Enumerating the junction trees of a decomposable graph

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Abstract

We derive methods for enumerating the distinct junction tree representations for any given decomposable graph. We discuss the relevance of the method to estimating conditional independence graphs of graphical models and give an algorithm that, given a junction tree, will generate uniformly at random a tree from the set of those that represent the same graph. Programs implementing these methods are included as supplemental material.

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1 Introduction

Decomposable or triangulated or chordal graphs are of interest in many areas of mathematics. Our primary interest is in their role as the conditional independence graphs of decomposable graphical models. In particular, we are interested in estimating decomposable graphical models from observed data using Markov chain Monte Carlo, or MCMC, schemes that traverse the space of decomposable graphs in order to sample from, or maximize, the posterior probability distribution defined by the data. The underlying approach is the same whether the data is continuous (Giudici & Green 1999, Jones, Carvalho, Dobra, Hans, Carter & West 2005) or discrete (Thomas & Camp 2004, Thomas 2005). A common feature of such schemes is that, given an incumbent decomposable graph G, we propose a new graph G' which is then accepted or rejected according to probabilities that depend on the distribution being sampled (Metropolis, Rosenbluth, Rosenbluth & Teller 1953, Hastings 1970, Kirkpatrick, Gellatt & Vecchi 1982). However, there are no known proposal schemes that guarantee in advance that G' will be decomposable, even if G is. Hence, it is necessary to test G' for decomposability before evaluating the usual acceptance probability. While such tests can be very quick (Giudici & Green 1999), for all practical methods for proposing a random G' of which we are aware, the probability that G' is decomposable decreases rapidly with the size of the graph, making this approach infeasible for large problems. For instance, in the genetic examples considered by Thomas (2009), involving up to 20,000 variables, the probability of proposing a decomposable G' decreases roughly as the inverse of the number of variables. Given these circumstances, it would be very useful to have an alternative representation of the problem that avoids the test for decomposability. It is with this in mind that we consider what follows.

It is often convenient in graphical modeling to operate not on the graph itself, but on its derived representation as a *junction tree*. This raises the prospect of discarding the underlying conditional independence graph entirely and defining MCMC schemes on the space of junction trees. As each junction tree uniquely defines a decomposable graph, this might avoid the expensive need to propose non-decomposable models. However, decomposable graphs have multiple equivalent junction tree representations and moreover the number is variable from graph to graph. Therefore, sampling the space of junction trees will over-sample decomposable graphs with a large number of such representations. This can be corrected for if the number of junction trees for any particular decomposable graph can be evaluated and this is the motivation for the method we present here.

We begin by reviewing some definitions and standard properties of decomposable graphs and junction trees. For more complete information on these see Golumbic (1980) and Lauritzen (1996), whose terminology we have adopted. We then consider the number of ways that sets of links of a junction tree that correspond to the same clique intersection can be rearranged. These counts are then combined to give the total number of junction trees. A simple algorithm is then presented that will take a junction tree and select an equivalent one uniformly at random from the set of all possible equivalents. Finally, we discuss the computational complexity of our method showing that it is faster than existing algorithms, and outline potential junction tree sampling methods.

2 Definitions and preliminary results

Consider a graph G = G(V, E) with vertices V and edges E. A subset of vertices $U \subseteq V$ defines an *induced subgraph* of G which contains all the vertices U and any edges in E that

connect vertices in U. A subgraph induced by $U \subseteq V$ is *complete* if all pairs of vertices in U are connected in G. A *clique* is a complete subgraph that is maximal, that is, it is not a subgraph of any other complete subgraph of G.

Definition 1 A graph G is decomposable if and only if the set of cliques of G can be ordered as (C_1, C_2, \ldots, C_c) so that for each $i = 1, 2, \ldots, c-1$

if
$$S_i = C_i \cap \bigcup_{j=i+1}^{c} C_j$$
 then $S_i \subset C_k$ for some $k > i$. (1)

This is called the running intersection property and is equivalent to the requirement that every cycle in G of length 4 or more is chorded. The sets $S_1, \ldots S_{c-1}$ are called the separators of the graph. The set of cliques $\{C_1, \ldots C_c\}$ and the collection of separators $\{S_1, \ldots S_{c-1}\}$ are uniquely determined from the structure of G; however, there may be many orderings that have the running intersection property. The cliques of G are distinct sets, but the separators are generally not all distinct.

Definition 2 The junction graph of a decomposable graph has nodes $\{C_1, \ldots, C_c\}$ and every pair of nodes is connected. Each link is associated with the intersection of the two cliques that it connects, and has a weight, possibly zero, equal to the cardinality of the intersection.

Note that for clarity we will reserve the terms vertices and edges for the elements of G, and call those of the junction graph and its subgraphs nodes and links.

Definition 3 Let J be any spanning tree of the junction graph. J has the junction property if for any two cliques C and D of G, every node on the unique path between C and D in J contains $C \cap D$. In this case J is said to be a junction tree.

Figure 1: A decomposable graph containing 23 vertices in 4 disjoint components.

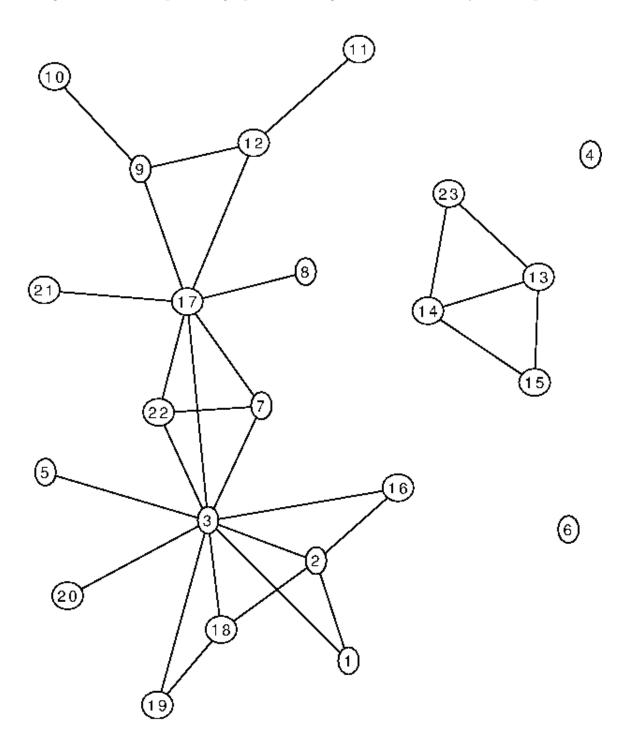


Figure 2: One possible junction tree for the graph shown in Figure 1. The 16 cliques are the vertices of the junction tree and are shown as ovals. The 15 clique separators are represented by the edges of the graph and each edge is associated with the intersection of its incident vertices. These intersections are shown as rectangles. Note that some of these intersections are empty.

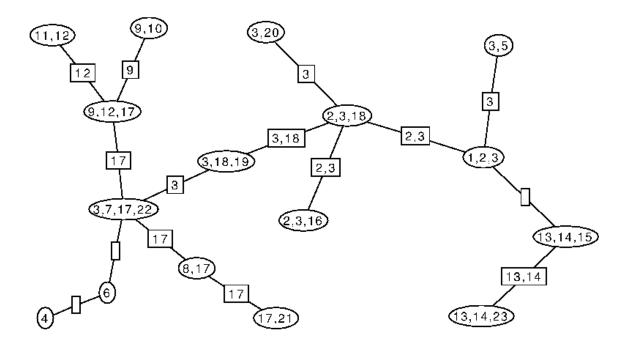


Figure 1 gives an example of a decomposable graph while Figure 2 shows one of its possible junction trees. The lexicographic search method of Tarjan & Yannakakis (1984) will find a junction tree for a given decomposable graph in time and storage of order |V| + |E|.

Note that some authors first partition a graph into its disjoint components before making a junction tree for each component, combining the result into a *junction forest*. The above definition, however, will allow us to state results more simply without having to make special provision for nodes in separate components. In effect, we have taken a conventional junction forest and connected it into a tree by adding links between the components. Each of these new links will be associated with the empty set and have zero weight. Clearly, this tree has the junction property. Results for junction forests can easily be recovered from the results we present below for junction trees.

As Lauritzen (1996) describes more fully, a junction tree for G will exist if and only if G is decomposable, and the collection of clique intersections associated with the c-1 links of any junction tree of G is equal to the collection of separators of G. Also, the junction property can be equivalently stated as the property that the subgraph of a junction tree induced by the set of cliques that contain any set $U \subseteq V$ is a single connected tree.

As stated in Definition 2, we can consider each link of the junction graph to have a weight. Thus, any subgraph of it, and in particular any spanning tree, can also be associated with a weight defined by the sum of the weights of the links included. Jensen (1988) exploits this to give the following useful characterization of a junction tree.

Theorem 4 A spanning tree of the junction graph is a junction tree if and only if it has maximal weight.

From this it is clear that any tree with the cliques of G as its nodes and for which the collection of clique intersections associated with the links is equal to the collection of separators of G, is a junction tree of G, since such a tree must span the junction graph and have maximal weight. Therefore, the problem of enumerating junction trees for a given graph G is equivalent to enumerating the ways that links of a given junction tree can be rearranged so that the result is also a tree, and the collection of clique intersections defined by the links of the tree is unchanged.

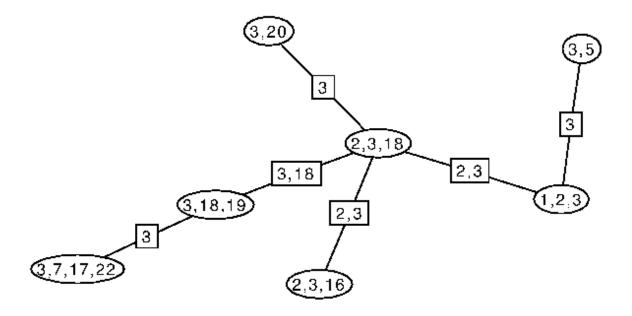
3 Rearranging the links for the set of separators with the same intersection

As noted above, the separators of G are not generally distinct. For example, in Figure 2 three links are associated with the clique intersection $\{17\}$ and two with the intersection $\{2,3\}$. We now consider the effect of rearranging all the links that are associated with the same clique intersection. Let J be any junction tree of G and S one of its separators. Define T_S to be the subtree of J induced by the cliques that contain S. The junction property ensures that T_S is a single connected subtree of J.

Clearly, any rearrangement of the links associated with S in J must be a rearrangement among certain links of T_S , since both cliques joined by such a link must contain S. For illustration, Figure 3 shows $T_{\{3\}}$, the subtree defined by the separator $\{3\}$ for the graph in Figure 1. If we now rearrange the links that are associated with S to produce a new graph, T'_S say, and replace T_S in J by T'_S to give a new graph J', J' will be a junction tree of G if and only if

• T'_S is a tree, and hence so is J', and

Figure 3: $T_{\{3\}}$, the connected subtree of the junction graph shown in Figure 2 induced by the cliques that contain the separator $\{3\}$.



• T'_S has the same weight as T_S , so that J' has the same weight as J.

In fact the second condition is redundant: all cliques in T_S contain S so their intersection must also do so, and any pair of cliques whose intersection is a superset of S cannot be joined in a tree T'_S unless already joined in T_S as T'_S would then have greater weight than T_S , and T'_S greater weight than T_S thus violating the latter's maximal weight property. So we need only count the number of ways of rearranging the links of T_S associated with T'_S is a tree.

Consider F_S to be the forest obtained by deleting all the links associated with S from T_S . For example, Figure 4 shows $F_{\{3\}}$, the forest obtained by deleting links associated with the separator $\{3\}$ from the tree $T_{\{3\}}$ shown in Figure 3. Define $\nu(S)$ to be the number of ways that the components of F_S can be connected into a single tree by adding the appropriate number of links. This value is given by a theorem by Moon (1970) which can be restated as follows.

Theorem 5 The number of distinct ways that a forest of p nodes comprising q subtrees of sizes $r_1 ldots r_q$ can be connected into a single tree by adding q-1 links is

$$p^{q-2} \prod_{i=1}^{q} r_i. \tag{2}$$

If the number of links associated with a given separator S is m_S we know that F_S will contain $m_S + 1$ components. Let these be of sizes $f_1, f_2, \ldots f_{m_S+1}$. Let the number of nodes in T_S be t_S which is simply the number of cliques of G that contain S. Then, directly from theorem 5 we obtain the following.

Theorem 6

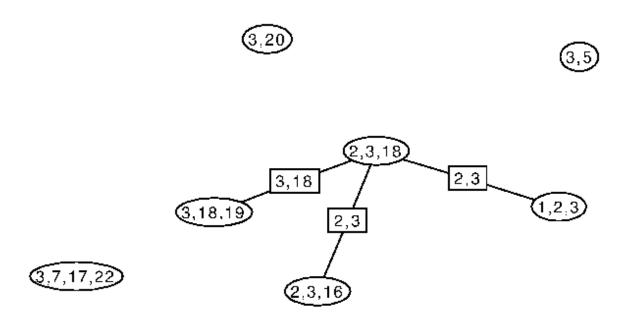
$$\nu(S) = t_S^{m_S - 1} \prod_{j=1}^{m_S + 1} f_j. \tag{3}$$

For example, the forest in Figure 4 has 7 nodes in 4 components of sizes 1, 1, 1 and 4. This forest, $F_{\{3\}}$, can be reconnected into a single tree by adding 3 links in $7^2 \times 1 \times 1 \times 1 \times 4 = 196$ different ways.

4 The number of junction trees for a decomposable graph

The final step in enumerating junction trees is to note that $\nu(S)$ depends only on the sizes of the components of F_S , not on their particular structure. These sizes are determined by the sets of cliques that contain separators that are supersets of S. Since the set of cliques and collection of separators are uniquely determined and independent of any particular

Figure 4: $F_{\{3\}}$, the forest obtained by from the tree shown in Figure 3 by deleting edges associated with the separator $\{3\}$.



junction tree, $\nu(S)$ is independent of J. Hence, the links associated with one particular separator can be reallocated independently of the links associated with another. Thus we obtain the following result.

Theorem 7 Consider a decomposable graph G with separators $S_1, \ldots S_{c-1}$. Let $S_{[1]}, \ldots S_{[s]}$ be the distinct sets contained in the collection of separators. The number of junction trees of G is

$$\mu(G) = \prod_{i=1}^{s} \nu(S_{[i]}). \tag{4}$$

As an example, the number of distinct junction trees for the graph shown in Figure 1 is 57,802,752. The calculations needed to obtain this are given in Table 1.

Table 1: The computations that enumerate the distinct junction trees for the decomposable graph given in Figure 1.

Separator S	m_S	t_S	$\{f_S\}$	$\nu(S)$
Ø	3	16	1, 1, 2, 12	6144
$\{13, 14\}$	1	2	1, 1	1
{3}	3	7	$1,\ 1,\ 1,\ 4$	196
$\{2, 3\}$	2	3	1, 1, 1	3
${3,18}$	1	2	1, 1	1
{9}	1	2	1, 1	1
{12}	1	2	1, 1	1
{17}	3	4	$1,\ 1,\ 1,\ 1$	16

$$\mu(G) = 6144 \times 1 \times 196 \times 3 \times 1 \times 1 \times 1 \times 16 = 57802752$$

As noted above, we can retrieve from this result the count of the number of possible conventional junction forests that a decomposable graph has. This is given simply by

$$\frac{\mu(G)}{\nu(\emptyset)}$$
,

which for the example is 57802752/6144 = 9408.

5 Randomizing the junction tree

Theorem 5 is similar in style to Prüfer's constructive proof (Prüfer 1918) of Cayley's result that there are n^{n-2} distinct labeled trees of n vertices (Cayley 1889). A similar construction lets us choose uniformly at random from the set of possible trees containing a given forest as follows:

1. Label each vertex of the forest $\{i, j\}$ where $1 \le i \le q$ and $1 \le j \le r_i$, so that the first index indicates the subtree the vertex belongs to and the second reflects some

ordering within the subtree. The orderings of the subtrees and of vertices within subtrees are arbitrary.

- 2. Construct a list v containing q-2 vertices each chosen at random with replacement from the set of all p vertices.
- 3. Construct a set w containing q vertices, one chosen at random from each subtree.
- 4. Find in w the vertex x with the largest first index that does not appear as a first index of any vertex in v. Since the length of v is 2 less than the size of w there must always be at least 2 such vertices.
- 5. Connect x to y, the vertex at the head of the list v.
- 6. Remove x from the set w, and delete y from the head of the list v.
- 7. Repeat from 4 until v is empty. At this point w contains 2 vertices. Connect them.

Given any particular junction tree representation J for a decomposable graph G we can choose uniformly at random from the set of equivalent junction trees by applying the above algorithm to each of the forests F_S defined by the distinct separators of J.

6 Computational complexity

Jensen's characterization of a junction tree as a maximal spanning tree of the junction graph means that general methods for enumerating the optimal spanning trees of a graph can also be applied to enumerating junction trees. The method of Broder & Mayr (1997) does precisely this. Recalling the notation used in section 2, the junction graph will have c nodes and c(c-1)/2 links. Broder and Mayr's method would require O(M(c)) elementary

operations to enumerate its maximal spanning trees, where M(c) is the number of operations needed to multiply $c \times c$ matrices. Asymptotically, the best algorithm for matrix multiplication is that of Coppersmith & Winograd (1990) which requires $O(c^{2.376})$ operations, although for realistically sized matrices the best practical methods, based on that of Strassen (1969), need $O(c^{2.807})$ operations. Letting n = |V|, the number of vertices in G, we note that c can be as large as n and typically grows linearly with n. Hence, Broder and Mayr's algorithm is at best an $O(n^{2.376})$ method.

However, as noted above, Jensen's characterization is not the only route to obtaining a junction tree. The lexicographic search of Tarjan & Yannakakis (1984) will find a simple elimination scheme, and hence a junction tree, in time O(n+m), where m=|E| the number of edges in G. While $m=O(n^2)$ in the worst case, typical graphical models are sparse and the time required is closer to linear in n. The enumeration method presented here depends only on knowing a single junction tree for G. The time required to carry it out is dominated by the time needed to find each $T_{S_{[4]}}$. We note that any link L of J will be a link in $T_{S_{[4]}}$ for each i such that $S_{[4]} \subseteq L$. Finding all the $T_{S_{[4]}}$ can be done by iterating over the c-1 links of J, and for each link checking for inclusion of each of the s distinct separators. Since both c and s can be O(n), the enumeration is an $O(n^2)$ algorithm in the worst case. Other ways of finding the $T_{S_{[4]}}$ will in practice be faster. For example, we can find $T_{S_{[4]}}$ by starting with a node that contains $S_{[4]}$ and searching outwards in J until we hit nodes that don't contain the separator. Thus, if $T_{S_{[4]}}$ is small it will be found very quickly. While it is straightforward to construct examples where this approach is also $O(n^2)$, for more typical graphs it will be considerably faster.

In summary, applying Broder and Mayr's general method to the special case of enumerating junction trees is at best an $O(n^{2.376})$ method, and more realistically $O(n^{2.807})$. By

exploiting the junction property, our method improves this to a worst case of $O(n^2)$ which in practice is a very conservative upper bound.

7 MCMC samplers for junction trees

Given a distribution $\pi(G)$ from which we want to sample decomposable graphs G, the methods of Metropolis et al. (1953) and Hastings (1970) allow us to construct Markov chains with $\pi(G)$ as the ergodic distribution. For example, we can propose a new graph G' by choosing two random vertices of G: if they are connected in G we disconnect them in G' and vice versa. G' is then accepted with probability $\max(1, \pi(G')/\pi(G))$, with the special case that $\pi(G')$ is defined to be 0 if G' is not a decomposable graph. Intuitively, it is easy to see that this can be very inefficient. If we consider choosing two random vertices of G, it is quite likely that we pick two vertices that are not connected directly, but which are connected by several paths through other vertices. Adding an edge between the vertices is, therefore, likely to create cycles. Unless all the connecting paths are short, a cycle of length 4 or more may well be formed which prevents G' being decomposable. Thomas (2009) shows that for modeling linkage disequilibrium between genetic loci, the acceptance rate decreases approximately as 1/n, making the method infeasible for large numbers of variables. As genetic methods now routinely assay hundreds of thousands of loci on a single chromosome, the high rejection rate becomes increasingly problematic.

The motivation for our enumeration method is that it makes it possible to devise MCMC schemes over decomposable graphs that that are expressed as operations on junction trees. If we wish to sample decomposable graphs from $\pi(G)$, it is sufficient to sample junction

trees from

$$\rho(J) = \frac{\pi(G(J))}{\mu(G(J))}$$

since

$$P(G) = \sum_{J:G(J)=G} \frac{\pi(G(J))}{\mu(G(J))}$$
$$= \frac{\pi(G)}{\mu(G)} \sum_{J:G(J)=G} 1$$
$$= \pi(G).$$

For each J sampled from a Markov chain with ergodic distribution $\rho(J)$, we would derive the graph G(J) which would be sampled with probability $\pi(G)$, as required. This, of course, requires efficient enumeration, but note that the Metropolis-Hastings acceptance probability for a junction tree MCMC scheme depends only on $\mu(G(J'))/\mu(G(J))$ which, given the factorization in equation 4 above, might be computable from $\mu(G(J))$ more simply than direct enumeration of $\mu(G(J'))$.

Simulating general labeled trees is a relatively straightforward matter. For instance, Prüfer's construction (Prüfer 1918) makes independent realizations of trees of a given size from a uniform distribution easy. However, for the junction tree problem the labels on the nodes of J are the cliques of G and these must be connected so that the junction property holds, making for a more difficult problem in a constrained space. Nonetheless, we have been able to construct an irreducible MCMC sampling scheme over the space of junction trees for graphs of a given size n. This involves operations on the nodes and links of an incumbent junction tree J that correspond to either adding edges to or deleting edges from G when the edges are chosen from highly restricted sets of possibilities. Following these

perturbation rules, ensures that any proposal J' is a junction tree for some graph G' on n vertices. Since J' is junction tree, G' must by definition be decomposable, and, hence, we avoid both the need to test for decomposability and the inefficiency of proposing non-decomposable graphs. The randomization method described in section 5 above can also be included in the scheme; although, it is not necessary for irreducibility, it may improve the mixing properties of the chain. While the space of junction trees is larger than that of decomposable graphs, it is more tractable and may be more easily traversible. A complete description of the method and implementation is the subject of a future manuscript currently under preparation.

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9 Supplemental material

The following materials are available to accompany this paper.

jtree.jar: This file contains the source code and compiled classes required to run the programs described below. The code can be run by adding jtree.jar to the list of files accessed by the user's classpath, or by using the -classpath option of the java command. (Jar, java archive, file.)

docs: This is a directory of HTML pages automatically produced for the code in jtree.jar using the javadoc program, and can be viewed using a web browser. (Directory of HTML files.)

illustration: This is simple text file specifying the graph used in this paper to illustrate junction tree enumeration. (Text file.)

The programs included in jtree.jar are:

CountJTrees: This reads a graph as a simple text file from the standard input stream and outputs the number of equivalent junction trees it has. If the input graph is not decomposable, zero is output. For example, to replicate the enumeration shown in Table 1, first ensure that files jtenum.jar and illustration are in the current working directory and then run the command:

java -classpath jtenum.jar CountJTrees < illustration

The following programs can be run in a similar way.

CountLogJTrees: This reads a graph as a simple text file and outputs the log of the number of equivalent junction trees it has. If the input graph is not decomposable, negative infinity is output.

FindRandomJTree: This reads a graph and outputs a junction tree selected uniformly at random from all possible junction tree representations. If the input graph is not decomposable, there is no output.

RandomJTreeDemo: This reads a graph and draws a junction tree representation to the screen. Every 2 seconds the junction tree is replaced by randomly chosen equivalent junction tree. If the input graph is not decomposable, the program exits with an error message.

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