Today’s lecture

- Markov random fields
  1. Factor graphs
  2. Bayesian networks $\Rightarrow$ Markov random fields (*moralization*)

- Exact inference
  1. Worst-case complexity of probabilistic inference
  2. Elimination algorithm
  3. Running-time analysis of elimination algorithm (*treewidth*)
Undirected graphical models

- An alternative representation for joint distributions is as an **undirected graphical model**

- As in BNs, we have one node for each random variable

- Rather than CPDs, we specify (non-negative) **potential functions** over sets of variables associated with cliques $C$ of the graph,

  $$
  p(x_1, \ldots, x_n) = \frac{1}{Z} \prod_{c \in C} \phi_c(x_c)
  $$

  $Z$ is the **partition function** and normalizes the distribution:

  $$
  Z = \sum_{\hat{x}_1, \ldots, \hat{x}_n} \prod_{c \in C} \phi_c(\hat{x}_c)
  $$

  - Like CPD’s, $\phi_c(x_c)$ can be represented as a table, but it is not normalized

  - Also known as **Markov random fields** (MRFs) or Markov networks
Higher-order potentials

- The examples so far have all been pairwise MRFs, involving only node potentials $\phi_i(X_i)$ and pairwise potentials $\phi_{i,j}(X_i, X_j)$.
- Often we need higher-order potentials, e.g.

$$\phi(x, y, z) = 1[x + y + z \geq 1],$$

where $X, Y, Z$ are binary, enforcing that at least one of the variables takes the value 1.

- Although Markov networks are useful for understanding independencies, they hide much of the distribution’s structure:

Does this have pairwise potentials, or one potential for all 4 variables?
Factor graphs

- $G$ does not reveal the structure of the distribution: maximum cliques vs. subsets of them.

- A **factor graph** is a bipartite undirected graph with variable nodes and factor nodes. Edges are only between the variable nodes and the factor nodes.

- Each factor node is associated with a single potential, whose scope is the set of variables that are neighbors in the factor graph.

- The distribution is same as the MRF – this is just a different data structure.
Example: Low-density parity-check codes

- Error correcting codes for transmitting a message over a noisy channel (invented by Gallager in the 1960’s, then re-discovered in 1996)

- Each of the top row factors enforce that its variables have even parity:

  \[ f_A(Y_1, Y_2, Y_3, Y_4) = 1 \text{ if } Y_1 \otimes Y_2 \otimes Y_3 \otimes Y_4 = 0, \text{ and } 0 \text{ otherwise} \]

- Thus, the only assignments \( Y \) with non-zero probability are the following (called codewords):

  3 bits encoded using 6 bits

  000000, 011001, 110010, 101011, 111100, 100101, 001110, 010111

- \( f_i(Y_i, X_i) = p(X_i \mid Y_i) \), the likelihood of a bit flip according to noise model
The decoding problem for LDPCs is to find

$$\arg\max_y p(y \mid x)$$

This is called the **maximum a posteriori** (MAP) assignment.

Since $Z$ and $p(x)$ are constants with respect to the choice of $y$, can equivalently solve (taking the log of $p(y, x)$):

$$\arg\max_y \sum_{c \in C} \theta_c(y_c, x_c),$$

where $\theta_c(x_c) = \log \phi_c(y_c, x_c)$

This is a discrete optimization problem!
What is the equivalent Markov network for a hidden Markov model?

Many inference algorithms are more conveniently given for undirected models – this shows how they can be applied to Bayesian networks.
Moralization of Bayesian networks

- Procedure for converting a Bayesian network into a Markov network

- The **moral graph** $\mathcal{M}[G]$ of a BN $G = (V, E)$ is an undirected graph over $V$ that contains an undirected edge between $X_i$ and $X_j$ if
  1. there is a directed edge between them (in either direction)
  2. $X_i$ and $X_j$ are both parents of the same node

(term historically arose from the idea of “marrying the parents” of the node)

- The addition of the moralizing edges leads to the loss of some independence information, e.g., $A \rightarrow C \leftarrow B$, where $A \perp B$ is lost
Converting BNs to Markov networks

1. Moralize the directed graph to obtain the undirected graphical model:

\[
\begin{align*}
\text{Moralization}
\end{align*}
\]

2. Introduce one potential function for each CPD:

\[
\phi_i(x_i, x_{pa(i)}) = p(x_i | x_{pa(i)})
\]

- So, converting a hidden Markov model to a Markov network is simple:

- For variables having $>1$ parent, factor graph notation is useful
Today we consider exact inference in graphical models.

In particular, we focus on conditional probability queries,

\[ p(Y|E = e) = \frac{p(Y, e)}{p(e)} \]

(e.g., the probability of a patient having a disease given some observed symptoms)

Let \( W = X - Y - E \) be the random variables that are neither the query nor the evidence. Each of these joint distributions can be computed by marginalizing over the other variables:

\[ p(Y, e) = \sum_w p(Y, e, w), \quad p(e) = \sum_y p(y, e) \]

Naively marginalizing over all unobserved variables requires an exponential number of computations.

Does there exist a more efficient algorithm?
Computational complexity of probabilistic inference

Here we show that, unless P=NP, there does not exist a more efficient algorithm.

We show this by reducing 3-SAT, which is NP-hard, to probabilistic inference in Bayesian networks.

3-SAT asks about the *satisfiability* of a logical formula defined on \( n \) literals \( Q_1, \ldots, Q_n \), e.g.

\[
(\neg Q_3 \lor \neg Q_2 \lor Q_3) \land (Q_2 \lor \neg Q_4 \lor \neg Q_5) \ldots
\]

Each of the disjunction terms is called a *clause*, e.g.

\[
C_1(q_1, q_2, q_3) = \neg q_3 \lor \neg q_2 \lor q_3
\]

In 3-SAT, each clause is defined on at most 3 literals.

Our reduction also proves that inference in Markov networks is NP-hard (why?)
Reducing satisfiability to MAP inference

- **Input:** 3-SAT formula with $n$ literals $Q_1, \ldots, Q_n$ and $m$ clauses $C_1, \ldots, C_m$

  - One variable $Q_i \in \{0, 1\}$ for each literal, $p(Q_i = 1) = 0.5$.
  - One variable $C_i \in \{0, 1\}$ for each clause, whose parents are the literals used in the clause. $C_i = 1$ if the clause is satisfied, and 0 otherwise:

    $$p(C_i = 1 \mid q_{pa(i)}) = 1[C_i(q_{pa(i)})]$$

  - Variable $X$ which is 1 if all clauses satisfied, and 0 otherwise:

    $$p(A_i = 1 \mid pa(A_i)) = 1[pa(A_i) = 1], \text{ for } i = 1, \ldots, m - 2$$
    $$p(X = 1 \mid a_{m-2}, c_m) = 1[a_{m-2} = 1, c_m = 1]$$
Reducing satisfiability to MAP inference

- **Input:** 3-SAT formula with \( n \) literals \( Q_1, \ldots, Q_n \) and \( m \) clauses \( C_1, \ldots, C_m \)

\[
\begin{align*}
Q_1 & \quad Q_2 & \quad Q_3 & \quad Q_4 & \quad \cdots & \quad Q_n \\
C_1 & \quad C_2 & \quad C_3 & \quad \cdots & \quad C_{m-1} & \quad C_m \\
A_1 & \quad A_2 & \quad \cdots & \quad A_{m-2} & \quad X
\end{align*}
\]

- \( p(q, c, a, X = 1) = 0 \) for any assignment \( q \) which does not satisfy all clauses
- \( p(Q = q, C = 1, A = 1, X = 1) = \frac{1}{2^n} \) for any satisfying assignment \( q \)

Thus, we can find a satisfying assignment (whenever one exists) by constructing this BN and finding the maximum a posteriori (MAP) assignment:

\[
\arg\max_{q, c, a} p(Q = q, C = c, A = a \mid X = 1)
\]

- This proves that MAP inference in Bayesian networks and MRFs is NP-hard
**Input:** 3-SAT formula with $n$ literals $Q_1, \ldots, Q_n$ and $m$ clauses $C_1, \ldots, C_m$

$$p(X = 1) = \sum_{q, c, a} p(Q = q, C = c, A = a, X = 1)$$ is equal to the number of satisfying assignments times $\frac{1}{2^n}$

Thus, $p(X = 1) > 0$ if and only if the formula has a satisfying assignment

This shows that *marginal inference* is also NP-hard
NP-hardness simply says that there exist difficult inference problems.

Real-world inference problems are not necessarily as hard as these worst-case instances.

The reduction from SAT created a very complex Bayesian network:

Some graphs are easy to do inference in! For example, inference in hidden Markov models and other tree-structured graphs can be performed in linear time.
Variable elimination (VE)

- Exact algorithm for probabilistic inference in any graphical model
- Running time will depend on the graph structure
- Uses dynamic programming to circumvent enumerating all assignments
- First we introduce the concept for computing marginal probabilities, $p(X_i)$, in Bayesian networks
- After this, we will generalize to MRFs and conditional queries
Basic idea

- Suppose we have a simple chain, $A \rightarrow B \rightarrow C \rightarrow D$, and we want to compute $p(D)$
- $p(D)$ is a set of values, $\{p(D = d), d \in \text{Val}(D)\}$. Algorithm computes sets of values at a time – an entire distribution
- By the chain rule and conditional independence, the joint distribution factors as

$$p(A, B, C, D) = p(A)p(B \mid A)p(C \mid B)p(D \mid C)$$

- In order to compute $p(D)$, we have to marginalize over $A, B, C$:

$$p(D) = \sum_{a, b, c} p(A = a, B = b, C = c, D)$$
Let’s be a bit more explicit...

\[
\begin{align*}
& P(a^1) \quad P(b^1 | a^1) \quad P(c^1 | b^1) \quad P(d^1 | c^1) \\
& + \quad P(a^2) \quad P(b^1 | a^2) \quad P(c^1 | b^1) \quad P(d^1 | c^1) \\
& + \quad P(a^1) \quad P(b^2 | a^1) \quad P(c^1 | b^2) \quad P(d^1 | c^1) \\
& + \quad P(a^2) \quad P(b^2 | a^2) \quad P(c^1 | b^2) \quad P(d^1 | c^1) \\
& + \quad P(a^1) \quad P(b^1 | a^1) \quad P(c^2 | b^1) \quad P(d^1 | c^2) \\
& + \quad P(a^2) \quad P(b^1 | a^2) \quad P(c^2 | b^1) \quad P(d^1 | c^2) \\
& + \quad P(a^1) \quad P(b^2 | a^1) \quad P(c^2 | b^2) \quad P(d^1 | c^2) \\
& + \quad P(a^2) \quad P(b^2 | a^2) \quad P(c^2 | b^2) \quad P(d^1 | c^2)
\end{align*}
\]

- There is structure to the summation, e.g., repeated \( P(c^1 | b^1) P(d^1 | c^1) \)
- Let’s modify the computation to first compute
  \[
P(a^1)P(b^1 | a^1) + P(a^2)P(b^1 | a^2)
\]
Let’s be a bit more explicit...

- Let’s modify the computation to first compute
  \[ P(a^1)P(b^1|a^1) + P(a^2)P(b^1|a^2) \]
  and
  \[ P(a^1)P(b^2|a^1) + P(a^2)P(b^2|a^2) \]

- Then, we get
  \[
  \begin{align*}
  & (P(a^1)P(b^1|a^1) + P(a^2)P(b^1|a^2)) \quad P(c^1|b^1) \quad P(d^1|c^1) \\
  + & (P(a^1)P(b^2|a^1) + P(a^2)P(b^2|a^2)) \quad P(c^1|b^2) \quad P(d^1|c^1) \\
  + & (P(a^1)P(b^1|a^1) + P(a^2)P(b^1|a^2)) \quad P(c^2|b^1) \quad P(d^1|c^2) \\
  + & (P(a^1)P(b^2|a^1) + P(a^2)P(b^2|a^2)) \quad P(c^2|b^2) \quad P(d^1|c^2) \\
  \end{align*}
  \]

- We define \( \tau_1 : \text{Val}(B) \to \mathbb{R} \), \quad \tau_1(b^i) = P(a^1)P(b^i|a^1) + P(a^2)P(b^i|a^2) \)
Let’s be a bit more explicit...

- We now have

\[
\begin{align*}
\tau_1(b^1) & \quad P(c^1 | b^1) & \quad P(d^1 | c^1) \\
+ \tau_1(b^2) & \quad P(c^1 | b^2) & \quad P(d^1 | c^1) \\
+ \tau_1(b^1) & \quad P(c^2 | b^1) & \quad P(d^1 | c^2) \\
+ \tau_1(b^2) & \quad P(c^2 | b^2) & \quad P(d^1 | c^2)
\end{align*}
\]

- We can once more reverse the order of the product and the sum and get

\[
\begin{align*}
\tau_1(b^1) & \quad P(c^1 | b^1) & \quad P(d^2 | c^1) \\
+ \tau_1(b^2) & \quad P(c^1 | b^2) & \quad P(d^2 | c^1) \\
+ \tau_1(b^1) & \quad P(c^2 | b^1) & \quad P(d^2 | c^2) \\
+ \tau_1(b^2) & \quad P(c^2 | b^2) & \quad P(d^2 | c^2)
\end{align*}
\]

- There are still other repeated computations!
Let’s be a bit more explicit...

- We define $\tau_2 : \text{Val}(C) \rightarrow \mathbb{R}$, with

  $$
  \tau_2(c^1) = \tau_1(b^1)P(c^1|b^1) + \tau_1(b^2)P(c^1|b^2)
  $$

  $$
  \tau_2(c^2) = \tau_1(b^1)P(c^2|b^1) + \tau_1(b^2)P(c^2|b^2)
  $$

- Now we can compute the marginal $p(D)$ as

  $$
  \begin{align*}
  \tau_2(c^1) \cdot P(d^1 | c^1) + \tau_2(c^2) \cdot P(d^1 | c^2) \\
  \tau_2(c^1) \cdot P(d^2 | c^1) + \tau_2(c^2) \cdot P(d^2 | c^2)
  \end{align*}
  $$
What did we just do?

- Our goal was to compute

\[ p(D) = \sum_{a,b,c} p(a, b, c, D) = \sum_{a,b,c} p(a)p(b \mid a)p(c \mid b)p(D \mid c) \]

\[ = \sum_c \sum_b \sum_a p(D \mid c)p(c \mid b)p(b \mid a)p(a) \]

- We can push the summations inside to obtain:

\[ p(D) = \sum_c p(D \mid c) \sum_b p(c \mid b) \sum_a p(b \mid a)p(a) \]

\[ \psi_1(a,b) \tau_1(b) \]

- Let’s call \( \psi_1(A, B) = P(A)P(B \mid A) \). Then, \( \tau_1(B) = \sum_a \psi_1(a, B) \)

- Similarly, let \( \psi_2(B, C) = \tau_1(B)P(C \mid B) \). Then, \( \tau_2(C) = \sum_b \psi_1(b, C) \)

- This procedure is dynamic programming: computation is inside out instead of outside in
Inference in a chain

- Generalizing the previous example, suppose we have a chain $X_1 \rightarrow X_2 \rightarrow \cdots \rightarrow X_n$ where each variable has $k$ states.

- In Problem Set 1 (question 2), you gave an algorithm to compute $p(X_i)$, for $k = 2$.

- For $i = 1$ up to $n - 1$, compute (and cache)
  
  $$p(X_{i+1}) = \sum_{x_i} p(X_{i+1} | x_i)p(x_i)$$

- Each update takes $k^2$ time (why?)

- The total running time is $\mathcal{O}(nk^2)$.

- In comparison, naively marginalizing over all latent variables has complexity $\mathcal{O}(k^n)$.

- We did inference over the joint without ever explicitly constructing it!
Summary so far

- Worst-case analysis says that marginal inference is NP-hard
- In practice, due to the structure of the Bayesian network, we can cache computations that are otherwise computed exponentially many times
- This depends on our having a good variable elimination ordering
Sum-product inference task

- We want to give an algorithm to compute $p(Y)$ for BNs and MRFs.
- This can be reduced to the following **sum-product** inference task:

  \[
  \tau(y) = \sum_{z} \prod_{\phi \in \Phi} \phi(z_{\text{Scope}[\phi] \cap z}, y_{\text{Scope}[\phi] \cap Y}) \quad \forall y,
  \]

  where $\Phi$ is a set of factors or potentials.

- For a BN, $\Phi$ is given by the conditional probability distributions for all variables,

  \[
  \Phi = \{\phi_{X_i}\}_{i=1}^n = \{p(X_i | X_{\text{Pa}(X_i)})\}_{i=1}^n,
  \]

  and where we sum over the set $Z = X - Y$.

- For Markov networks, the factors $\Phi$ correspond to the set of potentials which we earlier called $C$.
  - Sum-product returns an unnormalized distribution, so we divide by $\sum_y \tau(y)$. 


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Factor marginalization

- Let $\phi(X, Y)$ be a factor where $X$ is a set of variables and $Y \notin X$
- **Factor marginalization** of $\phi$ over $Y$ (also called “summing out $Y$ in $\phi$”) gives a new factor:

$$\tau(X) = \sum_Y \phi(X, Y)$$

For example,

<p>| | | | |</p>
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</tbody>
</table>

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Inference and Representation
Lecture 4, Sept. 29, 2015
27 / 41
Order the variables $Z$ (called the elimination ordering)

Iteratively marginalize out variable $Z_i$, one at a time

For each $i$,

1. Multiply all factors that have $Z_i$ in their scope, generating a new product factor
2. Marginalize this product factor over $Z_i$, generating a smaller factor
3. Remove the old factors from the set of all factors, and add the new one
Algorithm 9.1 Sum-Product Variable Elimination algorithm

Procedure Sum-Product-Variable-Elimination ( 
    \( \Phi \),  // Set of factors
    \( Z \),  // Set of variables to be eliminated
    \(<\)  // Ordering on \( Z \)
)

1. Let \( Z_1, \ldots, Z_k \) be an ordering of \( Z \) such that
2. \( Z_i < Z_j \) iff \( i < j \)
3. for \( i = 1, \ldots, k \)
4. \( \Phi \leftarrow \text{Sum-Product-Eliminate-Var}(\Phi, Z_i) \)
5. \( \phi^* \leftarrow \prod_{\phi \in \Phi} \phi \)
6. return \( \phi^* \)

Procedure Sum-Product-Eliminate-Var ( 
    \( \Phi \),  // Set of factors
    \( Z \)  // Variable to be eliminated
)

1. \( \Phi' \leftarrow \{ \phi \in \Phi : Z \in \text{Scope}[\phi] \} \)
2. \( \Phi'' \leftarrow \Phi \setminus \Phi' \)
3. \( \psi \leftarrow \prod_{\phi \in \Phi'} \phi \)
4. \( \tau \leftarrow \sum_{Z} \psi \)
5. return \( \Phi'' \cup \{ \tau \} \)
• What is \( p(\text{Job})? \) Joint distribution factorizes as:

\[
p(C, D, I, G, S, L, H, J) = p(C)p(D|C)p(I)p(G|D, I)p(L|G)P(S|I)P(J|S, L)p(H|J, G)
\]

with factors

\[
\Phi = \{\phi_C(C), \phi_D(C, D), \phi_I(I), \phi_G(G, D, I), \phi_L(L, G), \\
\phi_S(S, I), \phi_J(J, S, L), \phi_H(H, J, G)\}
\]

• Let’s do variable elimination with ordering \( \{C, D, I, H, G, S, L\} \) on the board!
Elimination ordering

- We can pick any order we want, but some orderings introduce factors with much larger scope

<table>
<thead>
<tr>
<th>Step</th>
<th>Variable eliminated</th>
<th>Factors used</th>
<th>Variables involved</th>
<th>New factor</th>
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<td>1</td>
<td>C</td>
<td>$\phi_C(C), \phi_D(D,C)$</td>
<td>C, D</td>
<td>$\tau_1(D)$</td>
</tr>
<tr>
<td>2</td>
<td>D</td>
<td>$\phi_G(G,I,D), \tau_1(D)$</td>
<td>G, I, D</td>
<td>$\tau_2(G,I)$</td>
</tr>
<tr>
<td>3</td>
<td>I</td>
<td>$\phi_I(I), \phi_S(S,I), \tau_2(G,I)$</td>
<td>G, S, I</td>
<td>$\tau_3(G,S)$</td>
</tr>
<tr>
<td>4</td>
<td>H</td>
<td>$\phi_H(H,G,J)$</td>
<td>H, G, J</td>
<td>$\tau_4(G,J)$</td>
</tr>
<tr>
<td>5</td>
<td>G</td>
<td>$\tau_4(G,J), \tau_3(G,S), \phi_L(L,G)$</td>
<td>G, J, L, S</td>
<td>$\tau_5(J,L,S)$</td>
</tr>
<tr>
<td>6</td>
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<td>$\tau_5(J,L,S), \phi_J(J,L,S)$</td>
<td>J, L, S</td>
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<tr>
<td>7</td>
<td>L</td>
<td>$\tau_6(J,L)$</td>
<td>J, L</td>
<td>$\tau_7(J)$</td>
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</table>

- Alternative ordering...

<table>
<thead>
<tr>
<th>Step</th>
<th>Variable eliminated</th>
<th>Factors used</th>
<th>Variables involved</th>
<th>New factor</th>
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<tbody>
<tr>
<td>1</td>
<td>G</td>
<td>$\phi_G(G,I,D), \phi_L(L,G), \phi_H(H,G,J)$</td>
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<td>S, I, D, L, J, H</td>
<td>$\tau_2(D,L,S,J,H)$</td>
</tr>
<tr>
<td>3</td>
<td>S</td>
<td>$\phi_J(J,L,S), \tau_2(D,L,S,J,H)$</td>
<td>D, L, S, J, H</td>
<td>$\tau_3(D,L,J,H)$</td>
</tr>
<tr>
<td>4</td>
<td>L</td>
<td>$\tau_3(D,L,J,H)$</td>
<td>D, L, J, H</td>
<td>$\tau_4(D,J,H)$</td>
</tr>
<tr>
<td>5</td>
<td>H</td>
<td>$\tau_4(D,J,H)$</td>
<td>D, J, H</td>
<td>$\tau_5(D,J)$</td>
</tr>
<tr>
<td>6</td>
<td>C</td>
<td>$\tau_5(D,J), \phi_D(D,C)$</td>
<td>D, J, C</td>
<td>$\tau_6(D,J)$</td>
</tr>
<tr>
<td>7</td>
<td>D</td>
<td>$\tau_6(D,J)$</td>
<td>D, J</td>
<td>$\tau_7(J)$</td>
</tr>
</tbody>
</table>
How to introduce evidence?

- Recall that our original goal was to answer *conditional* probability queries,

\[ p(Y|E = e) = \frac{p(Y, e)}{p(e)} \]

- Apply variable elimination algorithm to the task of computing \( P(Y, e) \)

- Replace each factor \( \phi \in \Phi \) that has \( E \cap \text{Scope}[\phi] \neq \emptyset \) with

\[ \phi'(x_{\text{Scope}[\phi] - E}) = \phi(x_{\text{Scope}[\phi] - E}, e_{E \cap \text{Scope}[\phi]}) \]

- Then, eliminate the variables in \( X - Y - E \). The returned factor \( \phi^*(Y) \) is \( p(Y, e) \)

- To obtain the conditional \( p(Y | e) \), normalize the resulting product of factors – the normalization constant is \( p(e) \)
Let $n$ be the number of variables, and $m$ the number of initial factors.

At each step, we pick a variable $X_i$ and multiply all factors involving $X_i$, resulting in a single factor $\psi_i$.

Let $N_i$ be the number of variables in the factor $\psi_i$, and let $N_{\text{max}} = \max_i N_i$.

The running time of VE is then $O(mk^{N_{\text{max}}})$, where $k = |\text{Val}(X)|$. Why?

The primary concern is that $N_{\text{max}}$ can potentially be as large as $n$. 

---

**Running time of variable elimination**

<table>
<thead>
<tr>
<th>Step</th>
<th>Variable eliminated</th>
<th>Factors used</th>
<th>Variables involved</th>
<th>New factor</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>$C$</td>
<td>$\phi_C(C), \phi_D(D,C)$</td>
<td>$C, D$</td>
<td>$\tau_1(D)$</td>
</tr>
<tr>
<td>2</td>
<td>$D$</td>
<td>$\phi_G(G,I,D), \tau_1(D)$</td>
<td>$G, I, D$</td>
<td>$\tau_2(G,I)$</td>
</tr>
<tr>
<td>3</td>
<td>$I$</td>
<td>$\phi_I(I), \phi_S(S,I), \tau_2(G,I)$</td>
<td>$G, S, I$</td>
<td>$\tau_3(G,S)$</td>
</tr>
<tr>
<td>4</td>
<td>$H$</td>
<td>$\phi_H(H,G,J)$</td>
<td>$H, G, J$</td>
<td>$\tau_4(G,J)$</td>
</tr>
<tr>
<td>5</td>
<td>$G$</td>
<td>$\tau_4(G,J), \tau_3(G,S), \phi_L(L,G)$</td>
<td>$G, J, L, S$</td>
<td>$\tau_5(J,L,S)$</td>
</tr>
<tr>
<td>6</td>
<td>$S$</td>
<td>$\tau_5(J,L,S), \phi_I(J,L,S)$</td>
<td>$J, L, S$</td>
<td>$\tau_6(J,L)$</td>
</tr>
<tr>
<td>7</td>
<td>$L$</td>
<td>$\tau_6(J,L)$</td>
<td>$J, L$</td>
<td>$\tau_7(J)$</td>
</tr>
</tbody>
</table>
Let’s try to analyze the complexity in terms of the graph structure

$G_\Phi$ is the undirected graph with one node per variable, where there is an edge $(X_i, X_j)$ if these appear together in the scope of some factor $\phi$

Ignoring evidence, this is either the original MRF (for sum-product VE on MRFs) or the moralized Bayesian network:
Elimination as graph transformation

When a variable $X$ is eliminated,

- We create a single factor $\psi$ that contains $X$ and all of the variables $Y$ with which it appears in factors.
- We eliminate $X$ from $\psi$, replacing it with a new factor $\tau$ that contains all of the variables $Y$, but not $X$. Let’s call the new set of factors $\Phi_X$.

How does this modify the graph, going from $G_\Phi$ to $G_{\Phi_X}$?

- Constructing $\psi$ generates edges between all of the variables $Y \in Y$.
- Some of these edges were already in $G_\Phi$, some are new.
- The new edges are called **fill edges**.
- The step of removing $X$ from $\Phi$ to construct $\Phi_X$ removes $X$ and all its incident edges from the graph.
Example

(Graph)

(Elim. C)

(Elim. D)

(Elim. I)
We can summarize the computation cost using a single graph that is the union of all the graphs resulting from each step of the elimination.

We call this the **induced graph** \( \mathcal{I}_{\Phi, \prec} \), where \( \prec \) is the elimination ordering.
Example

(Induced graph)

(Induced graph)

(Maximal Cliques)
Properties of the induced graph

**Theorem:** Let $\mathcal{I}_{\Phi, \prec}$ be the induced graph for a set of factors $\Phi$ and ordering $\prec$, then

1. Every factor generated during VE has a scope that is a clique in $\mathcal{I}_{\Phi, \prec}$
2. Every maximal clique in $\mathcal{I}_{\Phi, \prec}$ is the scope of some intermediate factor in the computation

(see Koller & Friedman for proof)

Thus, $N_{\text{max}}$ is equal to the size of the largest clique in $\mathcal{I}_{\Phi, \prec}$

The running time, $O(mk^{N_{\text{max}}})$, is exponential in the size of the largest clique of the induced graph
Induced width

- The width of an induced graph is \#nodes in largest clique - 1
- We define the induced width \( w_{G,\prec} \) to be the width of the graph \( I_{G,\prec} \) induced by applying VE to \( G \) using ordering \( \prec \)
- The treewidth, or “minimal induced width” of graph \( G \) is
  \[ w_G^* = \min_{\prec} w_{G,\prec} \]
- The treewidth provides a bound on the best running time achievable by VE on a distribution that factorizes over \( G \): \( O(mk^{w_G^*+1}) \)
- Unfortunately, finding the best elimination ordering (equivalently, computing the treewidth) for a graph is NP-hard
- In practice, heuristics are used to find a good elimination ordering
Choosing an elimination ordering

Set of possible heuristics:

- **Min-fill**: the cost of a vertex is the number of edges that need to be added to the graph due to its elimination.

- **Weighted-Min-Fill**: the cost of a vertex is the sum of weights of the edges that need to be added to the graph due to its elimination. Weight of an edge is the product of weights of its constituent vertices.

- **Min-neighbors**: The cost of a vertex is the number of neighbors it has in the current graph.

- **Min-weight**: the cost of a vertex is the product of weights (domain cardinality) of its neighbors.

Which one better?

- None of these criteria is better than others.
- Often will try several.