

# Lecture 6:

## Exact Inference via Clique-Tree Algorithms

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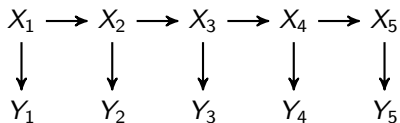
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Graphical Models PhD Course  
WASP

Graph structure gives us complexity bounds on VE.

Graph structure (specifically of chordal graphs) can also give us exact inference algorithms with nice computational benefits.

**Example (Hidden Markov Model).** Suppose we have a distribution Markov to the following graph:

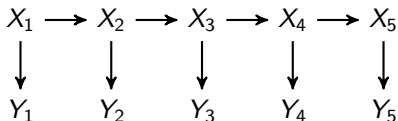


$$X_i | X_{i-1} = x_{i-1} \sim U(\{x_{i-1}/m, \dots, (m-1)x_{i-1}/m\}) \quad m > 1 \text{ fixed and } i = 1, \dots, n$$

$$Y_i | X_i = x_i \sim \text{Bin}(n, x_i) \quad n > 0 \text{ fixed and } i = 1, \dots, n$$

$[X_1, \dots, X_5]^T$  is a joint prior for  $\mathbf{Y} = [Y_1, \dots, Y_5]^T$ . model for sequences

We observe data  $\mathbf{y} = [y_1, \dots, y_5]^T$ . Can we estimate the (marginal) posterior distributions  $f_{X_i|\mathbf{Y}}(x_i|\mathbf{y})$ ?



**One answer:** Run VE.

Need many runs to compute all of  $f_{X_i|\mathbf{Y}}(x_i|\mathbf{y})$

Alternatively, we reuse  $\tau_i$ 's that we compute in one run in another run. This is the basic idea of the **Clique-Tree Algorithm**.

Start without evidence and assume we have estimated the conditional distributions:  $X_i|X_{i-1} = x_{i-1}$  and  $Y_i|X_i = x_i$ .

Elimination order:

$\prec = (Y_5, X_5, Y_4, X_4, Y_3, X_3, Y_2, X_2, Y_1)$

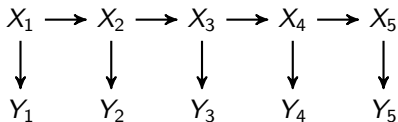
get  $f_{X_1}(x_1)$  with VE

$i$ -th step creates factors:  $\psi_i$  and  $\tau_i$

$$\psi_1(x_5, y_5) = f_{Y_5|X_5}(y_5|x_5) \quad \Rightarrow \quad \tau_1(x_5)$$

$$\psi_2(x_4, x_5) = \tau_1(x_5)f_{X_5|X_4}(x_5|x_4) \quad \Rightarrow \quad \tau_2(x_4)$$

$\vdots$

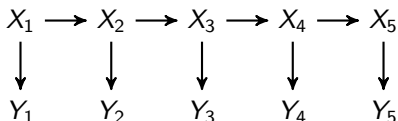


$\tau_i$  is a **message** being passed from node  $i$  to its parent  $j$ .

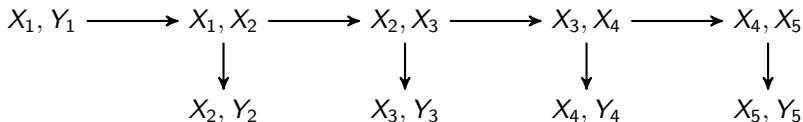
The parent node then uses  $\tau_i$  in the computation of its own factor  $\psi_{i-1}$  called its **potential**.

$$\begin{array}{llll}
 \text{potential for } Y_5: & \psi_1(x_5, y_5) & \Rightarrow & \tau_1(x_5) \quad \text{message from } Y_5 \text{ to } X_5 \\
 \text{potential for } X_5: & \psi_2(x_4, x_5) & \Rightarrow & \tau_2(x_4) \quad \text{message from } X_5 \text{ to } X_4
 \end{array}$$

Since the  $\tau_i$ 's are messages getting passed from one node  $i$  to another node  $j$  we denote them  $m_{i \rightarrow j}$ .



Messages are not really getting passed between **nodes**, but really **cliques** of nodes used in the potential that receives the message:



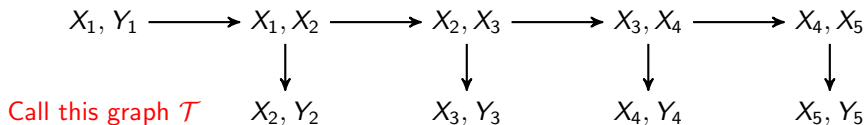
**Cliques:**

$$\begin{aligned}
 \mathcal{C} = \{ & C_1 = \{X_1, Y_1\}, C_2 = \{X_1, X_2\}, C_3 = \{X_2, X_3\}, C_4 = \{X_3, X_4\}, \\
 & C_5 = \{X_4, X_5\}, C_6 = \{X_2, Y_2\}, C_7 = \{X_3, Y_3\}, C_8 = \{X_4, Y_4\}, C_9 = \{X_5, Y_5\} \}.
 \end{aligned}$$

**Initial Factors:**

$$\begin{aligned}
 \Phi = \{ & f(x_1), f(x_2|x_1), f(x_3|x_2), f(x_4|x_3), f(x_5|x_4), \\
 & f(y_1|x_1), f(y_2|x_2), f(y_3|x_3), f(y_4|x_4), f(y_5|x_5) \}.
 \end{aligned}$$

$$\begin{aligned}
 & \alpha : \Phi \rightarrow \mathcal{C} \text{ such that} \\
 & \text{Scope}[\phi] \subseteq \alpha(C)
 \end{aligned}$$



$\alpha : \Phi \longrightarrow \mathcal{C}$  defines the **initial potentials**:  $\psi_i = \prod_{\phi: \alpha(\phi)=C_i} \phi$ .

$$\psi_1(x_1, y_1) = f(x_1)f(y_1|x_1)$$

initial potential for  $C_1 = \{X_1, Y_1\}$

$$\psi_2(x_1, x_2) = f(x_2|x_1)$$

initial potential for  $C_2 = \{X_1, X_2\}$

$$\psi_3(x_2, x_3) = f(x_3|x_2)$$

initial potential for  $C_3 = \{X_2, X_3\}$

$$\psi_4(x_3, x_4) = f(x_4|x_3)$$

initial potential for  $C_4 = \{X_3, X_4\}$

$$\psi_5(x_5, x_4) = f(x_5|x_4)$$

initial potential for  $C_5 = \{X_4, X_5\}$

$$\psi_6(x_2, y_2) = f(y_2|x_2)$$

initial potential for  $C_6 = \{X_2, Y_2\}$

$$\psi_7(x_2, y_2) = f(y_3|x_3)$$

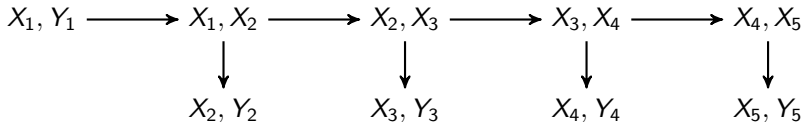
initial potential for  $C_7 = \{X_3, Y_3\}$

$$\psi_8(x_2, y_2) = f(y_4|x_4)$$

initial potential for  $C_8 = \{X_4, Y_4\}$

$$\psi_9(x_2, y_2) = f(y_5|x_5)$$

initial potential for  $C_9 = \{X_5, Y_5\}$



$\mathcal{T}$  is a rooted tree with source node  $C_1 = \{X_1, Y_1\}$ :

Every node  $C_i$  in  $\mathcal{T}$  has a unique parent.

VE according to any linear extension of this rooted tree will compute for us  $f_{X_1, Y_1}(x_1, y_1)$ .

Along the way we compute the **sum-product messages**:

$$\delta_{i \rightarrow j}(S_{i,j}) = \sum_{C_i \setminus S_{i,j}} \psi_i(C_i) \prod_{k \in \text{Ne}_{\mathcal{T}}(C_i) \setminus \{C_j\}} \delta_{k \rightarrow i}(S_{i,k}),$$

where  $S_{i,j} = C_i \cap C_j$  (a **separator set**).

The root clique  $C_r$  produces a factor called the **beliefs**:

$$\beta_r(C_r) = \psi_r(C_r) \prod_{k \in \text{Ne}_{\mathcal{T}}(C_r)} \delta_{k \rightarrow r}(S_{r,k}).$$

In our example, we started with a DAG and did VE, so we have  $r = 1$  and get

$$\beta_r(C_r) = \beta_r(x_1, y_1) = \psi_1(x_1, x_2) \delta_{2 \rightarrow 1}(x_1) = f_{X_1, Y_1}(x_1, y_1)$$

We have that:

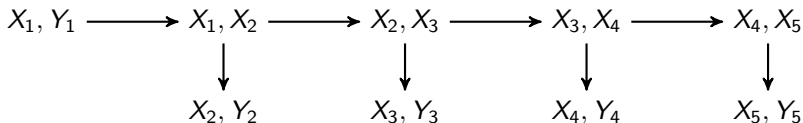
$$\sum_{y_1 \in \text{Val}[Y_1]} \beta_1(x_1, y_1) = f_{X_1}(x_1).$$

This is one of the marginals we wanted!

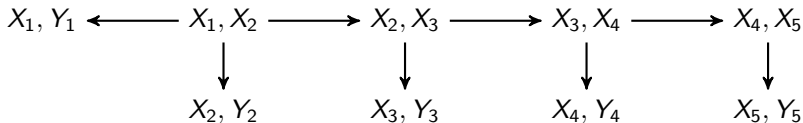
(essentially computed via VE in a fancy language)



$\mathcal{T}$  was used to compute  $f_{X_1}(x_1)$ :



If we change the root node we get a different marginal. The following orientation  $\mathcal{T}'$  can be used to compute  $f_{X_2}(x_2)$ :

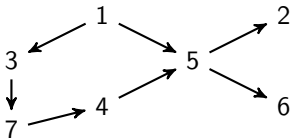


Note that all the messages computed here are exactly the same except for now we compute  $\delta_{1 \rightarrow 2}(x_1)$  and the beliefs  $\beta_2(x_1, x_2)$ .

This means that we can reuse messages computed in our previous run for  $\mathcal{T}$ !

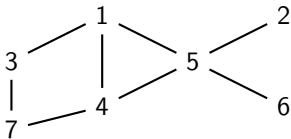
This algorithm works over any clique tree and hence is called the **clique-tree algorithm**.

Suppose we have a distribution  $\mathbf{X} = [X_1, X_2, X_3, X_4, X_5, X_6, X_7]^T$  that is Markov to the following DAG  $\mathcal{G}$ :



$$f_{\mathbf{X}}(\mathbf{x}) = f(x_1)f(x_2|x_5)f(x_3|x_1)f(x_4|x_7)f(x_5|x_1, x_4)f(x_6|x_5)f(x_7|x_3).$$

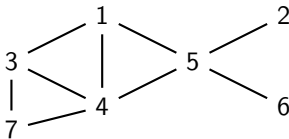
The distribution is also Markov to its **moralization**  $\mathcal{G}^m$ :



$$f_{\mathbf{X}}(\mathbf{x}) = h_1(x_1)h_2(x_2, x_5)h_3(x_1, x_3)h_4(x_4, x_7)h_5(x_1, x_4, x_5)h_6(x_5, x_6)h_7(x_3, x_7).$$

$$h_i = f(x_i | \mathbf{x}_{\text{pa}_{\mathcal{G}}(i)})$$

Similarly, it is Markov to any **chordal cover (triangulation)**  $\mathcal{G}^c$  of  $\mathcal{G}^m$ :



$$f_{\mathbf{x}}(\mathbf{x}) = \psi_{347}(x_3, x_4, x_7) \psi_{134}(x_1, x_3, x_4) \psi_{145}(x_1, x_4, x_5) \psi_{25}(x_2, x_5) \psi_{56}(x_5, x_6).$$

where

$$\psi_{347}(x_3, x_4, x_7) = h_4(x_4, x_7) h_7(x_3, x_7),$$

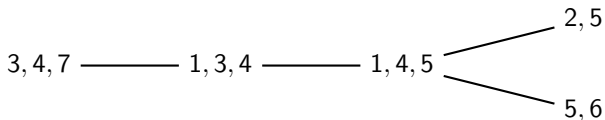
$$\psi_{134}(x_1, x_3, x_4) = h_1(x_1) h_3(x_1, x_3),$$

$$\psi_{145}(x_1, x_4, x_5) = h_5(x_1, x_4, x_5),$$

$$\psi_{25}(x_2, x_5) = h_2(x_2, x_5),$$

$$\psi_{56}(x_5, x_6) = h_6(x_5, x_6).$$

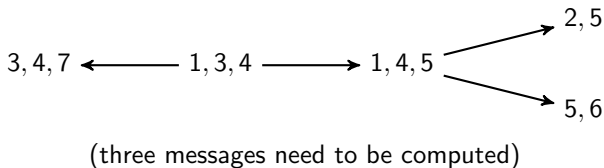
Since  $\mathcal{G}^c$  is chordal we can find a clique-tree for  $\mathcal{G}^c$ , denoted  $\mathcal{T}$ :



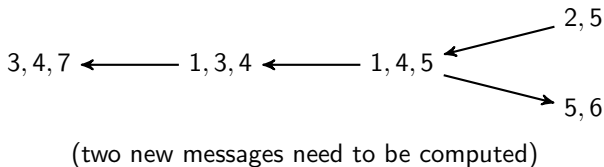
Cliques:  $\mathcal{C} = \{C_1 = \{3, 4, 7\}, C_2 = \{1, 3, 4\}, C_3 = \{1, 4, 5\},$   
 $C_4 = \{2, 5\}, C_5 = \{5, 6\}\}.$

Initial potentials:  $\Phi = \{\psi_{347}, \psi_{134}, \psi_{145}, \psi_{25}, \psi_{56}\}.$

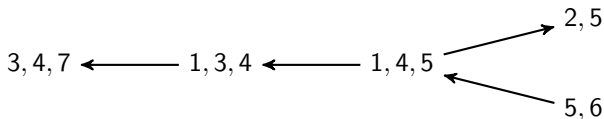
To compute the marginal distributions  $f_{X_1}(x_1)$ ,  $f_{X_3}(x_3)$ ,  $f_{X_4}(x_4)$ , compute the beliefs  $\beta_2(x_1, x_3, x_4)$  using the orientation:



Then to compute  $f_{X_2}(x_2)$  and  $f_{X_5}(x_5)$  use the orientation:

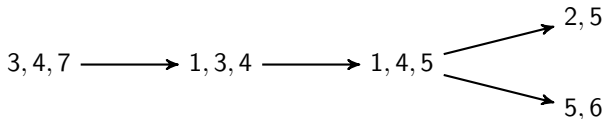


Then to compute  $f_{X_6}(x_6)$  use the orientation:



(one new message needs to be computed)

Finally, to compute  $f_{X_7}(x_7)$  use the orientation:



(one new message needs to be computed)

**7 messages** computed in total.

Using multiple runs of VE to compute all seven marginals would mean computing **16 messages**. 4 messages to compute each of:

$$f_{X_3, X_4, X_7}(x_3, x_4, x_7)$$

$$f_{X_5, X_6}(x_5, x_6)$$

$$f_{X_3, X_4, X_7}(x_1, x_4, x_5)$$

$$f_{X_2, X_5}(x_2, x_5)$$

## Final notes.

- ① The equivalence of factorizations w.r.t. a graph and the markov properties for graphs give us multiple ways to interpret a graphical model (via factorization or conditional independence constraints).
- ② Using these interpretations, we can learn a DAG (up to Markov equivalence) that represents relations in our data-generating distribution
- ③ After fitting the parameters for the DAG model according to the data we can use the DAG for inference

## Final notes.

- ① VE and the Clique-Tree algorithms are exact inference algorithms for computing marginals and posteriors exactly.
- ② Combinatorics of the graph inform the complexity of these methods and speed up computations.
- ③ When the complexity of inference for the learned/constructed graph is too high we can consider **approximate inference algorithms**... To be continued in Module 2!