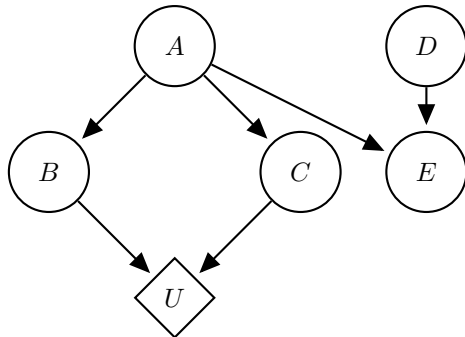


# CS188: Exam Practice Session 9

## Q1. VPI

(a) Consider a decision network with the following structure, where node  $U$  is the utility:



(i) For each of the following, choose the most specific option that is guaranteed to be true:

- |   |  |   |
|---|--|---|
| <input type="radio"/> $VPI(B) = 0$      | <input type="radio"/> $VPI(B) \geq 0$      | <input type="radio"/> $VPI(B) > 0$      |
| <input type="radio"/> $VPI(D) = 0$      | <input type="radio"/> $VPI(D) \geq 0$      | <input type="radio"/> $VPI(D) > 0$      |
| <input type="radio"/> $VPI(E) = 0$      | <input type="radio"/> $VPI(E) \geq 0$      | <input type="radio"/> $VPI(E) > 0$      |
| <input type="radio"/> $VPI(A E) = 0$    | <input type="radio"/> $VPI(A E) \geq 0$    | <input type="radio"/> $VPI(A E) > 0$    |
| <input type="radio"/> $VPI(E A) = 0$    | <input type="radio"/> $VPI(E A) \geq 0$    | <input type="radio"/> $VPI(E A) > 0$    |
| <input type="radio"/> $VPI(A B, C) = 0$ | <input type="radio"/> $VPI(A B, C) \geq 0$ | <input type="radio"/> $VPI(A B, C) > 0$ |

(ii) For each of the following, fill in the blank with the most specific of  $>$ ,  $\geq$ ,  $<$ ,  $\leq$ ,  $=$  to guarantee that the comparison is true, or write ? if there is no possible guarantee.

$VPI(B)$  \_\_\_\_\_  $VPI(A)$

$VPI(B, C)$  \_\_\_\_\_  $VPI(A)$

$VPI(B, C)$  \_\_\_\_\_  $VPI(B) + VPI(C)$

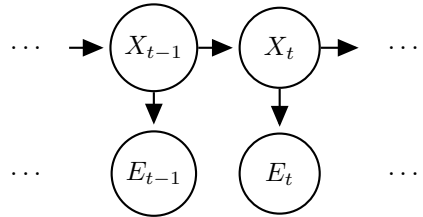
$VPI(B|C)$  \_\_\_\_\_  $VPI(A|C)$

## Q2. HMMs - Forward Algorithm

Below is the forward algorithm update equation for Hidden Markov Models. As seen in lecture, we used  $e_{1:t}$  to denote all the evidence variables  $e_1, e_2, \dots, e_t$ . Similarly,  $e_{1:t-1}$  denotes  $e_1, e_2, \dots, e_{t-1}$ . For reference, the Bayes net corresponding to the usual Hidden Markov Model is shown on the right side of the equation below.

$$P(x_t | e_{1:t}) \propto P(x_t, e_{1:t}) \quad (1)$$

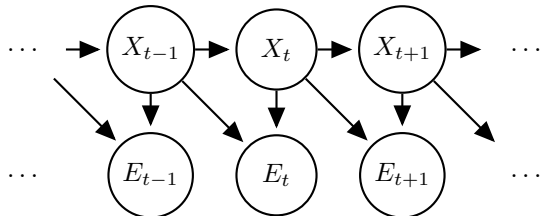
$$= P(e_t | x_t) \sum_{x_{t-1}} P(x_t | x_{t-1}) P(x_{t-1}, e_{1:t-1}) \quad (2)$$



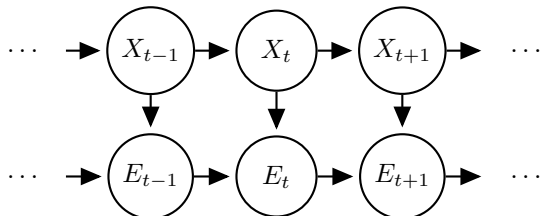
Hidden Markov Models can be extended in a number of ways to incorporate additional relations. Since the independence assumptions are different in these extended Hidden Markov Models, the forward algorithm updates will also be different.

Complete the forward algorithm updates for the extended HMMs specified by the following Bayes nets:

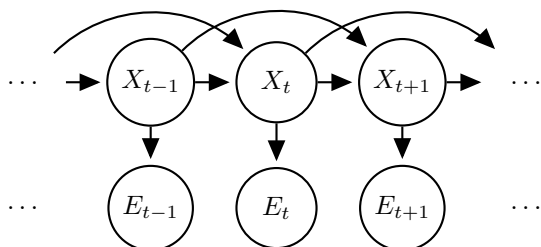
(a)  $P(x_t | e_{1:t}) \propto \sum_{x_{t-1}} P(x_{t-1}, e_{1:t-1}) \cdot \underline{\hspace{10em}}$



(b)  $P(x_t | e_{1:t}) \propto \sum_{x_{t-1}} P(x_{t-1}, e_{1:t-1}) \cdot \underline{\hspace{10em}}$



(c)  $P(x_t, x_{t+1} | e_{1:t}) \propto \sum_{x_{t-1}} P(x_{t-1}, x_t, e_{1:t-1}) \cdot \underline{\hspace{10em}}$

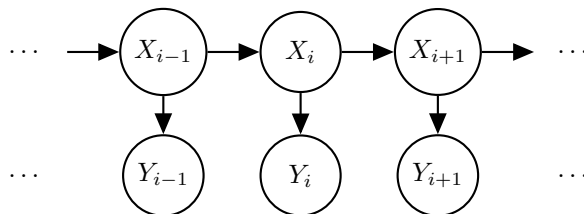


### Q3. DNA Sequencing

Suppose you want to model the problem of DNA sequencing using the following set-up:

- $X_i, Y_i \in \{A, T, C, G\}$
- $X_i$  :  $i$ th base of an individual
- $Y_i$  :  $i$ th base output by DNA sequencer

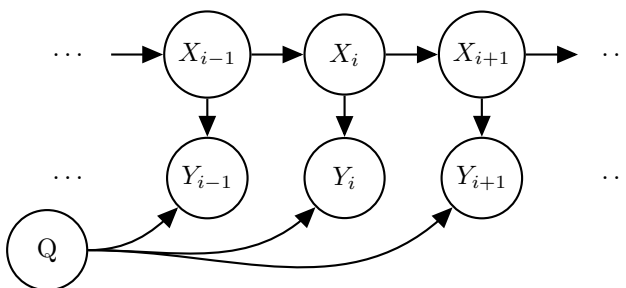
(a) First, you start by using a standard HMM model, shown below.



(i) Which of the following assumptions are made by the above HMM model

- |  |  |
|--|--|
| <input type="checkbox"/> $X_i \perp\!\!\!\perp X_j \quad \forall i \neq j$           | <input type="checkbox"/> $X_i \perp\!\!\!\perp Y_{i+1} \mid X_{i+1} \quad \forall i$ |
| <input type="checkbox"/> $Y_i \perp\!\!\!\perp Y_j \quad \forall i \neq j$           | <input type="checkbox"/> None of the provided options.                               |
| <input type="checkbox"/> $X_i \perp\!\!\!\perp Y_j \quad \forall i \neq j$           |  |
| <input type="checkbox"/> $X_{i-1} \perp\!\!\!\perp X_{i+1} \mid X_i \quad \forall i$ |  |

(b) Now you want to model the quality of your sequencer with a random variable  $Q$ , and decide to use the following modified HMM:



(i) Which of the following assumptions are made by the above modified HMM model?

- |  |  |
|--|--|
| <input type="checkbox"/> $X_i \perp\!\!\!\perp X_j \quad \forall i \neq j$           | <input type="checkbox"/> $Q \perp\!\!\!\perp X_i \quad \forall i$                      |
| <input type="checkbox"/> $Y_i \perp\!\!\!\perp Y_j \quad \forall i \neq j$           | <input type="checkbox"/> $Q \perp\!\!\!\perp X_i \mid Y_i \quad \forall i$             |
| <input type="checkbox"/> $X_i \perp\!\!\!\perp Y_j \quad \forall i \neq j$           | <input type="checkbox"/> $Q \perp\!\!\!\perp X_i \mid Y_1, \dots, Y_N \quad \forall i$ |
| <input type="checkbox"/> $X_{i-1} \perp\!\!\!\perp X_{i+1} \mid X_i \quad \forall i$ | <input type="checkbox"/> None of the provided options.                                 |
| <input type="checkbox"/> $X_i \perp\!\!\!\perp Y_{i+1} \mid X_{i+1} \quad \forall i$ |  |

(ii) You observe the sequencer output  $y_1, \dots, y_N$  and want to estimate probability distribution of the particular sequence of length  $c$  starting at base  $k$ :  $P(X_k \dots X_{k+c-1} \mid y_1, \dots, y_N)$ .

Select all elimination orderings which are maximally efficient with respect to the sum of the generated factors' sizes.

- |  |  |
|--|--|
| <input type="checkbox"/> $X_1, \dots, X_{k-1}, X_{k+c}, \dots, X_N, Q$ | <input type="checkbox"/> $X_1, \dots, X_{k-1}, Q, X_{k+c}, \dots, X_N$ |
| <input type="checkbox"/> $X_1, \dots, X_{k-1}, X_N, \dots, X_{k+c}, Q$ | <input type="checkbox"/> $X_1, \dots, X_{k-1}, Q, X_N, \dots, X_{k+c}$ |
| <input type="checkbox"/> $Q, X_1, \dots, X_{k-1}, X_{k+c}, \dots, X_N$ | <input type="checkbox"/> None of the provided options: _____           |
| <input type="checkbox"/> $Q, X_1, \dots, X_{k-1}, X_N, \dots, X_{k+c}$ |  |

(iii) How many entries are in the final conditional probability table  $P(X_k, \dots, X_{k+c-1} \mid y_1, \dots, y_N)$ ? The answer takes the form  $a^b$  – what are  $a$  and  $b$ ?

$a =$  \_\_\_\_\_

$b =$  \_\_\_\_\_