# Two Algorithms for Generating Weighted Spanning Trees in Order

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#### 1. Introduction

In many practical situations, one wants to generate the spanning trees of a graph in order of increasing weight. For example, consider this electrical wiring problem: n pins must be wired together with as little wire as possible; further, the wiring must satisfy complicated and diverse constraints, e.g., at most k wires can meet any pin, no two wires can be too close, etc. One way to solve the problem is to generate spanning trees in order of increasing length (weight) until a tree satisfying the constraints is found.

This paper presents an algorithm for generating the K smallest spanning trees of a connected graph, in order of increasing weight.  $\underline{K}$  may be known in advance or specified as trees are generated. The algorithm requires time  $O(KE\alpha(E,V)+E\log E)$  and space O(K+E). Here V is the number of vertices, E is the number of edges, and  $\alpha$  is Tarjan's inverse of Ackermann's function and is very slow-growing. A previously known algorithm for this problem [2] has slower run time,  $O(K \min(N^2, E(\log E)^2) + K\log K)$ . The algorithm works by using a minimum weight spanning tree as a "reference tree" [6], and exchanging edges to find other trees. It uses fast set merging algorithms [10], and a bounding technique for partitioning the solution space [5].

The algorithm is modified for the problem of generating all spanning trees of a connected graph, in order of increasing weight. The run time is O(NE) and the space is O(N+E), where N is the number of spanning trees of the graph. Efficient algorithms for generating all spanning trees of a graph without weights [7,8] can be applied to this problem.

The best known algorithm gives time bound O(NE) and space bound O(NV+E). Since N can be large, storage may be a limiting factor. So the improvement in space is significant.

Section 2 gives definitions from graph theory. Section 3 gives a result on edge exchanges that is the basis of the algorithm. Section 4 presents the algorithm, and analyzes time and space requirements. Section 5 describes the modified algorithm. Section 6 summarizes computational experience.

#### 2. <u>Definitions</u>

A graph G consists of a finite set of V vertices and a finite set of E edges. An edge is an (unordered) set of two distinct vertices. The edge containing vertices v and w is denoted (v,w); it joins v and w.

A <u>subgraph</u> of G is a graph whose vertices and edges are in G. If H is a collection of edges on the vertices of G, graph G-H consists of the vertices of G and all edges in G except H; graph GUH consists of all vertices of G and all edges in G or H.

A path from  $v_1$  to  $v_n$  is a sequence of edges  $(v_1, v_2), (v_2, v_3), \ldots$ ,  $(v_{n-1}, v_n)$ . If all vertices  $v_i$ ,  $1 \le i \le n$ , are distinct, the path is <u>simple</u>. A graph is <u>connected</u> if there is a path between any two distinct vertices. A bridge is an edge (v, w) that is in every path from v to w.

A <u>tree</u> is a connected graph, where any two distinct vertices are joined by a unique simple path. A <u>spanning tree</u> of G is a tree that is a subgraph containing all vertices of G.

A <u>rooted tree</u> is a tree T with one vertex r chosen as the <u>root</u>. Let v be a vertex in T, and let  $v = v_0, v_1, \ldots, v_n = r$  be the sequence of vertices in the simple path from v to r. Any vertex  $v_i$ , for  $0 \le i \le n$ , is an <u>ancestor</u> of v;  $v_i$  is the <u>father</u> of v; v is a <u>son</u> of  $v_i$ ; the <u>depth</u> of

v is n. The <u>first common ancestor</u> of two vertices v and w is the ancestor of both v and w that has greatest depth. Two distinct sons of the same vertex are <u>brothers</u>. In an <u>oriented tree</u>, the sons of a vertex are ordered from left to right.

In a <u>weighted graph</u>, every edge e has a number,  $\underline{w(e)}$ , called its  $\underline{weight}$ . If H is a set of edges (such as a spanning tree), the <u>weight of H</u> is  $\sum w(e)$ .  $e \in H$ 

#### 3. T-exchanges

This section describes how spanning trees can be derived by exchanging edges. T-exchanges are defined and a useful property is proved.

Let T be a spanning tree of graph G. A <u>T-exchange</u> is a pair of edges e,f where e $\epsilon$ T, f $\epsilon$ T, and T-e $\theta$ f is a spanning tree. The <u>weight of exchange</u> e,f is w(f)-w(e). So the weight of tree T-e $\theta$ f is the weight of exchange e,f.

A T-exchange can be used to derive one minimum weight spanning tree from another, as shown below.

<u>Lemma 1</u>: A spanning tree T has minimum weight if and only if no T-exchange has negative weight.

<u>Proof</u>: The necessity of this condition is obvious. Sufficiency is proved in [4].

Theorem 2: Let T be a minimum weight spanning tree of G, and let e be an edge in T. Let e,f be a T-exchange having the smallest weight of all T-exchanges e,f'. Then T-eVf is a minimum weight spanning tree of graph G-e.

<u>Proof</u>: Let S = T-eUf. Suppose S does not have minimum weight. By Lemma 1, there is an S-exchange g,h having negative weight. We derive a contradiction below.

We first show edges e,f,g,h are situated as in Figure 1. Let T-e consist of the two trees U,V. Edge e joins U and V. Edge f also joins U and V, since e,f is a T-exchange.

Edge h also joins U and V. For if not, assume without loss of generality that h joins two vertices in U. Since g,h is an S-exchange,

g is in tree U. Thus g,h is a T-exchange. But g,h has negative weight, and T is a minimum weight spanning tree. This contradiction proves h joins U and V.

Now we show edge  $g \in U \cup V$ . Since e,h is a T-exchange, its weight is no smaller than that of e,f. Thus

(1) 
$$w(f) \leq w(h) < w(g).$$

So  $g \neq f$ , and  $g \in S - f = U \cup V$ .

Assume without loss of generality that geU. This gives Figure 1.

Now let U-g consist of the two trees W,X. Edge e is incident to one of these trees, say W. Since T-e-gUfUh is a spanning tree, either f or h is incident to X. So either g,f or g,h is a T-exchange. But (1) implies both of these exchanges have negative weight, a contradiction. This contradiction shows the original assumption is false. Thus S has minimum weight.

Now let T be a minimum weight spanning tree. The Theorem shows a spanning tree with the next smallest weight is T-eUf; where e,f is a minimum weight T-exchange. This observation is the basis of the algorithm.

#### 4. Algorithm for K Spanning Trees

The algorithm for generating K spanning trees consists of procedures EX,GEN, and a main procedure GENK. This section describes these procedures in turn.

EX finds a minimum weight T-exchange e,f subject to certain constraints, IN,OUT. We describe EX, first assuming there are no constraints and then incorporating the constraints.

For each edge geT, EX finds a minimum weight T-exchange g,h; it sets e,f to the smallest of these exchanges. During the execution of EX, we call an edge geT eligible if a minimum weight exchange g,h has not been found.

EX works as follows. Originally all edges in T are eligible. A list L contains all edges  $h_1, h_2, \ldots, h_E$  of G, sorted in order of increasing weight. EX begins by finding every T-exchange  $g, h_1$ . This is a minimum exchange for g, since  $h_1$  has the smallest weight possible. Now all edges g become ineligible.

Edge  $h_2$  is processed similarly. EX finds every T-exchange  $g,h_2$ , where g is eligible. This is a minimum exchange for g. The edges g become ineligible.

The procedure is repeated for every edge  $h_{k}$  in L. The smallest exchange found is the minimum T-exchange e,f.

The tree T is represented by a father array F. Vertex 1 is the root; for any vertex  $v \ne 1$ , F(v) is the father of v. Call a vertex  $v \ne 1$  eligible if edge (v,F(v)) is eligible, i.e., a minimum exchange for (v,F(v)) has not been found; call vertex 1 eligible too.

Now suppose EX examines the edge  $h_k = (x,y)$ , as shown in Figure 2. Vertex  $\underline{a}$  is the first common ancestor of x and y that is eligible; vertices  $x_i$ , for  $1 \le i \le n$ , are the other eligible vertices on the path from x to  $\underline{a}$ ; vertices  $y_j$ , for  $1 \le j \le m$ , are defined similarly. Edge  $h_k$  gives a minimum exchange for each edge  $(x_i, F(x_i))$ ,  $(y_j, F(y_j))$ , and no others.

EX finds the vertices  $x_i, y_j$ , by using set merging techniques. A family of sets partitions the vertices of G. Each set contains a unique eligible vertex, which is used as the name of the set. Vertex w is in the set named v when v is the first eligible ancestor of w. (Since 1 is eligible, v exists for any w.) EX uses two procedures to manipulate these sets: FIND (v) computes the name of the set containing vertex v; UNION (v,w,x) combines sets named v and w into a new set named x (destroying sets v and w).

EX computes the vertices  $x_i$  using the equations  $x_1 = FIND(x)$ ,  $x_{i+1} = FIND(x_i)$ . Vertices  $y_j$  are computed similarly. Vertex  $\underline{a}$  is the first common value,  $\underline{a} = x_{n+1} = y_{m+1}$ . In this manner, EX finds all minimum exchanges for edge  $h_{\nu}$ .

Now we discuss the constraint lists IN and OUT. These lists prevent trees from being generated twice; their construction is described fully below IN is a list of edges in T that must remain in IT; OUT is a list of edges out of T that must remain out of T. Thus EX must find a T-exchange e,f, such that esT-IN, fsG-OUT, and e,f has the smallest weight possible.

To do this, EX begins by making all IN edges (x,y) ineligible. This is done by placing x and y in the same set (since they have the same first eligible ancestor). Also, EX marks all OUT edges in the list L, so they are not considered for exchanges. These edges are later unmarked, for subsequent calls to EX. In this manner, the constraints are handled.

The final array used is W, which specifies the weight W(e) of each edge e.

The procedure below sets global variables e,f and r, so e,f is a minimum weight exchange subject to constraints IN,OUT, and r is the weight of e,f.

### procedure EX(F,IN,OUT); begin

- 1. <u>initialize</u>:  $r \leftarrow \infty$ ; make each vertex v the only element in a set named v;
- 2. IN edges: for edge (x,y) in IN do begin wlog assume F(x) = y;
- 3.  $y \leftarrow FIND(y)$ ; UNION(x,y,y);

end;

```
4.
     OUT edges:
                         for edge (x,y) in OUT do mark (x,y) in L;
                         for edge (x,y) in L do comment edges in L are in
 5.
     L edges:
                         increasing weight order;
 6.
                           if(x,y) is marked then unmark (x,y)
 7.
                           else if F(x) \neq y and F(y) \neq x then begin
 8.
     find exchanges:
                        let a be the first eligible common ancestor of x and y;
 9.
                         for v1 \leftarrow x,y do begin v \leftarrow FIND(v1);
10.
                           while v ≠ a do begin
                                  rl \leftarrow W(x,y)-W(v,F(v)); comment rl is the
11.
     check exchange:
                                  exchange's weight;
                                  if rl < r then begin r \leftarrow rl; e \leftarrow (v,F(v));
12.
                                  f \leftarrow (x,y), end;
                                  u \leftarrow FIND(F(v)); UNION(v,u,u); v \leftarrow u;
13.
     advance:
                         end end end ex;
```

Table I illustrates EX. Input parameters describe  ${\rm T_1}$  for the graph of Figure 3. The exchanges found by EX in step 8 are listed, along with the output values describing  ${\rm T_2}$ .

<u>Lemma 3</u>: Let T be a minimum weight spanning tree subject to constraints IN CTCG-OUT. Then EX, called with a father array for T, sets e,f so T-evf is the next smallest spanning tree subject to the same constraints.

<u>Proof:</u> Theorem 2 shows T-e**U**f has the desired property if e,f is a minimum weight T-exchange subject to constraints e£IN, f£OUT. The remarks preceding EX show EX finds such an exchange.

Lemma 4: EX runs in time  $O(E_{\infty}(E,V))$ .

<u>Proof:</u> First we bound the time for set operations. Lines 3 and 13 together do at most one UNION for each vertex. So they do O(V) UNIONs, and also O(V) FINDs. Line 9 does at most two FINDs for each edge in G-T. Thus, a total of O(V) UNIONs and O(E) FINDs are done. If fast set merging algorithms are used, the total time is  $O(E\alpha(E,V))$  [10].

Line 4 can be done in time O(E). For OUT contains at most E edges e; e can be marked (in L) in time O(1), if OUT contains a pointer to e in L.

Next consider line 8. Vertex <u>a</u> can actually be computed as exchanges for vertices v are found (lines 9-13). To do this, the paths from x to <u>a</u> and y to <u>a</u> are processed simultaneously. Referring to Figure 2, suppose  $x_i$  and  $y_j$  have been found, for some isn, jsm; if  $x_i \neq y_j$ , the vertex with greater depth precedes <u>a</u>, and so dis  $x_{i+1}$  or  $y_{j+1}$ . In this manner, the exchanges for vertices v, and the vertex <u>a</u>, are found.

To do this efficiently, we use an array D, that gives the depth D(v) of vertex v in T. D can be computed from F in time O(V). Thus, line 8 requires only O(V) extra time.

The remaining lines in EX require time O(E+V). This gives the desired time bound.

Procedures GEN and GENK call EX to generate spanning trees in correct order. A technique resembling branch and bound, described by Lawler [5], is used.

Let  $T_i$ , for  $1 \le i \le N$ , denote the spanning trees of G, in order of increasing weight. Suppose the algorithm has output the first j-1 trees  $\{i,j\}$ . The remaining trees have been partitioned into j-1 sets of the form trees one been partitioned.

$$P_i^{j-1} = \{T_k \mid k > j-1; e_1, e_2, ..., e_r \in T_k; e_{r+1}, ..., e_s \notin T_k\},$$

for  $1 \le i < j$  (Here r and s vary over sets). Tree  $T_i$  satisfies all conditions of this set except the first, k > j-1. The smallest tree in this set is known; it is the result of a  $T_i$ -exchange e,f.

Procedure GEN finds the next smallest tree  $T_j$  from among the j-1 smallest trees in these sets. Suppose  $T_j$  is in the above set  $P_i^{j-1}$ , and  $T_j = T_i - e U f$ . GEN computes  $T_j$  and outputs it. Then a new partition  $P_k^j$ ,  $1 \le k \le j$ , is formed, by subdividing  $P_i^{j-1}$ . For all  $k \ne i$  in  $1 \le k < j$ , set  $P_k^j = P_k^{j-1}$ ; further,

$$P_{i}^{j} = \{T_{k} \mid k>j; e_{1}, ..., e_{r}, e \in T_{k}; e_{r+1}, ..., e_{s} \notin T_{k}\},$$

$$P_{i}^{j} = \{T_{k} \mid k>j; e_{1}, ..., e_{r} \in T_{k}; e_{r+1}, ..., e_{s}, e \notin T_{k}\}.$$

 $T_i$  satisfies all conditions of  $P_i^j$  except the first. GEN finds the smallest tree in  $P_i^j$  by executing EX on  $T_i$ , with edges  $e_1, \ldots, e_r$ ,  $e_r$ , in IN, and  $e_{r+1}, \ldots, e_s$  in OUT. (Lemma 3 shows EX works correctly.)  $P_j^j$  is processed similarly, using tree  $T_j$ . In this way, the partition is updated. Now  $T_{i+1}$  can be found.

The algorithm uses a list P to represent the partition. A set  $P_i^j$  is represented by a tuple (t,e,f,F,IN,OUT) in P. Here t is the weight of the smallest tree in  $P_i^j$ ; F is the father array for  $T_i$ ; the smallest tree in  $P_i^j$  results from the  $T_i$ -exchange e,f; IN is the list of edges that are in all trees of  $P_i^j$ ; OUT is the list of edges that are out of all trees of  $P_i^j$ .

Procedure GEN outputs the next smallest tree  $T_j$  from set  $P_i^{j-1}$ , and puts the new sets  $P_i^j, P_j^j$  in the partition.

## procedure GEN; begin

- 1.  $\frac{\text{find P}_{1}^{j-1}}{\text{i}}$ : remove the tuple (t,e,f,F,IN,OUT) with smallest weight t from P;
- 2. if  $t = \infty$  then stop; comment all spanning trees have been output;

- 3. output  $T_i$ : Fj  $\leftarrow$  F; modify Fj so edge f replaces e; output (Fj);
- 4.  $ti \leftarrow t-W(f)+W(e)$ ; comment ti is the weight of  $T_i$ ;
- form list INi by adding e to IN; form list OUT; by adding e to OUT;
- 6.  $\underline{\text{form P}}_{i}^{j}$ : EX(F,INi,OUT); add (ti+r,e,f,F,INi,OUT) to P;  $\underline{\text{comment}}$  EX sets e,f,r for the minimum weight exchange;
- 7.  $\underline{\text{form P}}_{j}^{j}$ : EX(Fj,IN,OUTj); add (t+r,e,f,Fj,IN,OUTj) to P; end GEN;

Table II illustrates GEN, by showing the partition after tree  $T_2$  for Figure 3 is output.

Lemma 5: Let GEN be called, with P containing tuples for the sets  $P_k^{j-1}$ ,  $1 \le k \le j$ . Then GEN outputs  $T_j$ , and changes P to tuples for the sets  $P_k^j$ ,  $1 \le k \le j$ .

Proof: The Lemma follows from the remarks preceding GEN.

<u>Lemma 6</u>: GEN runs in time  $O(E\alpha(E,V))$ .

<u>Proof</u>: Lines 1, 6, and 7 remove and add tuples to P. These operations can be done in time O(E). For this, we use a heap [1]. The tuples in P are numbered, by assigning i to  $P_i^j$ . An entry in the heap is a number i; entries are ordered on the weight t of the corresponding tuple. The heap contains at most K entries. So an entry can be removed or added in time  $O(\log K)$ . Since  $K<2^E$ , this is O(E).

Line 3 requires time O(V) to derive Fj. Referring to Figure 2, let  $e = (x_i, F(x_i))$ , f = (x,y). To derive Fj, array F need only be changed on the path from x to  $x_i$ .

The rest of the time is dominated by the two calls to EX. This gives the desired bound.

The main procedure GENK finds a minimum spanning tree, and calls GEN for the next (K-1) trees.

#### procedure GENK(K); begin

- 1. make L a list of the edges of G, sorted in order of increasing weight;
- 2.  $\underline{\text{find } T_1}$ : make F a father array for a minimum weight spanning tree, having weight t;  $\underline{\text{output}}(F)$ ;
- 3. form  $P_1^1$ : EX  $(F, \phi, \phi)$ ; make  $(t+r, e, f, F, \phi, \phi)$  the only tuple in P;
- 4. generate  $T_j$ : for  $j \leftarrow 2$  to K do GEN; end GENK;

Table III shows the partition after all minimum weight spanning trees for Figure 3 are output.

<u>Lemma 7</u>: GENK generates the K smallest weight spanning trees in order, in time  $O(KE\alpha(E,V) + E \log E)$ .

<u>Proof:</u> The correctness of GENK follows by induction, using Lemma 5. Now we analyze the time. Line 1 requires time  $O(E \log E)$  to sort E edges. Line 2 can be done in time  $O(E \log E)$ , using Kruskal's algorithm [1]. The rest of the time is dominated by O(K) calls to GEN, requiring time  $O(KE\alpha(E,V))$  by Lemma 6.

When K is small, the term E log E in the run time is significant. It can be reduced to E log log E. This is done by using a faster minimum spanning tree algorithm [3,12], and changing EX so the edges of G need not be sorted. For the latter, the techniques of [11] are used.

Now we examine the space requirements. The dominating requirement is for P. Since a tuple can contain O(E) words, P can use O(KE) words. Below we modify the data structure to reduce this bound, without changing the run time.

Lemma 8: GENK uses O(K+E) space.

<u>Proof</u>: We show the structures F,IN,OUT in a tuple can be replaced by 6 words, and still be computed in time O(E). The extra time is spent in line 1 of GEN, and does not change the algorithm's time bound; P is reduced to O(K) words. So this suffices to prove the Lemma.

Suppose the first k trees  $T_j$ ,  $1 \le j \le k$ , have been output. Make these trees the vertices of an oriented tree  $\boldsymbol{\mathcal{T}}$ , as follows. Make  $T_1$  the root of  $\boldsymbol{\mathcal{T}}$ . For j > 1, make  $T_j$  a son of  $T_i$  if GEN derives  $T_j$  from a  $T_i$ -exchange. Let  $e_j$ ,  $f_j$  denote this  $T_i$ -exchange. Arrange the sons  $T_j$  of  $T_i$  so j increases from left to right.

For a set  $P_j^k$ , the structures F, IN, and OUT can be derived from  ${\cal T}$ , as follows. Let A be the path in  ${\cal T}$  from  $T_j$  to the root  $T_1$ . Consider these sets:

$$\begin{array}{l} 0_1 = \{e_i \mid T_i \text{ is in A and i>1}\},\\ 0_2 = \{f_i \mid T_i \text{ is in A and i>1}\},\\ T = (T_1 U 0_2) - 0_1,\\ I_1 = \{e_i \mid T_i \text{ is a left brother of some } T_\ell \text{ in A}\},\\ I_2 = \{e_i \mid T_i \text{ is a son of } T_j\}. \end{array}$$

T contains the edges of  $T_j$ ; the father array F for  $T_j$  is easily derived.  $I_1 U I_2$  contains the edges of IN, and  $O_1$  contains the edges of OUT.

Now for each set  $P_j^k$ , replace F,IN,OUT by the 6 words j,i,ej,fj,s,b. Here j is the index in  $P_j^k$ ; i is the index of the father  $T_i$  of  $T_j$ ; ej,fj is the exchange that derives  $T_j$  from  $T_i$ ; s is the index of the rightmost son (if any) of  $T_j$ ; b is the index of the brother (if any) immediately to the left of  $T_j$ .

These 6 words are easily computed in GEN. Let the tuple found in line 1 be (t,e,f,i,il,ei,fi,s,b). Then line 6 adds the tuple (ti+r,e,f,i,il,ei,fi,j,b) to P; line 4 saves e,f as ej,fj, and line 8 adds the tuple (t+r,e,f,j,i,ej,fj,0,s) to P.

Finally, it is easy to see the five sets above, and F,IN,OUT, can be constructed (in line 1 of GEN) in time O(E).

Table IV shows how the partition in Table III is changed using this scheme.

#### 5. Algorithm for All Spanning Trees

This section describes a modification of the algorithm that generates all spanning trees of a connected graph, in order of increasing weight. The run time is O(NE), a slight improvement over GENK. The three main changes are described below.

The first change eliminates extra calls to EX, by modifying the tuples for sets  $P_i^j$  in P. The tuple's minimum  $T_i$ -exchange e,f is replaced by X, a list of  $T_i$ -exchanges. For each edge  $g\epsilon T_i$ -IN, X contains g,h, the smallest exchange with  $h\epsilon G$ -OUT. So e,f is the smallest exchange in X.

EX is modified to generate X. X is initialized to an empty list (in line 1), and a line is added after 12:  $\frac{1}{2}$  add  $\frac{1}{2}$  add  $\frac{1}{2}$  add exchange (v,F(v)),(x,y) to X;

GEN is modified to use X. In line 1, the minimum exchange e,f is found by examining X. In line 6, the call to EX is eliminated; the tuple

for  $P_i^j$  is formed by removing e,f from X, and computing the new minimum weight t, using X.

The second change, in the set merging algorithms, speeds up FIND at the expense of UNION. Sets are represented by an array VSET, where VSET(w) = v when v is the name of the set containing w. With this approach, FIND takes time O(1) and UNION takes time O(V).

EX uses these algorithms to manipulate the sets of vertices. Also, lines 2-3 are modified to avoid calls to UNION:

- 2'. IN edges: sort the edges (x,y) of IN, so F(x) = y and the depth of x increases;
- 3'. for edge (x,y) in IN do  $VSET(x) \leftarrow VSET(y)$ ;

When (x,y) is processed in line 3', VSET(y) already has its final value. So lines  $2^{\frac{1}{2}}3^{\frac{1}{2}}$  initialize the sets correctly.

The third change involves the main procedure. A new main procedure is used to insure all trees are generated:

# procedure GENA; begin GENK(∞) end;

Note the algorithm halts in line 2 of GEN.

Further, GENK is modified to reduce the set-up time in lines 1-2. Line 1 is changed:

1'. Make B a list of the bridges of G; make L a list of the edges of G-B, sorted in order of increasing weight;

In line 2, the bridges B are placed in the spanning tree, before the minimum spanning tree algorithm is called.

The remaining changes in the algorithm are typographical.

<u>Lemma 9</u>: GENA generates all spanning trees in order, in time O(NE) and space O(N + E), where N is the number of spanning trees.

<u>Proof</u>: The correctness of GENA follows from Lemma 7 and the discussion above. Now we analyze the time, beginning with GENK.

Let E' be the number of edges that are not bridges. Then lines 1'-2 of GENK use time  $O(E + E' \log E')$ . For in line 1', the bridges B can be found in time O(E) [9]; L can be sorted in time  $O(E' \log E')$ . Line 2, using Kruskal's algorithm with the bridges already in the tree, uses time  $O(E' \log E')$ .

Now we show E' log E' is O(NE). It suffices to show E' < 2N. Let T be a spanning tree. We can list T-exchanges e,f, so each edge that is not a bridge occurs in the list. Since each exchange represents a distinct tree, E' < 2N. Thus, lines 1'-2 use time O(NE).

The rest of GENK is dominated by O(N) calls to GEN. As before, the heap operations in GEN (lines 1,6,7) require time O(NE); the rest of GEN is dominated by O(N) calls to EX (line 7).

In EX, line 2' can be done once in time O(V). First the depth of each vertex in the tree is computed, using F. Then the edges of IN are sorted on depth, using a bucket sort [1] with one bucket for each depth. Thus, line 2' uses total time O(NV).

Line 13 of EX, using the modified FIND and UNION procedures, is done once in time O(V). It is executed N times, once for each spanning tree. So the total time is O(NV).

The rest of EX uses time 0 (NE). This gives the desired time bound for GENA.

Now we examine the space. The X lists in P require a total of O(N) words, since each exchange in an X list represents a distinct spanning tree. Using the modification of Lemma 8, the rest of P uses O(N) words. So O(N + E) space is used.

Minty [7] and Read and Tarjan [8] give an algorithm that generates all spanning trees of a graph without weights. It requires time O(NE) and space O(E), assuming each tree is output as soon as it is generated. We can generate all spanning trees in a weighted graph, by first applying this algorithm, and then sorting the trees in order of increasing weight. This method requires time O(NE). The space is O(NV + E), since each tree must be stored until the sort is done. GENA has the same time bound but uses less space. Since N can be large  $(N = V^{V-2})$  in a complete graph, space is likely to be a limiting factor, so this improvement is significant.

#### 6. <u>Computational Experience</u>

A version of GENK has been programmed in PASCAL and run on the CDC 6400. The program stores tuples in a hybrid form (using the notation defined above, a tuple is (t,X,i,fj,F,IN)). The time bound for the program is  $O(KE\alpha(E,V) + E \log E)$ , and the space is O(KV + E).

Experiments were conducted on random connected graphs with random integer edge weights between 50 and 10000. Tables V-VI show results. The run time is specified by  $\overline{T}$ , the average time in milliseconds to generate one spanning tree. (Experiments show for a given graph, the time per spanning tree is approximately constant, as expected.) Table V shows  $\overline{T}$  for graphs that are almost complete;  $\overline{T}$  is computed from K = 15. Table VI shows  $\overline{T}$  for graphs with 60 vertices; again K = 15. When  $\overline{T}$  is plotted against E, the data is almost linear. Least square fits

for the two tables have slopes .72 and .68, respectively. Almost linear performance is expected from the asymptotic time bound, since the factor  $\alpha(E,V)$  is constant in this relatively small range.

#### 7. Acknowledgments

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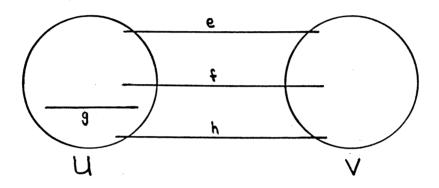


Fig. 1 Edges in Theorem 1

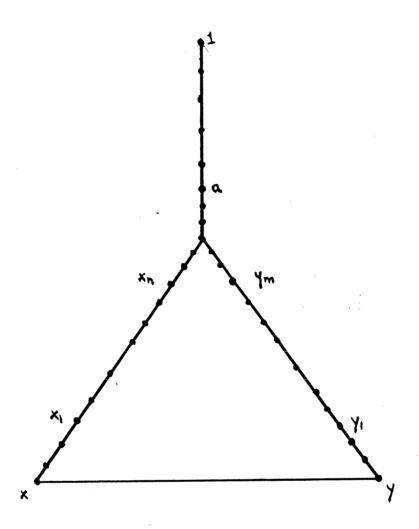


Fig. 2 Exchanging (x,y) into T

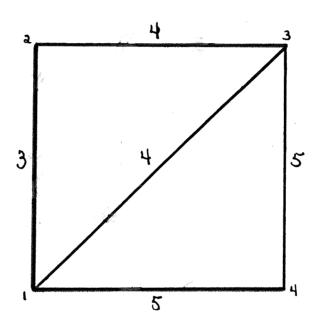


Fig. 3 Example graph

	t	F(2)	F(3)	F(4)	ΙN	OUT		ŕ	е	f
Ь	2	1	2	3	-	-	(2,1),(3,1);(3,2),(3,1);(4,3),(4,1)	0	(3,2)	(3,1)
			inp				exchanges	<b></b>	output	

Table I.

# EX processes $T_1$

<b>*</b>	t_	е	f	F(2)	F(3)	F(4)	IN	€ OUT
<b></b>		(4,3)				l		-
P <sub>2</sub> <sup>2</sup>	12	(4,3)	(4,1)	].	1	3	-	(3,2)

Table II.

Partition  $P_i^2$ 

	t	е	f	F(2)	F(3)	F(4)	IN	OUT
P1	13	(2,1)	(3,1)	1	2	3	(3,2),(4,3)	-
P <sub>2</sub>	13	(3,1)	(4,1)	1	1	3	(4,3)	(3,2)
P34	13	(2,1)	(3,1)	1	2	1	(3,2)	(4,3)
P4	∞	-	-	1	1	1	-	(3,2),(4,3)

Table III.

Partition P<sub>i</sub>

		j	i	еj	fj	S	- b
	P4 1	: 1	-	_	. <b>-</b>	3	<u>-</u>
Property and a second s	P <sub>2</sub> 4	2	7	(3,2)	(3,1)	4	
bearing and the state of the st	P <sup>4</sup> 3	3	1	(4,3)	(4,1)	-	2
Manhatan Manhatan Manhatan	P44	4	2	(4,3)	(4,1)		

Table IV

Changes in  $P_i^4$  using storage reduction scheme

٧	10	15	25	35	45	55	60
E	44	104	300	595	985	1476	1762
T	12	27	74	143	236	362	422

Table V

Time for generating one tree

E	159	336	513	690	867	1044	1221	1398	1575	1757
Ŧ	48	86	124	162	199	237	279	324	365	413

Table VI

Time for generating one tree, V = 60

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