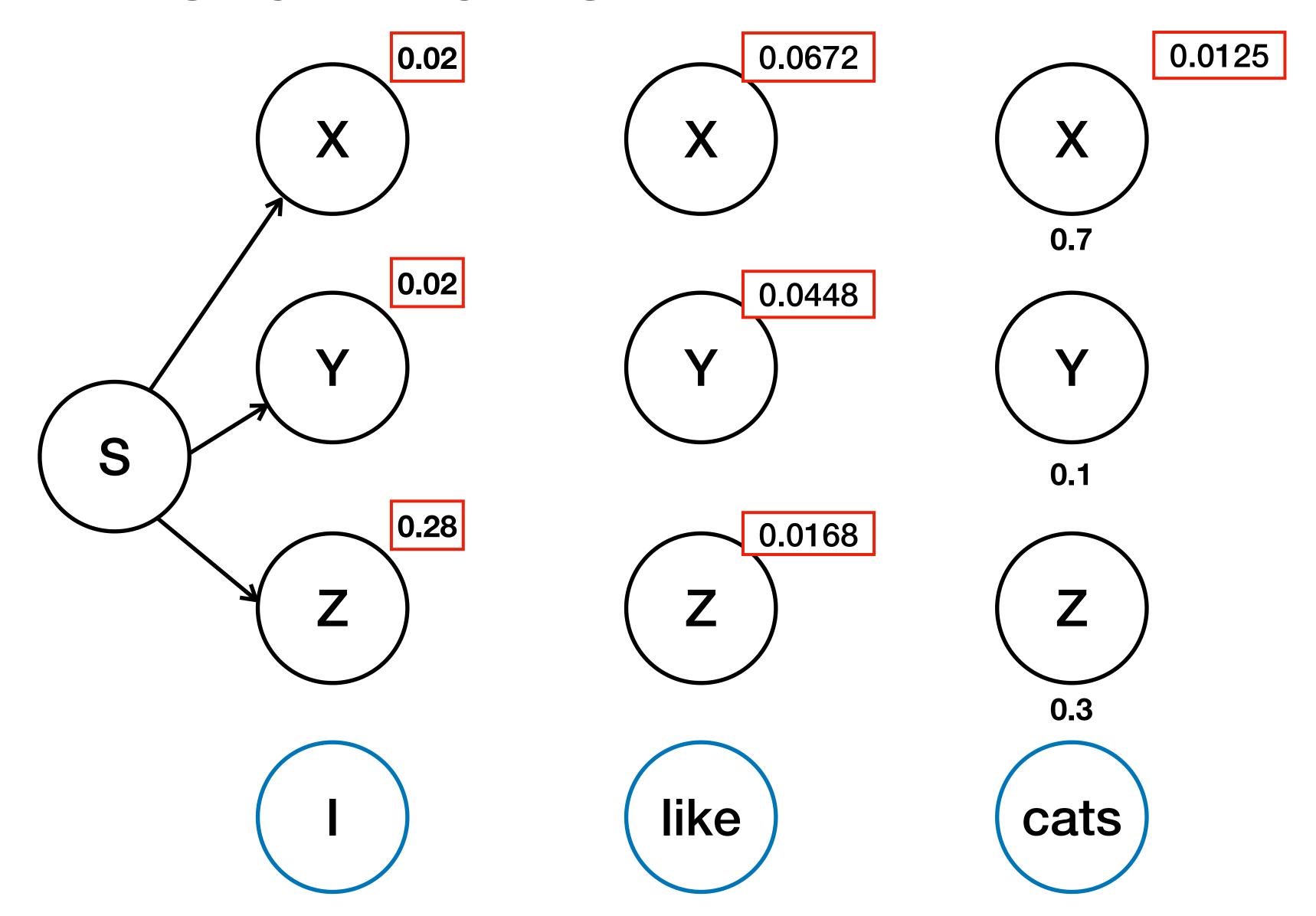


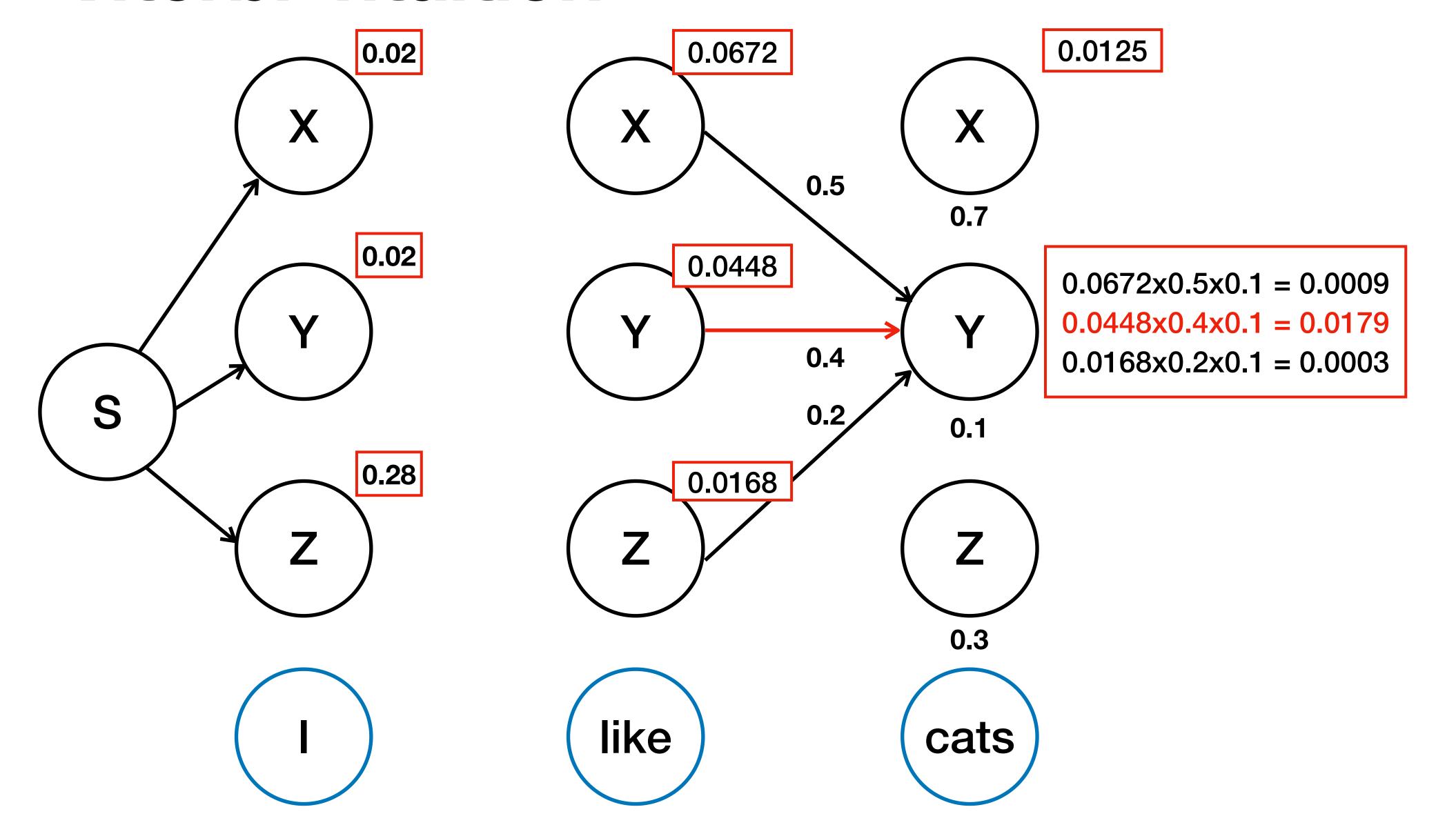


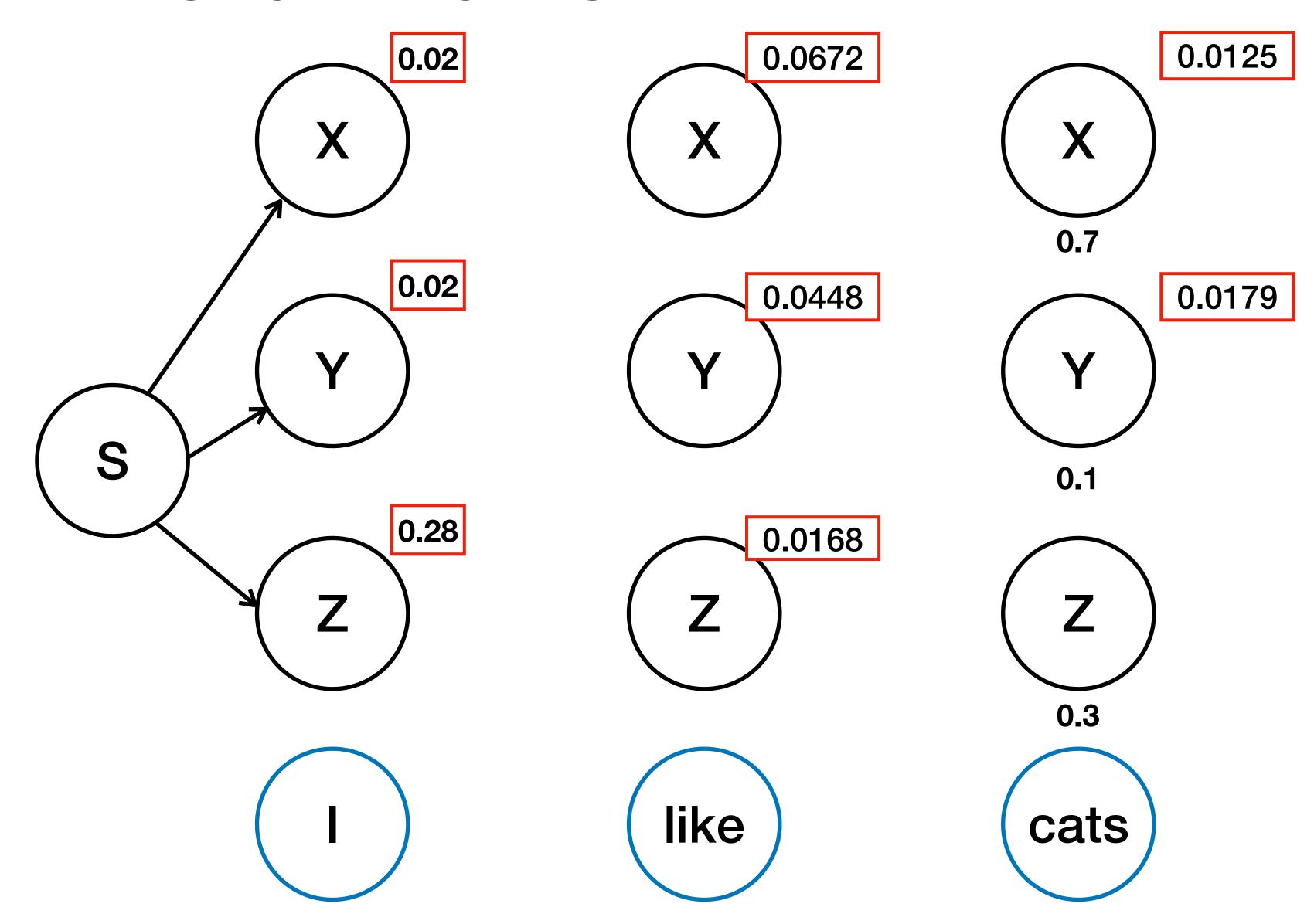
 $0.0672 \times 0.2 \times 0.7 = 0.0023$   $0.0448 \times 0.4 \times 0.7 = 0.0125$  $0.0168 \times 0.6 \times 0.7 = 0.0070$ 

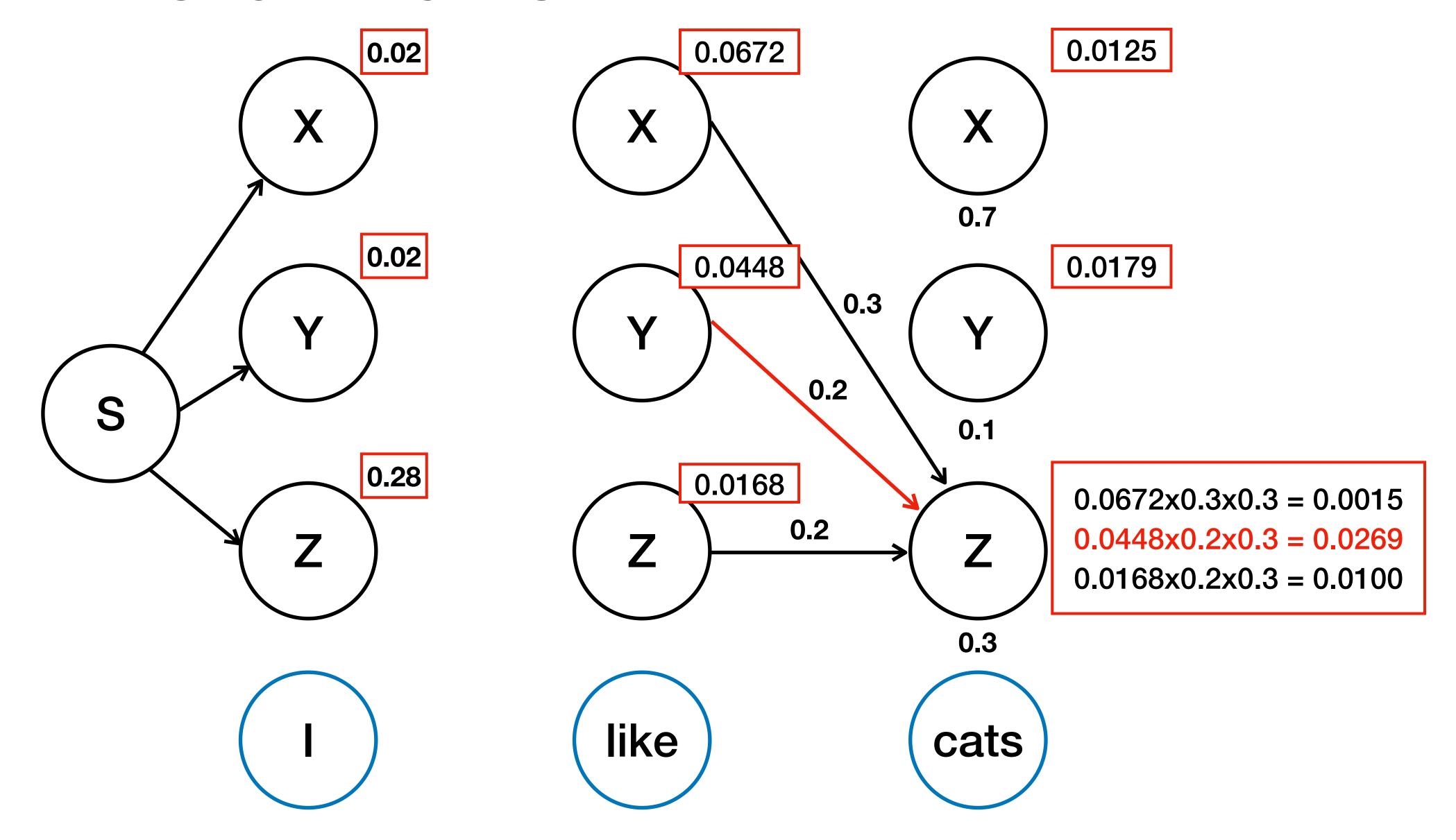


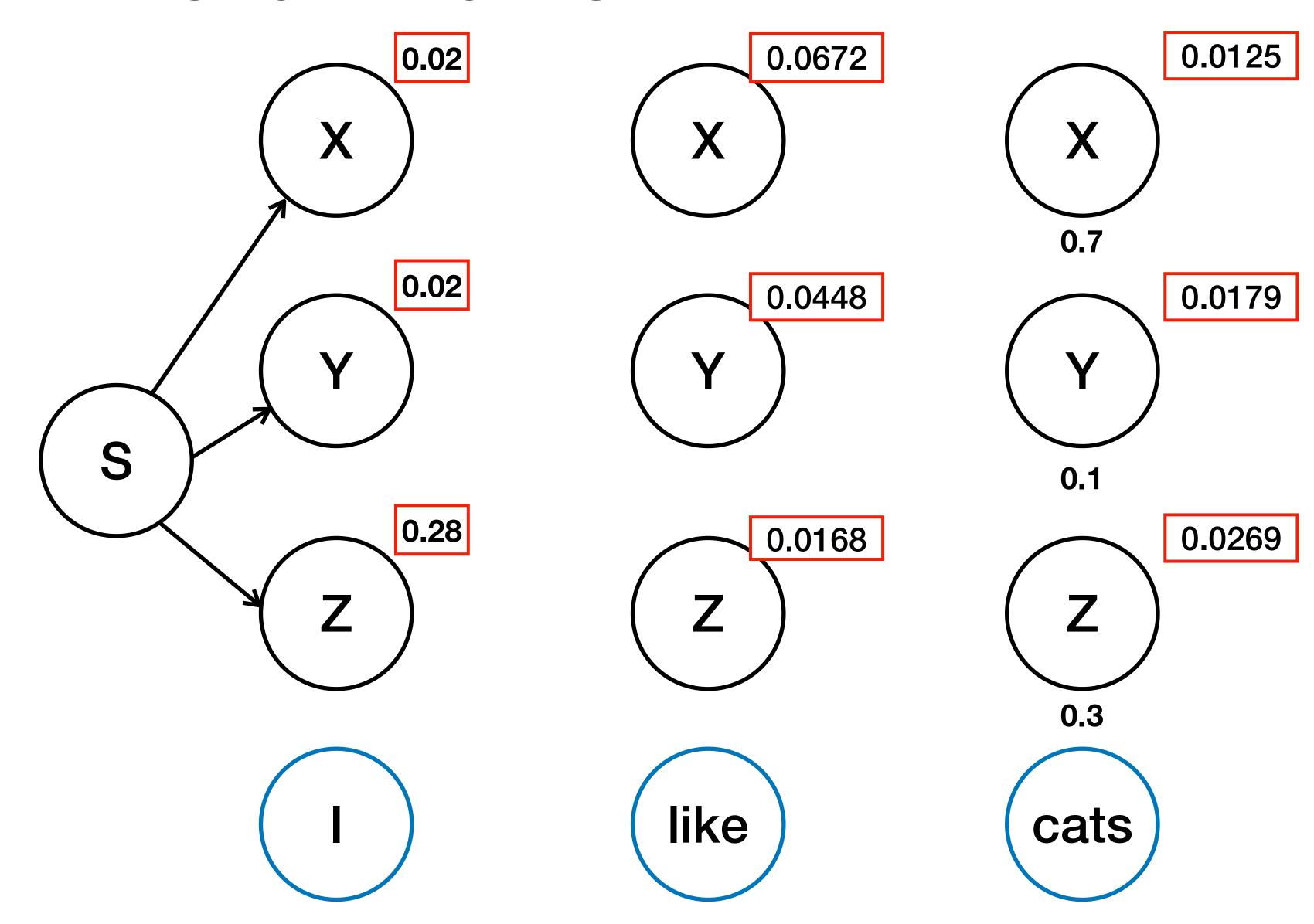
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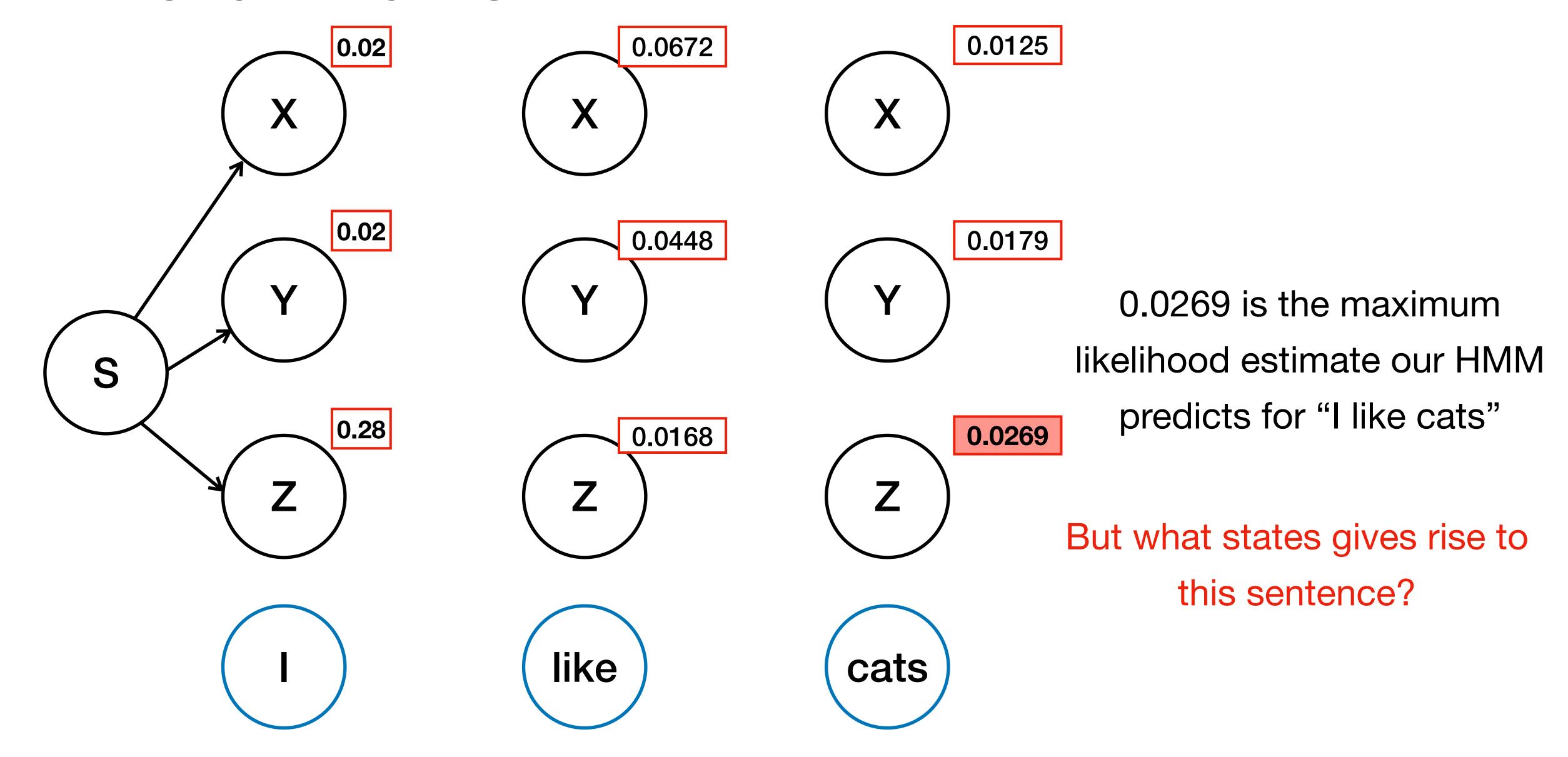




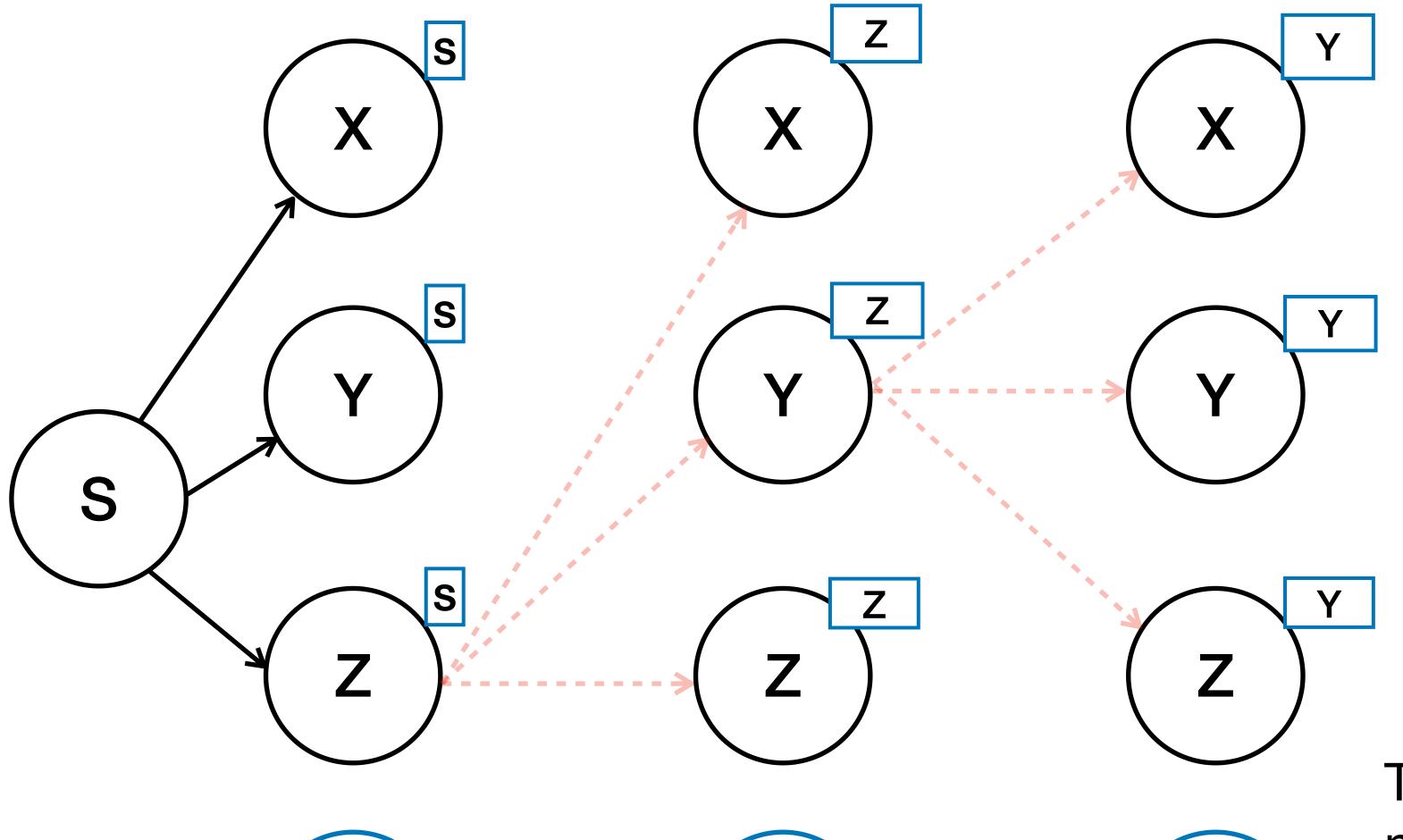








### Viterbi Intuition: Backtracking



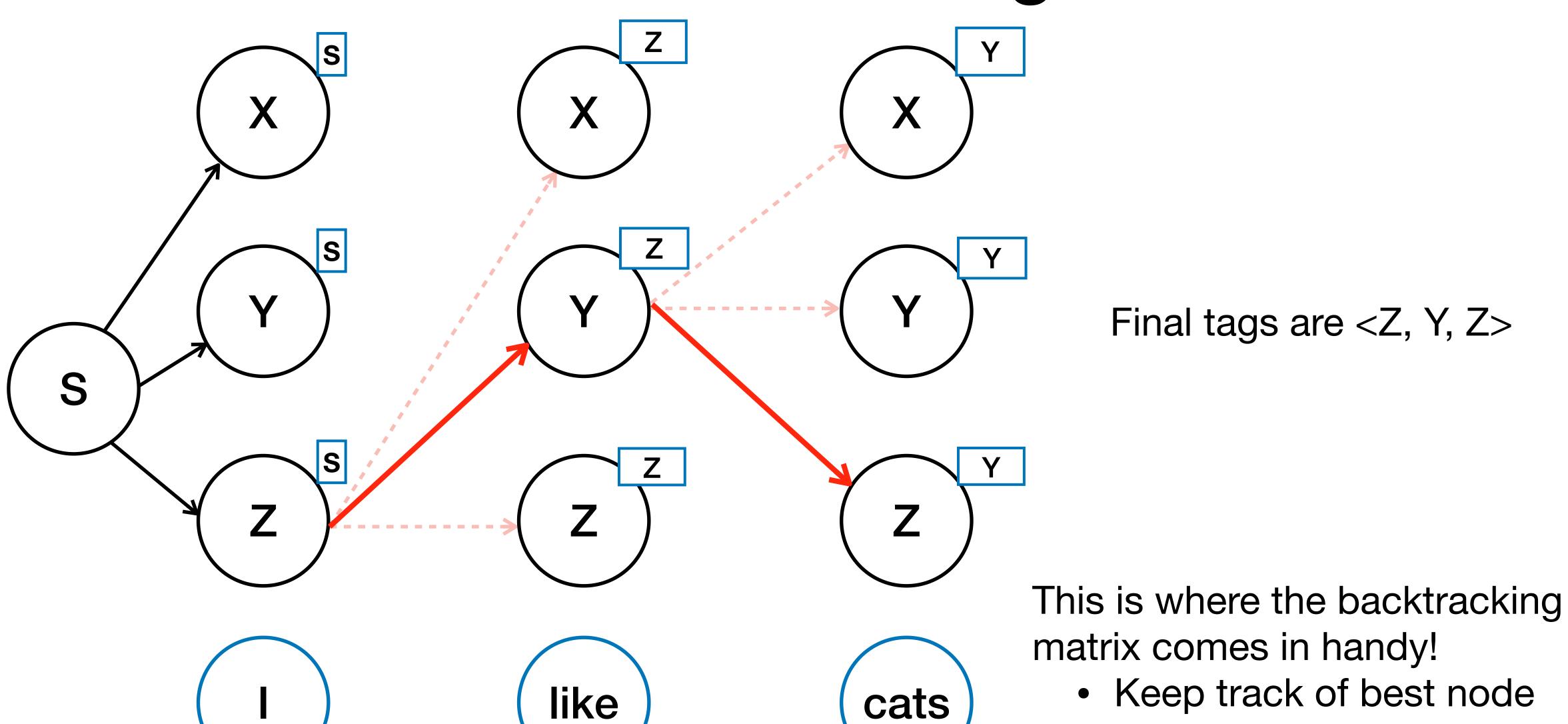
like

cats

This is where the backtracking matrix comes in handy!

 Keep track of best node from previous step

## Viterbi Intuition: Backtracking



from previous step

### Viterbi Understanding Check

How does Viterbi on a trigram HMM change? What about a 4-gram HMM?

### Viterbi Understanding Check

How does Viterbi on a trigram HMM change? What about a 4-gram HMM?

Key: Without Markov, we just need to look further back to calculate our likelihood!

. HMM extended to trigram, 4-gram etc: 
$$P(S,O) = \prod_{i=1}^n P(s_i \mid s_{i-1}, s_{i-2}) P(o_i \mid s_i)$$

. MLE estimate: 
$$P(s_i | s_{i-1}, s_{i-2}) = \frac{\text{Count}(s_i, s_{i-1}, s_{i-2})}{\text{Count}(s_{i-1}, s_{i-2})}$$

Viterbi:

$$M[i,j,k] = \max_{r} M[i-1,k,r] \ P(s_j | s_k, s_r) \ P(o_i | s_j) \quad 1 \le j, k, r \le K \quad 1 \le i \le n$$

- most probable sequence of states ending with state j at time i, and state k at i-1
- Time complexity =  $O(nK^3)$

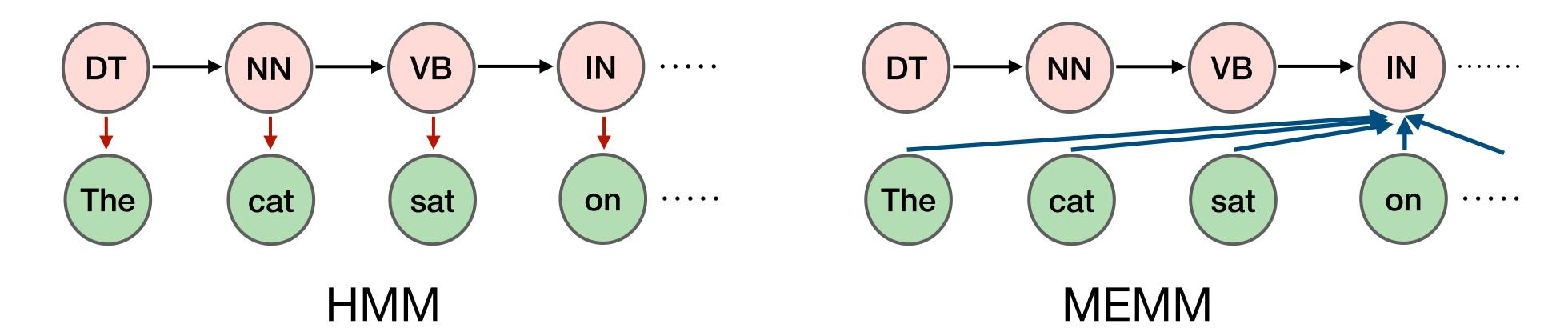
# Maximum Entropy Markov Models

#### Generative vs. Discriminative

- HMM is a generative model : we compute probability P(S, O)
- Can we model  $P(s_1, \ldots, s_n | o_1, \ldots, o_n)$  directly?

	Generative	Discriminative
Text classification	Naive Bayes: $P(c) P(d \mid c)$	Logistic Regression: $P(c \mid d)$
Sequence prediction	HMM: $P(s_1, \ldots s_n) P(o_1, \ldots o_n   s_1, \ldots s_n)$	MEMM: $P(s_1, \ldots s_n   o_1, \ldots o_n)$

## MEMM Basics

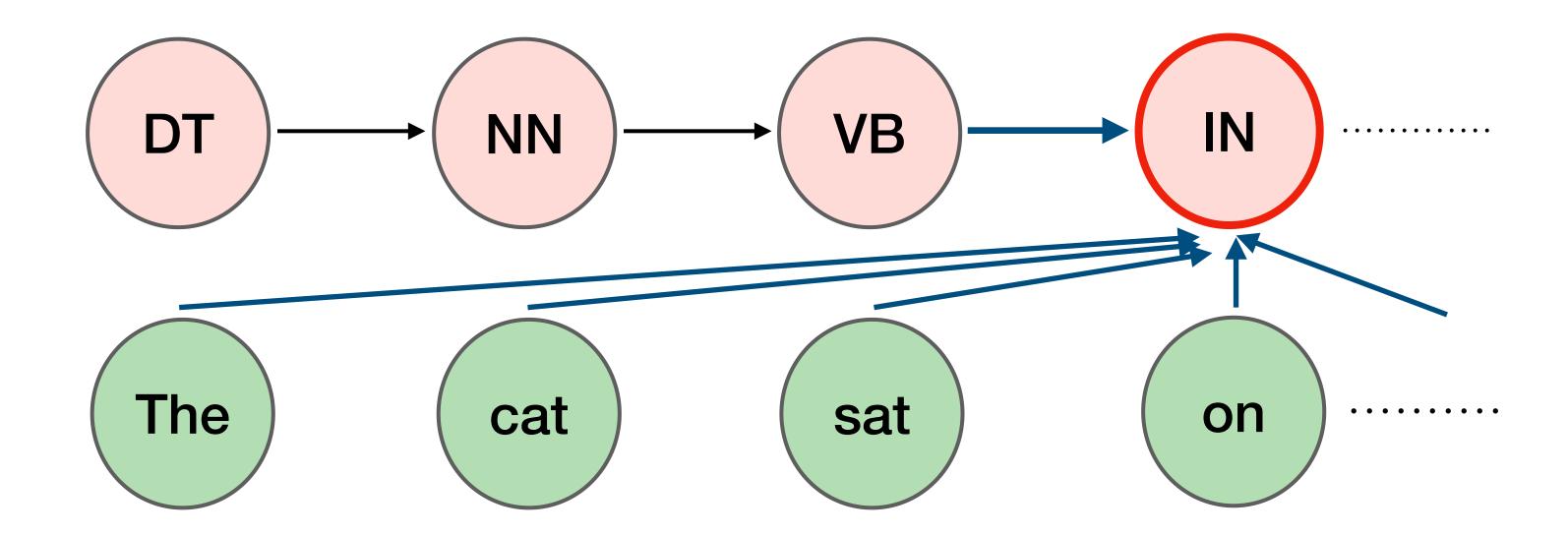


$$P(S \mid O) = \prod_{i=1}^{n} P(s_i \mid s_{i-1}, s_{i-2}, ..., s_1, O)$$

$$= \prod_{i=1}^{n} P(s_i \mid s_{i-1}, O)$$
Markov assumption:
Bigram MEMM

Instead of learning how to model observations given states, directly learn to predict states given observations.

## MEMM Basics



- To predict the red node, the bigram MEMM conditions on the "prior tag" (VB) and the observations in the window (The, cat, sat, on)
- Prior tags and observations will be transformed into features (some sort of vector representation) just like logistic regression

## (Bigram) MEMM Formulation

- To make the equivalence b/w MEMMs & logistic regression clearer, we will depart slightly from the lecture notation
- Our primary objective:  $P(S \mid O) = \prod_{i=1}^{n} P(s_i \mid s_{i-1}, s_{i-2}, ..., s_1, O) = \prod_{i=1}^{n} P(s_i \mid s_{i-1}, O)$

#### Logistic Regression

*Input*: documents *d* 

Features:  $\vec{\mathbf{x}} = \mathbf{f}(d)$  (#words, patterns, ...)

Output: 
$$z_i = \mathbf{w}^{(i)} \cdot \vec{\mathbf{x}} \quad (\mathbf{w}^{(i)} \in \mathbb{R}^d)$$

$$P(y = i \mid d) = \operatorname{softmax}(\mathbf{\vec{z}})_i$$

$$\propto \exp(\mathbf{w}^{(i)} \cdot \mathbf{f}(d))$$
weights features

#### **MEMM**

*Input*: state  $s_{i-1}$ , observations O, position i

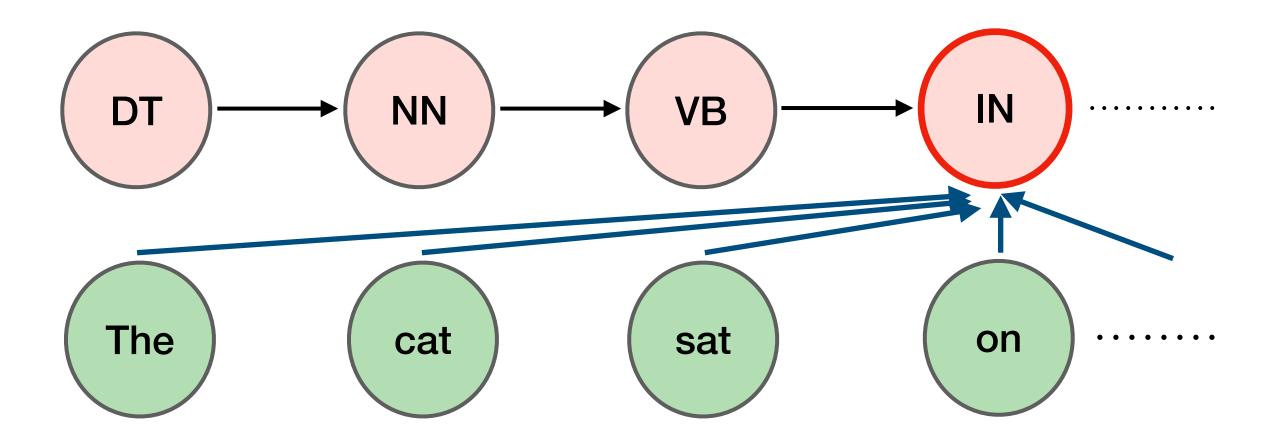
Features:  $\mathbf{f}(s_{i-1}, O, i)$  (1 if  $\langle man, the \rangle$  else 0)

Output: 
$$z_s = \mathbf{w}^{(s)} \cdot \mathbf{f}(s_{i-1}, O, i) \quad (\mathbf{w}^{(s)} \in \mathbb{R}^d)$$

$$P(s_i = s \mid s_{i-1}, O) = \operatorname{softmax}(\vec{\mathbf{z}})_s$$

$$\propto \exp(\mathbf{w}^{(s)} \cdot \mathbf{f}(s_{i-1}, O, i))$$
weights features

### MEMM Formulation

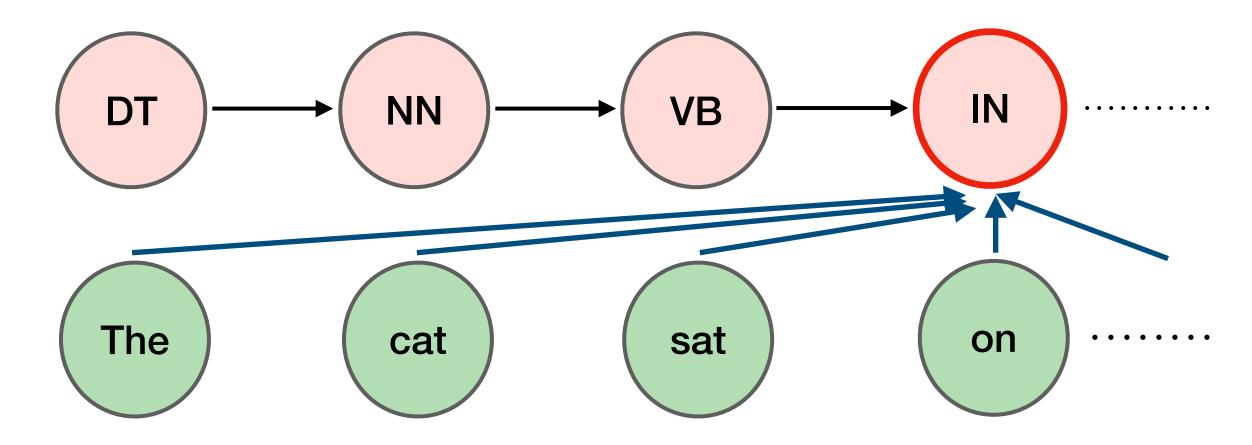


Examples of binary features and potential weights:

- $(o_{i-2} = \text{animal}, s_{i-1} = \text{VB})$ :  $w_{IN} = 3, w_{VB} = -1, w_{DT} = 0, w_{NN} = 1$
- (tri-gram)  $(s_{i-2} = NN, s_{i-1} = VB)$ :  $w_{IN} = 4,...$

**f** would look like [1,0,0,1,...], and  $\mathbf{w}_{\text{IN}} = [3,0,1,4,...]$ 

### MEMM Formulation

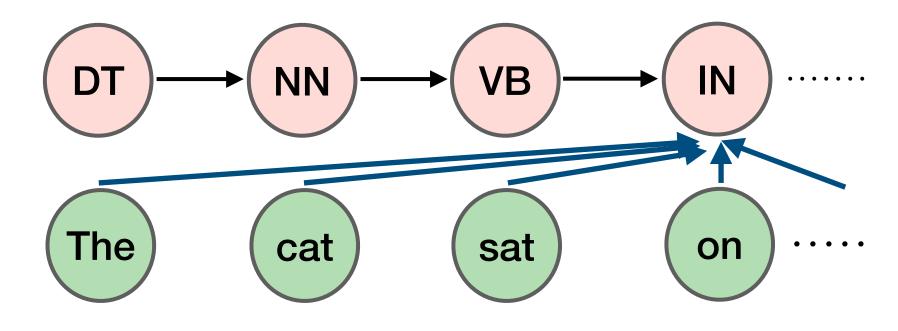


Can also define generic feature templates:

• 
$$\langle s_i, o_{i-2} \rangle$$
,  $\langle s_i, o_{i-1} \rangle$ ,  $\langle s_i, o_i \rangle$ ,  $\langle s_i, o_{i+1} \rangle$ ,  $\langle s_i, o_{i+2} \rangle$ 

• 
$$\langle s_i, s_{i-1} \rangle$$
,  $\langle s_i, s_{i-1}, s_{i-2} \rangle$ 

## MEMM Basics



MEMM

$$P(S \mid O) = \prod_{i=1}^{n} P(s_i \mid s_{i-1}, O)$$

$$P(s_i = s \mid s_{i-1}, O) \propto \exp(\mathbf{w}^{(s)} \cdot \mathbf{f}(s_{i-1}, O, i))$$
weights features

$$P(s_i = s \mid s_{i-1}, O) = \frac{\exp(\mathbf{w}^{(s)} \cdot \mathbf{f}(s_{i-1}, O, i))}{\sum_{s'=1}^{K} \exp(\mathbf{w}^{(s')} \cdot \mathbf{f}(s_{i-1}, O, i))}$$

#### Markov assumption: Bigram MEMM

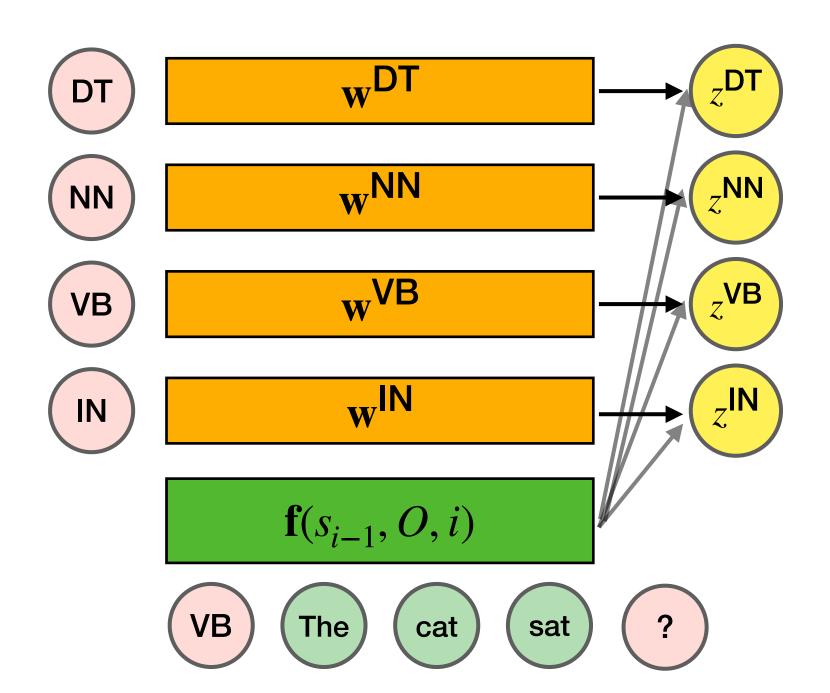
Important: you can define features over entire word sequence O!

## Quick Aside: Two Equivalent Formulations

This formulation is identical to one in lecture; it's just different in featurization.

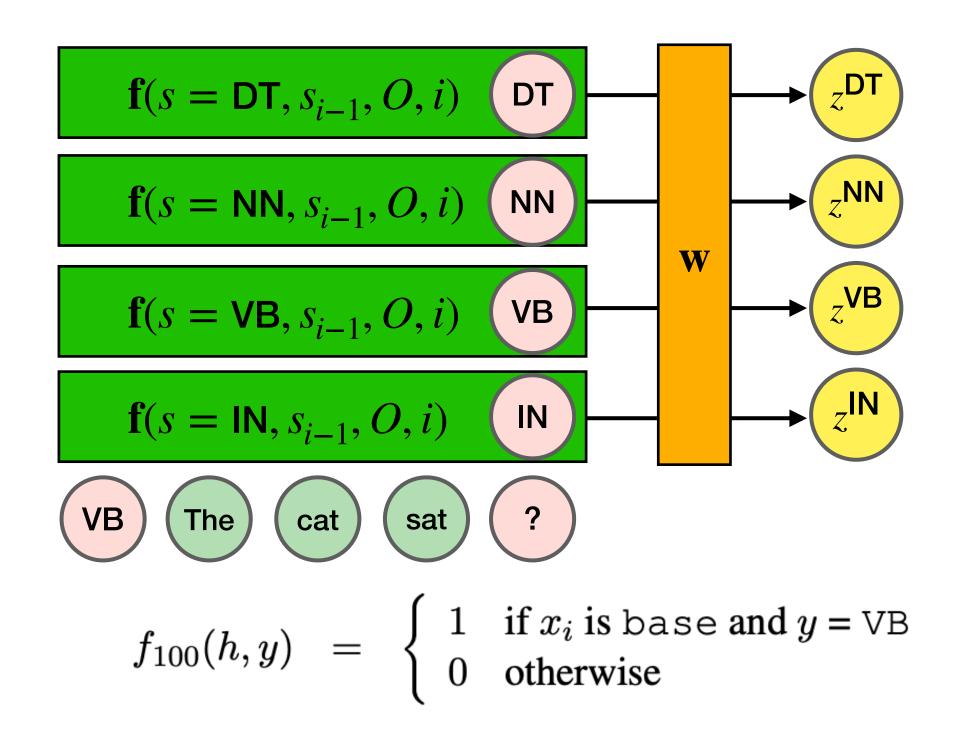
$$P(s_i = s \mid s_{i-1}, O) \propto \exp(\mathbf{w}^{(s)} \cdot \mathbf{f}(s_{i-1}, O, i))$$

Parameters:  $\mathbf{w}^{(s)} \in \mathbb{R}^d$  for s = 1, 2, ..., K

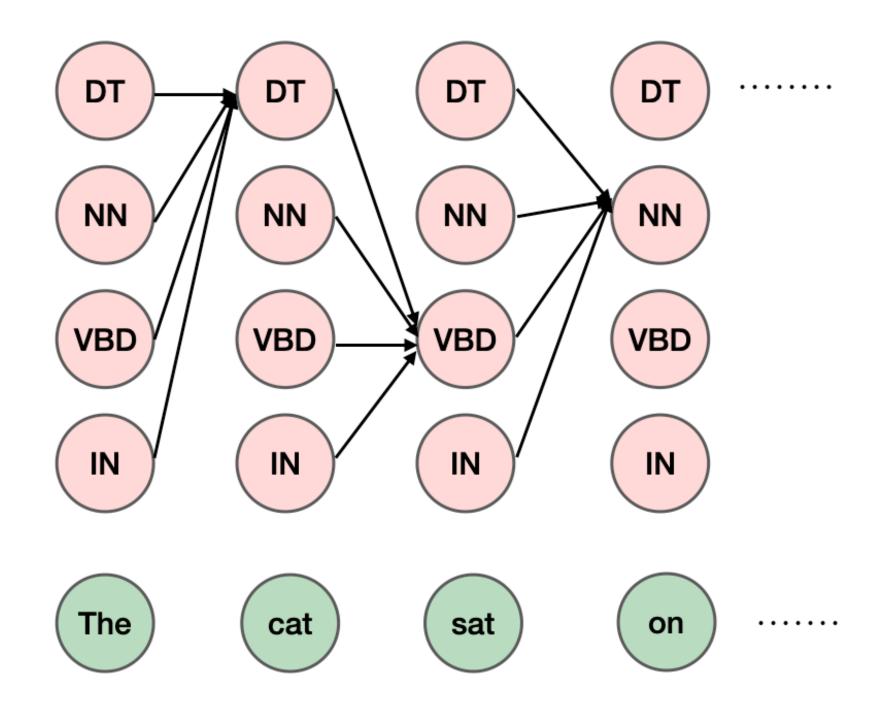


$$P(s_i = s \mid s_{i-1}, O) \propto \exp(\mathbf{w} \cdot \mathbf{f}(s = s_i, s_{i-1}, O, i))$$

Parameters:  $\mathbf{w} \in \mathbb{R}^d$ , same for all features



## Viterbi Decoding for MEMMs



M[i,j] stores joint probability of most probable sequence of states ending with state j at time i

$$M[i,j] = \max_{k} M[i-1,k] P(s_i = j | s_{i-1} = k, O)$$
  $1 \le k \le K$   $1 \le i \le n$ 

Backward: Pick  $\max_{k} M[n, k]$  and backtrack using B

### MEMMs vs. HMMs

- HMM models the joint P(S, O) while MEMM models the required prediction  $P(S \mid O)$
- MEMM has more expressivity
  - accounts for dependencies between neighboring states and entire observation sequence
  - allows for more flexible features
- HMM may hold an advantage if the dataset is small

# [SP23 Midterm] Problem 5: Lie Detection

You are building a lie detector taking as input a stream of recorded behaviors:

•  $x_t \in \{a, b\}$ , i.e., face touching (a) or blinking (b)

Detector at every moment then predicts one of four labels:

•  $y_t \in \{N, U, L, H\}$ , i.e., Neutral (N), Unclear (U), Lying (L), and Honest (H)

Dataset is triplets  $(y_{t-1}, y_t, x_t)$ : 9x of (N, L, a), 9x of (U, L, b), 1x of (N, H, b)

• Labels  $y_t$  come from body language experts' annotation

**Q1:** You use the above data to build an HMM. What is  $P(x_t = b \mid y_{t-1} = N)$ ?

## [SP23 Midterm] Problem 5: Lie Detection

(3) (4 points) Now assume you design the system with an MEMM model that predicts the label  $y_t$ :

$$P(y_{t} = \hat{y}_{t} \mid y_{t-1} = N, x_{t} = b) = \frac{\exp\left(\mathbf{w} \cdot \mathbf{f}(y_{t} = \hat{y}_{t}, y_{t-1} = N, x_{t} = b)\right)}{\sum_{y'_{t}} \exp\left(\mathbf{w} \cdot \mathbf{f}(y_{t} = y'_{t}, y_{t-1} = N, x_{t} = b)\right)},$$

where  $\mathbf{f}(\cdot)$  is defined by the feature templates:  $\langle y_{t-1}, y_t \rangle$ ,  $\langle y_t, x_t \rangle$ , and  $\langle y_{t-1}, y_t, x_t \rangle$ . Assume that, after the model is trained, the learned weight vector  $\mathbf{w}$  is:

$$w_{i} = \begin{cases} 0.2, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = L] \\ 0.1, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = H] \\ 0.6, & \text{for } \mathbb{1}[y_{t} = L, x_{t} = b] \\ 0.5, & \text{for } \mathbb{1}[y_{t} = H, x_{t} = b] \\ 0.7, & \text{for } \mathbb{1}[y_{t-1} = U, y_{t} = L, x_{t} = b] \\ 0.3, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = H, x_{t} = b] \\ 0, & \text{otherwise} \end{cases}$$

where  $w_i$  is the element in **w**. Which tag will this model predict at time step t if we know  $y_{t-1} = N$  and  $x_t = b$ ? With the model predicting  $\hat{y}_t$ , does this triplet  $(y_{t-1} = N, y_t = \hat{y}_t, x_t = b)$  have a higher count than all other triplets  $(y_{t-1} = N, y_t = y'_t, x_t = b)$  in the past annotated recordings?

#### Labels are (N, H, L, U)

## [SP23 Midterm] Problem 5: Lie Detection

(a) Just add up each of the  $z_i$  for each label; unnormalized is good enough

• 
$$\tilde{P}(y_t = H \mid y_{t-1} = N, x_t = b) = 0.1 + 0.5 + 0.3 = 0.9$$

• 
$$\tilde{P}(y_t = L \mid y_{t-1} = N, x_t = b) = 0.2 + 0.6 = 0.8$$

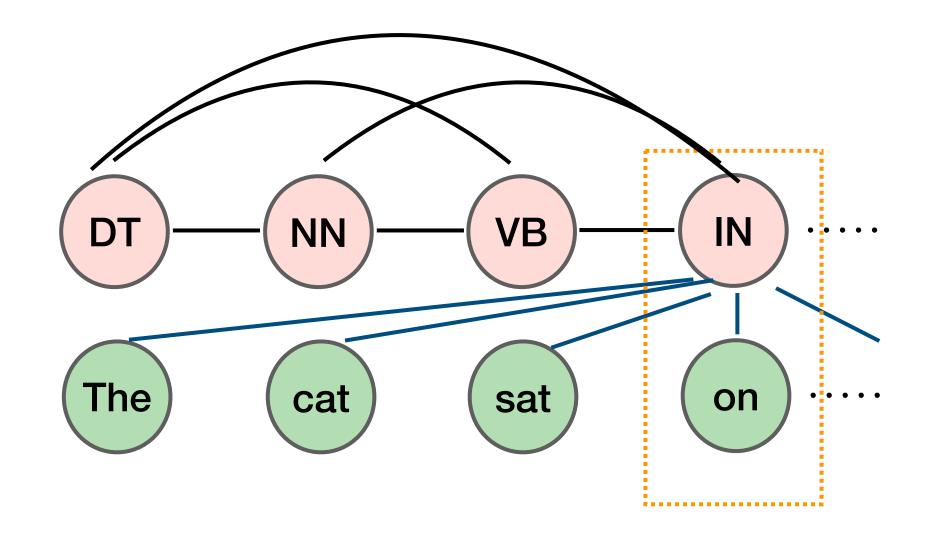
Therefore  $\hat{y}_t = H$  is most likely.

(b) Yes. The 0.3 on  $(y_{t-1}=N,y_t=H,x_t=b)$  shows that.

$$w_{i} = \begin{cases} 0.2, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = L] \\ 0.1, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = H] \\ 0.6, & \text{for } \mathbb{1}[y_{t} = L, x_{t} = b] \\ 0.5, & \text{for } \mathbb{1}[y_{t} = H, x_{t} = b] \\ 0.7, & \text{for } \mathbb{1}[y_{t-1} = U, y_{t} = L, x_{t} = b] \\ 0.3, & \text{for } \mathbb{1}[y_{t-1} = N, y_{t} = H, x_{t} = b] \\ 0, & \text{otherwise} \end{cases}$$

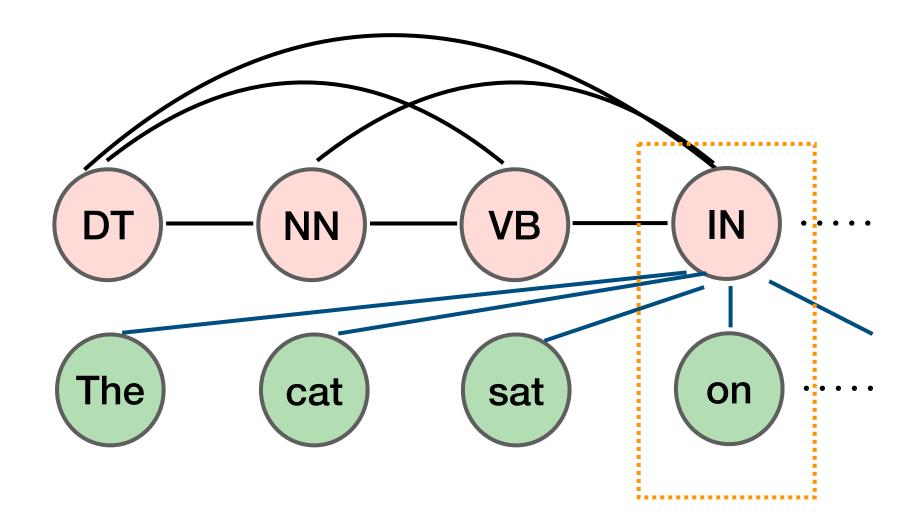
## Conditional Random Field

- Model  $P(s_1, \ldots, s_n | o_1, \ldots, o_n)$  directly
- No Markov assumption
  - Map entire sequence of states S and observations O to a global feature vector
    - Normalize over entire sequences
- Generalization of MEMMs



$$P(S \mid O) = \frac{\exp(\mathbf{w} \cdot \mathbf{f}(S, O))}{\sum_{S'} \exp(\mathbf{w} \cdot \mathbf{f}(S', O))} = \frac{\exp(\mathbf{w} \cdot \mathbf{f}(S, O))}{Z(O)}$$

#### Features



$$1\{x_i = the, y_i = DET\}$$

$$1\{y_i = PROPN, x_{i+1} = Street, y_{i-1} = NUM\}$$

$$1\{y_i = VERB, y_{i-1} = AUX\}$$

$$P(S \mid O) = \frac{\exp(\sum_{k=1}^{m} w_k \cdot F_k(S, O))}{\sum_{S'} \exp(\sum_{k=1}^{m} w_k \cdot F_k(S', O))}$$

- Each  $F_k$  in f is a global feature function
- · Can be computed as a combination of local

features: 
$$F_k = \sum_{i=1}^{n} f_k(s_{i-1}, s_i, O, i)$$

Each local feature only depends on previous and current states

#### CRFs vs. MEMMs

- MEMM models the required prediction  $P(S \mid O)$  using the Markov assumption, while the CRF does not
- CRF uses global features while MEMM features are localized
- Feature design is flexible in both models
- CRF is computationally more complex