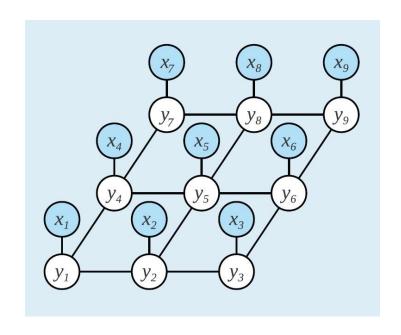


# Probabilistic Graphical Models in Bioinformatics

Lecture 13: Clique trees; MAP inference



#### What you have learned so far and remaining topics



#### Representation

- Bayesian networks
- Markov networks
- Conditional independencies; conditional probability distributions; continuous and discrete data
- Temporal models: HMM and dynamic Bayesian networks

#### Inference

- Variable elimination
- Clique trees
- MAP inference
- Sampling methods: Gibbs sampling and MCMC

#### Learning

- Parameter learning: MLE, Bayesian inference
- Structure learning: constraint-based and score-based methods
- In the case of partially observed data: gradient ascent methods; EM

#### Applications

- Gene regulatory networks
- Motif finding
- Profile HMM for sequence alignment



Exact inference: clique tree



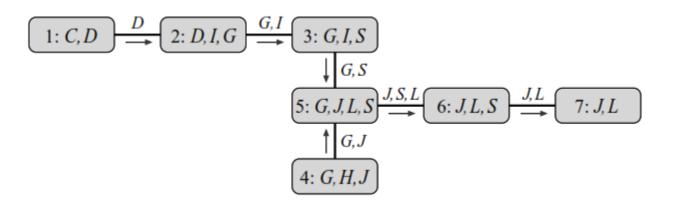
### Exact inference: clique tree

- We already showed how variable elimination can perform exact inference effectively.
  - Key insight: performing local operations on the factors defining the distribution, rather than generating the entire joint distribution.
- Today we present an alternative implementation of the same insight.
  - The algorithm uses a global data structure, called clique tree, for scheduling manipulations of factors.
- In the variable elimination,
  - each step in the computation creates a factor  $\psi_i$  by multiplying existing factors.
  - A variable is them eliminated in  $\psi_i$  to generate a new factor  $\tau_i$ .
- In this section, we consider a factor  $\psi_i$  to be a computational data structure
  - takes "messages"  $au_i$  generated by other factor  $\psi_i$
  - and generates a message  $au_i$  used by another factor  $\psi_l$ .

#### Cluster tree definition



- A cluster graph U for a set of factors  $\Phi$  over  $\mathcal X$  is an undirected graph
  - each of whose nodes *i* is associated with a subset  $C_i \subset \mathcal{X}$
  - each edge between a pair of clusters  $C_i$  and  $C_j$  is associated with a sepset  $S_{i,j} \subseteq C_i \cap C_j$ .
- A cluster graph must be family-preserving
  - Each factor  $\phi \in \Phi$  must be associated with a cluster  $C_i$





### Variable elimination and cluster graph

#### **Example revisited**

- We have seven factors  $\psi_1, ..., \psi_7$ .
- The message  $\tau_1(D)$  is generated from  $\psi_1(C,D)$ .
- $au_1(D)$  participates in the computation of  $\psi_2$ .
- Hence, there is an edge between  $C_1$  and  $C_2$ .

Step	Variable eliminated	Factors used	Variables involved	New factor
1	C	$\phi_C(C)$ , $\phi_D(D,C)$	C,D	$\tau_1(D)$
2	D	$\phi_G(G, I, D), \tau_1(D)$	G, I, D	$\tau_2(G,I)$
3	I	$\phi_{I}(I), \phi_{S}(S, I), \tau_{2}(G, I)$	G, S, I	$ au_3(G,S)$
4	H	$\phi_H(H,G,J)$	H,G,J	$\tau_4(G, J)$
5	G	$\tau_4(G, J),  \tau_3(G, S),  \phi_L(L, G)$	G, J, L, S	$\tau_5(J, L, S)$
6	S	$\tau_5(J,L,S), \phi_J(J,L,S)$	J, L, S	$\tau_6(J,L)$
7	L	$\tau_6(J,L)$	J, L	$ au_7(J)$

### Running intersection property



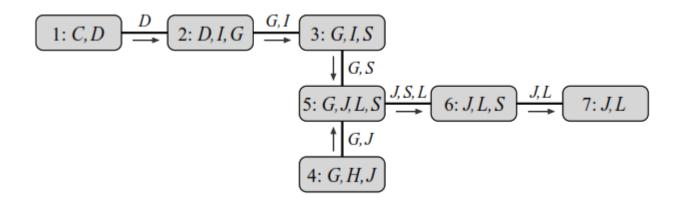
• We say a cluster tree has the running intersection property if  $X \in C_i$  and  $X \in C_j$ , then X is also in **every** cluster in the **unique** path between  $C_i$  and  $C_j$ .

Note that running intersection property implies that  $S_{i,j} = C_i \cap C_j$ .

#### Clique tree



• A cluster tree that satisfies the running intersection property is called a clique tree, also called a junction tree or a joint tree.



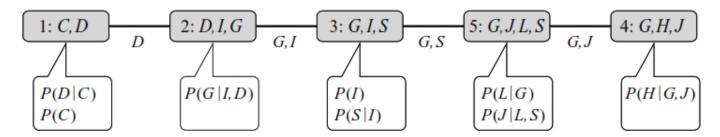


Nonmaximal cliques are absent in this simplified clique tree.

#### Variable elimination in a clique tree



Consider one possible clique tree for the Student network



• The first step is to generate a set of *initial potentials* associated with the different cliques by multiplying the initial factors assigned each clique

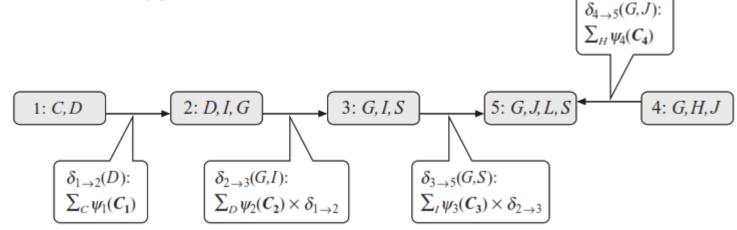
$$\psi_5(J, L, G, S) = \phi_L(L, G) \cdot \phi_J(J, L, S)$$

- Assume our task is to compute P(J). Then we want to do VE so the J is not eliminated.
- We select as our root clique some clique that contains J, for example C5

#### Variable elimination in a clique tree



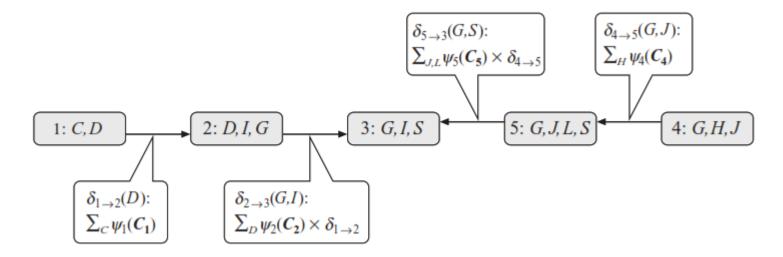
- We then execute the following steps:
  - 1. In  $C_1$ : we eliminate C by performing  $\sum_C \psi_1(C, D)$ . The resulting factor has scope D and send it as a message  $\delta_{1\to 2}(D)$  to  $C_2$ .
  - 2. In  $C_2$ : we eliminate D. The resulting factor has scope G, I and send it as a message  $\delta_{2\to 3}(G,I)$  to  $C_3$ .
  - 3. In  $C_3$ : we eliminate I. The resulting factor has scope G, S and send it as a message  $\delta_{3\to 5}(G,S)$  to  $C_5$ .
  - 4. In  $C_4$ : we eliminate H. The resulting factor has scope G, J and send it as a message  $\delta_{4\to5}(G,J)$  to  $C_5$ .
  - 5. In  $C_5$ : we define  $\beta_5(G,J,S,L) = \delta_{3\to 5}(G,S)$ .  $\delta_{4\to 5}(G,J)$ .  $\psi_5(G,J,S,L)$ . By summing out G, L, and S we obtain P(J).





### Another example: computing P(G)

• Messages sent from  $C_1$  to  $C_2$ , from  $C_2$  to  $C_3$ , and from  $C_4$  to  $C_5$  are the same as the previous run

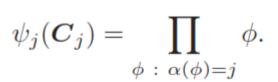


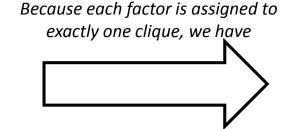
 Ready clique: a clique is ready when it has received all of its incoming messages.

#### Clique-tree message passing



- 1. Let T be a clique tree with the cliques  $C_1, \dots, C_k$ .
- 2. We begin by multiplying the factors assigned to each clique, resulting in our initial potentials.





$$\prod_{\phi} \phi = \prod_{j} \psi_{j}.$$

3. We use clique-tree data structure to pass messages between neighboring cliques, sending all messages toward to the root clique.

$$\delta_{i\to j} = \sum_{\boldsymbol{C}_i - \boldsymbol{S}_{i,j}} \psi_i \cdot \prod_{k \in (\mathrm{Nb}_i - \{j\})} \delta_{k\to i}.$$

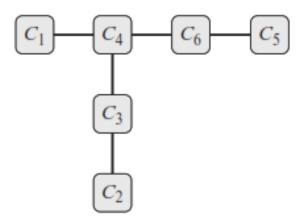
4. When the root clique has received all messages, it multiplies them with its own initial potential. The result is a factor called the *beliefs*, denoted  $\beta_r(C_r)$ . It represents

$$\tilde{P}_{\Phi}(C_r) = \sum_{\mathcal{X} - C_r} \prod_{\phi} \phi.$$

#### Algorithm 10.1 Upward pass of variable elimination in clique tree

```
Procedure CTree-SP-Upward (
            // Set of factors
             // Clique tree over \Phi
            // Initial assignment of factors to cliques
   C_r // Some selected root clique
   Initialize-Cliques
  while \boldsymbol{C}_r is not ready
      Let C_i be a ready clique
      \delta_{i \to p_r(i)}(\boldsymbol{S}_{i,p_r(i)}) \leftarrow \text{SP-Message}(i,p_r(i))
  \beta_r \leftarrow \psi_r \cdot \prod_{k \in \text{Nb}_{C_r}} \delta_{k \to r}
  return \beta_r
Procedure Initialize-Cliques (
   for each clique C_i
      \psi_i(\mathbf{C}_i) \leftarrow \prod_{\phi_i : \alpha(\phi_i)=i} \phi_i
Procedure SP-Message (
           // sending clique
         // receiving clique
 \psi(\boldsymbol{C}_i) \leftarrow \psi_i \cdot \prod_{k \in (\mathrm{Nb}_i - \{j\})} \delta_{k \to i}\tau(\boldsymbol{S}_{i,j}) \leftarrow \sum_{\boldsymbol{C}_i - \boldsymbol{S}_{i,j}} \psi(\boldsymbol{C}_i)
  return \tau(S_{i,j})
```





Assume we have selected  $C_6$  as our root clique. There are multiple legitimate orderings.

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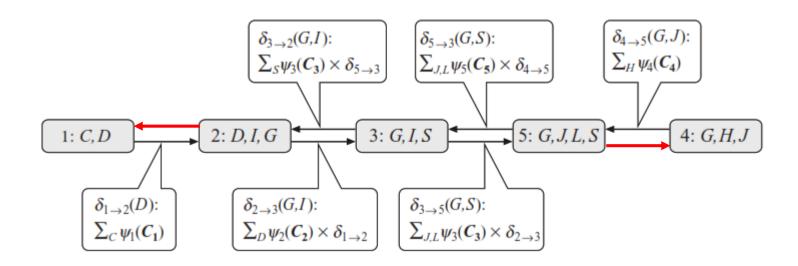
### Clique tree calibration



- We showed how to use the same clique tree to compute the probability of any variable in the input variables.
- We often wish to estimate the probability of a large number of variables.
  - For example, we want to compute the probability of several possible diseases in medical diagnosis setting.
- Consider the task of computing the posterior distribution over each random variable in the network
  - Naïve approach: doing inference separately for each variable.
  - Better approach: **sum-product belief propagation** algorithm.



### Sum-Product Belief Propagation





### Sum-Product Belief Propagation

#### Algorithm 10.2 Calibration using sum-product message passing in a clique tree

```
Procedure CTree-SP-Calibrate ( \Phi, // Set of factors \mathcal{T} // Clique tree over \Phi )

1 Initialize-Cliques while exist i,j such that i is ready to transmit to j \delta_{i \to j}(S_{i,j}) \leftarrow \text{SP-Message}(i,j) for each clique i \beta_i \leftarrow \psi_i \cdot \prod_{k \in \text{Nb}_i} \delta_{k \to i} return \{\beta_i\}
```

#### Scheduling for sending messages:

- Asynchronously
- Upward pass and downward pass



### Constructing a clique tree



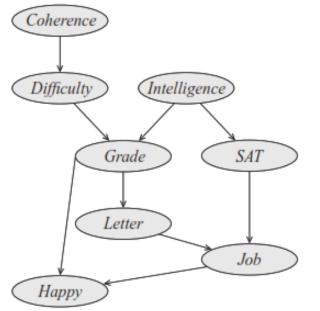
#### Constructing a clique tree

• How do we construct a clique tree for a set of factors, or equivalently, for its underlying undirected graph  $H_{\Phi}$ ? (See section 9.4.2.1 for the definition of  $H_{\Phi}$ .)

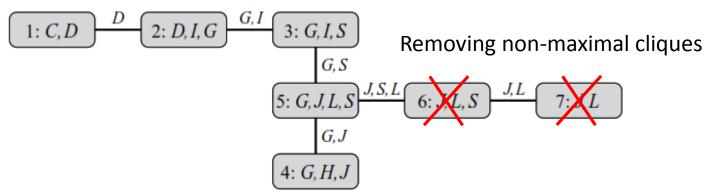
- Two basic approaches
  - Clique trees from variable elimination
  - Clique trees from chordal graphs

#### Clique trees from variable elimination





Step	Variable eliminated	Factors used	Variables involved	New factor
1	C	$\phi_C(C)$ , $\phi_D(D,C)$	C,D	$\tau_1(D)$
2	D	$\phi_G(G, I, D), \tau_1(D)$	G, I, D	$\tau_2(G,I)$
3	I	$\phi_{I}(I), \phi_{S}(S, I), \tau_{2}(G, I)$	G, S, I	$ au_3(G,S)$
4	H	$\phi_H(H,G,J)$	H,G,J	$\tau_4(G,J)$
5	G	$\tau_4(G, J),  \tau_3(G, S),  \phi_L(L, G)$	G, J, L, S	$\tau_5(J, L, S)$
6	S	$\tau_5(J,L,S), \phi_J(J,L,S)$	J, L, S	$\tau_6(J, L)$
7	L	$\tau_6(J,L)$	J, L	$\tau_7(J)$

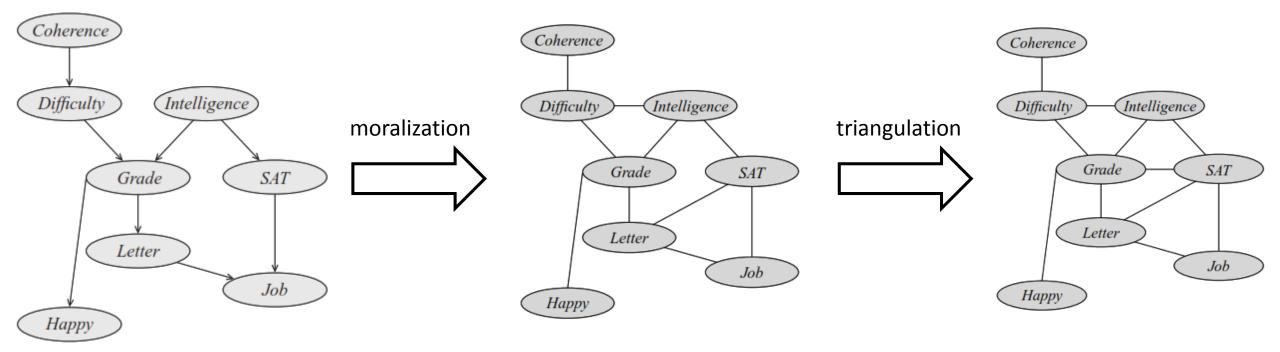


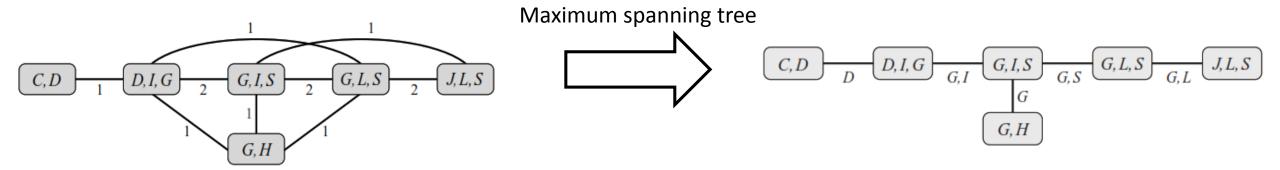
- Each cluster is defined by  $C_i = \text{Scope}[\psi_i]$
- There is an edge between  $C_i$  and  $C_i$  of the message  $\tau_i$  is used in computing  $\psi_i$

### Clique trees from chordal graphs



- 1. Find the minimal Markov network I-map  $H_{\Phi}$ 
  - For a Bayesian network, in the case without evidence, it is simply the moralized graph
- 2. Triangulation to obtain a chordal graph H\*
- 3. Find cliques in  $H^*$ , and make each one a node in the cluster graph
- 4. Run the maximum spanning tree algorithm on the cluster graph to construct a tree







#### MAP inference



#### MAP inference

- A MAP query aims to find the most likely assignment to all of the (non-evidence) variables.
- A marginal MAP query aims to find the most likely assignment to a subset of the variables, marginalizing over the rest.
- Examples:
  - Speech recognition: to decode the most likely utterance given the (noisy) acoustic signal.
- For addressing MAP queries, we need to compute

$$\xi^{map} = \arg\max_{\xi} P_{\Phi}(\xi) = \arg\max_{\xi} \frac{1}{Z} \tilde{P}_{\Phi}(\xi) = \arg\max_{\xi} \tilde{P}_{\Phi}(\xi).$$

#### Max-Product Variable Elimination



• Consider the Bayesian network  $A \rightarrow B$ . Assume our goal is to compute:

$$\max_{a,b} P(a,b) = \max_{a,b} P(a)P(b \mid a)$$
$$= \max_{a} \max_{b} P(a)P(b \mid a).$$

• For any choice of a, the value of B must be chosen so as to maximize  $P(b \mid a)$ . We must also choose the value of A appropriately

$$\max_{b} P(a)P(b \mid a) = P(a) \max_{b} P(b \mid a).$$

• Let  $\phi(a)$  denote  $\max_b P(b \mid a)$ . For example, for the following assignments

•  $\phi(a^1) = \max_b P(b \mid a^1) = 0.55$  and  $\phi(a^0) = \max_b P(b \mid a^0) = 0.9$ . Finally

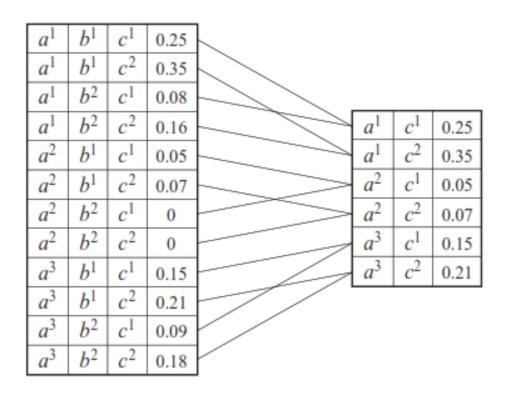
$$\max_{a} P(a)\phi(a) = \max [0.4 \cdot 0.9, 0.6 \cdot 0.55] = 0.36.$$





Let  $\phi(X,Y)$  be a factor. We define the factor maximization of Y in  $\phi$  as

$$\psi(\boldsymbol{X}) = \max_{Y} \phi(\boldsymbol{X}, Y).$$



Algorithm 13.1 Variable elimination algorithm for MAP. The algorithm can be used both in its max-product form, as shown, or in its max-sum form, replacing factor product with factor addition.

```
Procedure Max-Product-VE (
         // Set of factors over X
        // Ordering on X
  Let X_1, \ldots, X_k be an ordering of X such that
     X_i \prec X_j iff i < j
     for i = 1, \ldots, k
        (\Phi, \phi_{X_i}) \leftarrow \text{Max-Product-Eliminate-Var}(\Phi, X_i)
     x^* \leftarrow \text{Traceback-MAP}(\{\phi_{X_i} : i = 1, ..., k\})
     return x^*, \Phi // \Phi contains the probability of the MAP
  Procedure Max-Product-Eliminate-Var (
      Φ. // Set of factors
           // Variable to be eliminated
     \Phi' \leftarrow \{\phi \in \Phi : Z \in Scope[\phi]\}
     \Phi'' \leftarrow \Phi - \Phi'
     \psi \leftarrow \prod_{\phi \in \Phi'} \phi
     \tau \leftarrow \max_{Z} \psi
     return (\Phi'' \cup \{\tau\}, \psi)
  Procedure Traceback-MAP (
      \{\phi_{X_i} : i = 1, ..., k\}
     for i = k, ..., 1
       u_i \leftarrow (x_{i+1}^*, \dots, x_k^*) \langle Scope[\phi_{X_i}] - \{X_i\} \rangle
           // The maximizing assignment to the variables eliminated after
             X_i
       x_i^* \leftarrow \arg \max_{x_i} \phi_{X_i}(x_i, u_i)
           ||x_i^*| is chosen so as to maximize the corresponding entry in
             the factor, relative to the previous choices u_i
     return x
```



Theorem 13.4

The algorithm of algorithm 13.1 returns

$$x^* = \arg \max_{x} \prod_{\phi \in \Phi} \phi,$$

and  $\Phi$ , which contains a single factor of empty scope whose value is:

$$\max_{x} \prod_{\phi \in \Phi} \phi$$
.

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## A run of max-product variable elimination TEHRAN



Step	Variable	Factors	Intermediate	New
	eliminated	used	factor	factor
1	S	$\phi_S(I,S)$	$\psi_1(I,S)$	$ au_1(I)$
2	I	$\phi_I(I), \phi_G(G, I, D), \tau_1(I)$	$\psi_2(G,I,D)$	$ au_2(G,D)$
3	D	$\phi_D(D),  \tau_2(G,D)$	$\psi_3(G,D)$	$ au_3(G)$
4	L	$\phi_L(L,G)$	$\psi_4(L,G)$	$\tau_4(G)$
5	G	$\tau_4(G)$ , $\tau_3(G)$	$\psi_5(G)$	$ au_5(\emptyset)$

For example, the first step would compute

$$\tau_1(I) = \max_s \phi_S(I, s)$$

The final factor,  $\tau_5(0)$ , is simply a number, whose value is

$$\max_{S,I,D,L,G} P(S,I,D,L,G).$$

#### Decoding or finding the most probable assignment

- Traceback of the solution
- We begin by computing g

$$g^* = \arg \max_g \psi_5(g)$$

$$l^* = \arg \max_l \psi_4(g^*, l)$$

$$d^* = \arg \max_d \psi_3(g^*, d)$$

$$i^* = \arg \max_i \psi_2(g^*, i, d^*)$$

$$s^* = \arg \max_s \psi_1(i^*, s).$$

Step	Variable	Factors	Intermediate	New
	eliminated	used	factor	factor
1	S	$\phi_S(I,S)$	$\psi_1(I,S)$	$\tau_1(I)$
2	I	$\phi_I(I)$ , $\phi_G(G, I, D)$ , $\tau_1(I)$	$\psi_2(G,I,D)$	$ au_2(G,D)$
3	D	$\phi_D(D)$ , $\tau_2(G,D)$	$\psi_3(G,D)$	$ au_3(G)$
4	L	$\phi_L(L,G)$	$\psi_4(L,G)$	$\tau_4(G)$
5	G	$\tau_4(G)$ , $\tau_3(G)$	$\psi_5(G)$	$ au_5(\emptyset)$