# OPTIMAL PARAMETRIC SEARCH ALGORITHMS IN TREES I: TREE PARTITIONING

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# Optimal Parametric Search Algorithms in Trees I: Tree Partitioning

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Abstract. Linear-time algorithms are presented for partitioning a tree with weights on the vertices by removing k edges so as to either minimize the maximum weight component or maximize the minimum-weight component.

Key words and phrases. Data structures, max-min, min-max, p-center, parametric search, partial order, sorted matrices, tree partitioning.

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## 1. Introduction

Parametric search is a powerful technique for solving various optimization problems [AP], [C1], [C2], [CSSS], [FJ1], [FJ2], [Gu], [M], [R], [S], [Z]. To solve a given problem by parametric search, one must find a largest (or smallest) value that passes a certain feasibility test. In general, a set of candidate values is explicitly or implicitly identified, and the set is searched, choosing one value at a time upon which to test feasibility. As a result of the test, various values in the set will be discarded from further consideration. Eventually the search will converge on the optimal value.

In [M], Megiddo has emphasized the role that parallel algorithms can play in the implementation of such a technique. In [C2], Cole improves the time bounds for a number of problems addressed in [M], abstracting two techniques upon which parametric search can be based: sorting and performing independent binary searches. Despite the clever ideas elaborated in these papers, none of the algorithms presented in [M] and [C2] is known to be optimal. In fact, rarely have algorithms for parametric search problems been shown to be optimal. (For one that is optimal, see [CSSS].) In at least some cases the time required for feasibility testing matches known lower bounds for the parametric search problem. But in worst case  $\Omega(\log n)$  values must be tested for feasibility. Is this extra factor of at least  $\log n$  necessary? For several parametric search problems on trees, we show that a polylogarithmic penalty in the running time can be avoided, and give linear-time (and hence optimal) algorithms for these problems.

We first consider the max-min tree k-partitioning problem [PS]. Let T be a tree with n vertices and a nonnegative weight associated with each vertex. Let k < n be a positive integer. The problem is to delete k edges in the tree so as to maximize the weight of the lightest of the resulting subtrees. Perl and Schach introduced an algorithm that runs in  $O(k^2rd(T)+kn)$  time, where rd(T) is the radius of the tree [PS].

Megiddo and Cole first considered the special case in which the tree is a path, and give  $O(n(\log n)^2)$ -time and  $O(n\log n)$ -time algorithms, resp., for this case.<sup>1</sup> (Megiddo notes that an  $O(n\log n)$ -time algorithm is possible, using ideas from [FJ1].) For the problem on a tree, Megiddo presents an  $O(n(\log n)^3)$ -time algorithm [M], and Cole presents an  $O(n(\log n)^2)$ -time algorithm [C2]. The algorithm we present here runs in O(n) time, which is clearly optimal.

The second problem is the min-max tree k-partitioning problem [BPS]. The problem here is to delete k edges in the tree so as to minimize the weight of the heaviest of the resulting subtrees. This is closely related to the above problem. Becker, Perl and Schach give an algorithm that runs in  $O(k^3rd(T) + kn)$  time [BPS], and Perl and Vishkin give an efficient implementation of this algorithm that brings the time down to  $O(k rd(T)(k + \log \Delta) + n)$ , where  $\Delta$  is the maximum degree in the tree [PV]. Megiddo's and Cole's techniques apply to this problem with the bounds quoted above. Again, our techniques yield linear-time algorithms.

In [BP], Becker and Perl prove that the algorithms of [PS], [BPS] can be applied to tree partitioning problems that are subject to any one of a number of objective functions from a certain class. We suspect that our techniques can be applied to most (if not all) of the objective functions identified in [BP]. In [BP], a somewhat different algorithm is given for the tree partitioning problem in which the maximum diameter is minimized, with a running time of  $O(k^2rd(T)n)$ . We note that this problem is equivalent to the p-center problem E/V/k+1, which we discuss in [F2].

Our techniques can be extended and applied to the problem of finding a p-center in a tree [Hk], [Go1], [Go2], [Hn1], [Hn2], [KH], [CT], [FJ1]. In [F2] and [F3] we describe the extensions to our technique that yield O(n)-time algorithms for three variations of the p-center problem in trees. Also included in [F3] is a linear-time

<sup>&</sup>lt;sup>1</sup>All logarithms are to the base 2.

algorithm for the problem of computing recovery points in trees [CK], [CEIY].

Our results are a contribution to parametric search in several ways beyond merely producing a number of optimal algorithms. First, our results demonstrate that parallel algorithms are not essential or even that helpful in designing optimal algorithms for certain parametric search problems. We choose to ignore the link with parallel computation, and instead emphasize the generation and searching of partial orders. Instead of the two searching paradigms discussed in [C2], we focus on a third paradigm: searching within a collection of sorted matrices [FJ1], [FJ2]. Thus our work refocuses the emphasis on a carefully managed search. Second, we show how to manage the parametric search to produce intermediate information on which to base faster feasibility tests. To construct optimal algorithms for our problems, it appears to be necessary to interleave the narrowing of bounds on the optimal value with the construction of better feasibility tests. This is similar in spirit to the method in [CSSS], though the reasons that the feasibility tests can be sped up are quite different and the driving search procedure is quite different. Third, we design a pruning strategy for handling problems on trees, so that the feasibility test is not thwarted by trees that are exceptionally bushy.

Our paper is organized as follows. In section 2 we discuss some features of parametric search, and lay the foundation for the optimal algorithms that appear in subsequent sections. In section 3 we present our approach for solving the max-min partitioning problem on a path. In section 4 we build on the results in section 3 and present our approach for solving the max-min partitioning problem on a tree.

A preliminary version of this paper appeared in [F1].

# 2. Paradigms for parametric search

In this section we discuss some features of parametric search, and lay the foun-

dation for the optimal algorithms that appear in subsequent sections. We discuss path and tree partitioning problems in this section, note that the methods can be applied also to the p-center problem. We first review straightforward feasibility tests to test the feasibility of a search value in a tree. We then review how to represent all possible search values of a path within a sorted matrix. We next present our approach for searching using values from a collection of sorted matrices. Finally, we describe straightforward approaches for the path and the tree that are as good as any algorithms in [M] or [C2].

We first describe straightforward feasibility tests for cutting edges in a tree so as to either maximize the minimum size of any resulting component (max-min problem), or minimize the maximum size of any resulting component (min-max problem). Our faster feasibility tests will be built on these. For the max-min problem, the feasibility test takes a test value  $\lambda$ , and determines if (at least) k cuts can be placed in the tree such that no component has weight less than  $\lambda$ . For the max-min problem, let  $\lambda^*$  be the largest value of  $\lambda$  that passes the test. For the min-max problem, the feasibility test takes a test value  $\lambda$ , and determines if at most k cuts can be placed in the tree such that no component has weight greater than  $\lambda$ . For the min-max problem, let  $\lambda^*$  be the smallest value of  $\lambda$  that passes the test.

A straightforward test FTEST0 for the max-min problem in the tree is the following [M]. Root the tree at a vertex of degree 1, and initialize numcut to -1. Explore the tree starting at the root. At vertex v, set wgt(v) to the weight of v, and for each child w of v first explore the subtree rooted at w, and then add the resulting wgt(w) to wgt(v). After all children of v are explored, if  $wgt(v) \geq \lambda$ , then add 1 to numcut and reset wgt(v) to 0. Whenever numcut is incremented, except for the last time, the edge from v to the parent of v should be cut. When the traversal is complete, if  $numcut \geq k$ , then  $\lambda$  is feasible ( $\lambda \leq \lambda^-$ ), else  $\lambda$  is not feasible ( $\lambda > \lambda^-$ ). The

feasibility test uses O(n) time.

The feasibility test for the min-max problem is similar [KM]. Root the tree at a vertex of degree 1, and initialize numcut to 0. Explore the tree starting at the root. At vertex v, set wgt(v) to the weight of v, and explore the subtree rooted at each child w of v. Determine the child w' with largest  $wgt(\cdot)$  value such that  $wgt(v) + \sum_{wgt(w) \leq wgt(w')} wgt(w) \leq \lambda$ , using lexicography to break ties in  $wgt(\cdot)$  values. For each child w, if  $wgt(w) \leq wgt(w')$ , then add wgt(w) to wgt(v), else cut the edge from w to v and add 1 to numcut. Using a standard median-finding algorithm [BFPRT], the appropriate child w' can be found in time linear in the number of children. When the traversal is complete, if  $numcut \leq k$ , then  $\lambda$  is feasible ( $\lambda \leq \lambda^{-}$ ), else  $\lambda$  is not feasible ( $\lambda < \lambda^{-}$ ). Thus this feasibility test also uses O(n) time.

In a parametric search problem, the search can be restricted to considering values from a finite set of values. The desired value is the largest (in the case of a maxmin problem) or the smallest (in the case of a min-max problem) that passes the feasibility test. Let each value in the finite set be called a candidate value. We next discuss a data structure that contains all candidate values for problems on a path. This structure is based on ideas in [FJ1] and [M]. Let a matrix be called a sorted matrix if for every row the values are in nonincreasing order, and for every column the values are in nonincreasing order. Let the vertices on the path P be indexed from 1 to n. The set of values to be searched is the set of sums of weights of vertices from i to j, for all pairs  $i \leq j$ . The data structure contains these values, plus others, in the form of a sorted matrix that is succinctly represented and generated as follows. For  $i = 0, 1, \dots, n$ , let  $A_i$  be the sum of the weights of vertices 1 to i. Note that for any pair i, j with i < j, the sum of the weights from vertex i+1 to j is  $A_j - A_i$ . Let  $X_1$  be the sequence of sums  $A_n, A_{n-1}, \dots, A_1$ , and let  $X_2$  be the sequence of sums  $A_0, A_1, \dots, A_{n-1}$ . Then sorted matrix M(P) is the  $n \times n$  Cartesian matrix

 $X_1 - X_2$ , where the ij-th entry is  $A_{n+1-i} - A_{j-1}$ . In determining the above, we can use proper subtraction, i.e., a - b gives  $\max\{a - b, 0\}$ . Clearly, the values in any row of M(P) are in nonincreasing order, and the values in any column of M(P) are also in nonincreasing order. Note that M(P) can be represented in O(n) space in such a way that any matrix entry can be computed in constant time.

As an example, we show a vertex-weighted path P in Fig. 2.1, and its associated matrix M(P). The sequence  $X_1$  is listed vertically to the left of M(P), and the sequence  $X_2$  is listed horizontally above M(P), in such a way that the ij-th element of M(P) is to the right of the i-th element of  $X_1$  and beneath the j-th element of  $X_2$ .

We next describe a searching algorithm MSEARCH that combines the methods of [FJ1] and [FJ2], and adapts them for our purposes. Algorithm MSEARCH takes as its arguments a collection of sorted matrices, searching bounds  $\lambda_1$  and  $\lambda_2$ , and a stopping count. For a max-min problem,  $\lambda_1$  is feasible and  $\lambda_2$  is not. (The reverse holds for a min-max problem.) MSEARCH will produce a sequence of values one at a time to be tested for feasibility, with elements in the collection of matrices discarded as a result of each test. For a max-min problem, if  $\lambda > \lambda_1$  is feasible, then  $\lambda_1$  is reset to  $\lambda$ . Otherwise, if  $\lambda < \lambda_2$  is not feasible, then  $\lambda_2$  is reset to  $\lambda$ . (The reverse is done for a min-max problem.) MSEARCH will halt when the number of matrix elements remaining is no greater than the stopping count. The algorithm consists of a sequence of iterations. On each iteration the largest matrices are split into smaller submatrices, and three test values are produced in turn. On the basis of each test, certain of the submatrices are discarded. The stopping count can prevent the use of many iterations to discard relatively few elements by halting the search before all candidate values have been discarded. We assume that the product of the dimensions of each sorted matrix is a power of 4. If this is not the case, then the matrix can be padded out logically with big values. (Including these big values enlarges only the set of values to be tested, and does not change the underlying path.) Let the size of a matrix or a submatrix be the product of its dimensions.

We now specify MSEARCH for a max-min problem. Let a cell be a matrix or submatrix contained in a set C.

```
proc MSEARCH (set_of_matrices \mathcal{M}, integer stop_count, real \lambda_1, \lambda_2)
C \leftarrow \emptyset
Let cell_size equal the size of the largest matrix in M.
repeat
     Move from \mathcal{M} to \mathcal{C} all matrices whose size equals cell_size.
     if cell\_size > 1 then
          for each cell C in C do
               if both dimensions of C are greater than 1
               then Split C into four cells by halving each dimension.
                else Split C into four cells by quartering the larger dimension.
                endif
                Replace C by the resulting four cells.
          endfor
          cell\_size \leftarrow cell\_size/4
      endif
      repeat three times
          if cell\_size = 1
          then Let R be the multiset of values in the cells.
           else Let R be the multiset consisting of the smallest and the largest
                    element from each cell in C.
           endif
           Select the median element x in R.
           if \lambda_1 < x < \lambda_2
           then
                Use the designated feasibility test on element x.
                if x is feasible then \lambda_1 \leftarrow x else \lambda_2 \leftarrow x endif
                Discard from C any cell with no values in (\lambda_1, \lambda_2).
           endif
      endrepeat
until the total size of all matrices and cells in \mathcal{M} \cup \mathcal{C} is at most stop\_count
endproc
```

For the min-max problem MSEARCH is the same, except that  $\lambda_2$  is reset to x if x is feasible, and  $\lambda_1$  is reset to x if x is not feasible.

The following theorem is similar in spirit to Lemma 5 in [FJ1] and Theorem 2 in [FJ2], and its proof shares similarities with their proofs.

Theorem 2.1. Let  $\mathcal{M}$  be a collection of N sorted matrices  $\{M_1, M_2, \dots, M_N\}$  in which matrix  $M_j$  is of dimension  $m_j \times n_j$ ,  $m_j \leq n_j$ , and  $\sum_{j=1}^N m_j = m$ . Let q be the stopping count. The number of iterations needed by MSEARCH is  $O(\max\{\log \max_j \{n_j\}, \log(m/(q+1))\})$ , and the total time of MSEARCH exclusive of feasibility tests is  $O(\sum_{j=1}^N m_j \log(2n_j/m_j))$ .

**Proof.** Let  $M_j$  be a matrix in  $\mathcal{M}$ . Let i' be the last iteration in which cells are divided. Consider any iteration  $i \leq i'$ , and let c be the current cell size at the end of the iteration. If  $M_j$  is still in  $\mathcal{M}$  on the i-th iteration, let  $b_{ij} = 0$ . Otherwise, if  $c > n_j/m_j$ , then let  $b_{ij} = 2\sqrt{n_j m_j/c} - 1$ , else let  $b_{ij} = m_j$ . For any given value x, at most  $b_{ij}$  of the cells of  $M_j$  on iteration i will have smallest element less than x and largest element greater than x. Let  $B_i = \sum_{j=1}^N b_{ij}$ .

We claim that for all iterations in which cells are divided, the number of cells remaining at the end of the *i*-th iteration is no greater than  $2B_i + 2$ . The proof of the claim is by induction on *i*. Let  $ncells_i$  be the number of cells in C after cells have been quartered on iteration *i*, and let  $ncells_i'$  be the number of cells in C at the end of iteration *i*. Since  $B_0 = 0$  and  $ncells_0' = 0$ , the claim holds before the first iteration. For any iteration i,  $0 < i \le i'$ , there are, by the induction hypothesis,  $ncells_{i-1}' \le 2B_{i-1} + 2$  cells before the iteration begins. Suppose that  $K_i$  cells are transferred from M to C in iteration i. Then  $ncells_i \le 4(K_i + 2B_{i-1} + 2)$ . Note that  $b_{ij} \ge b_{i-1,j}$  for any  $M_j$  not in M on iteration i. (Equality holds if  $M_j$  was not in M on iteration i-1 and the cells resulting from  $M_j$  have one dimension equal to 1 on iteration i-1. Note that  $b_{ij} = b_{i-1,j} + 1$  for any matrix  $M_j$  whose cells have both dimensions greater than 1 on iteration i-1 but have one dimension equal to 1 on iteration i.) Thus  $B_{i-1} + K_i \le B_i$ , and thus  $ncells_i \le 8B_i + 8$ .

When the median is selected and cells are discarded, the number of cells discarded will be half of the following quantity: the current number of cells minus  $B_i$ . This follows since at most  $B_i$  cells have smallest element less than x and largest element greater than x, and thus x will induce a partition of the remaining cells into two equal size sets. The number of cells discarded on each of the three executions of the inner loop will be  $d_1 \geq \lfloor (ncells_i - B_i)/2 \rfloor$ ,  $d_2 \geq \lfloor (ncells_i - d_1 - B_i)/2 \rfloor$ , and  $d_3 \geq \lfloor (ncells_i - d_1 - d_2 - B_i)/2 \rfloor$ , resp. Then  $ncells_i' = ncells_i - d_1 - d_2 - d_3 \leq (ncells_i + 7B_i + 7)/8 \leq (8B_i + 8 + 7B_i + 7)/8 = (15/8)(B_i + 1) < 2B_i + 2$ . This concludes the proof of the claim.

Since the number of cells remaining at the end of the *i*-th iteration is no greater than  $2B_i + 2$ , the time to divide and select among cells on the *i*-th iteration is  $O(B_i)$ , which is  $O(\sum_{j=1}^N b_{ij})$ . It follows from the definition of  $b_{ij}$  that  $\sum_{i=1}^{i'} b_{ij}$  is  $O(m_j \log(2n_j/m_j))$ . Thus the time to perform all iterations  $i \leq i'$ , exclusive of feasibility tests, is  $O(\sum_{j=1}^N m_j \log(2n_j/m_j))$ . For each iteration  $i \geq i'$ , the cells selected among contain single elements, so that the number of elements is halved on each iteration. Thus all iterations i > i' will use O(m) time, exclusive of feasibility testing. This is dominated by the time to perform all iterations  $i \leq i'$ .

Since the current cell size is quartered on each iteration  $i \leq i'$ , the number of such iterations is  $O(\log \max_j \{n_j\})$ . Since the number of elements is halved for each iteration i > i', the number of such iterations is  $O(\log(m/(q+1)))$ .  $\square$ 

We are now ready to use MSEARCH to generate (relatively) simple algorithms for partitioning a path and partitioning a tree. These algorithms match the time of the algorithms in [C2], and set the stage for the improved algorithms that we present in the next two sections. We first describe a simple approach to the max-min problem on a path P. Initialize  $\lambda_1$  to 0 and  $\lambda_2$  to  $\infty$ . (For path and tree partitioning, we may take  $\infty$  to be the total weight of all vertices in the path or tree.) Run algorithm

MSEARCH on the set containing the single sorted matrix M(P) with stopping count 0. Use the straightforward feasibility test described above. When MSEARCH halts,  $\lambda^* = \lambda_1$ . (For the min-max problem,  $\lambda^* = \lambda_2$ .) By Theorem 2.1, MSEARCH will take O(n) total time, exclusive of the feasibility tests, and will produce a sequence of  $O(\log n)$  values to be tested. It follows that the total time for all feasibility tests is  $O(n \log n)$ . This corresponds to the times achieved by Megiddo and Cole for the path problem. We show how to do better in section 3.

We illustrate the above procedure (and also MSEARCH) using path P and in Fig. 2.1, with k=3. We set  $\lambda_1$  to 0 and  $\lambda_2$  to  $\infty$  and call MSEARCH on M(P). Matrix M(P) is first quartered into 4 cells of dimension  $4 \times 4$ . The set R formed from these cells is  $\{59, 3, 28, 0, 31, 0, 0, 0\}$ . The median of the set is 1.5 (the average of the lower median 0 and the upper median 3). For  $\lambda = 1.5$ , 6 cuts are required, so  $\lambda_1$  is reset to 1.5, and the cell with all values less than or equal to 1.5 is discarded, leaving three cells and a new set  $R = \{59, 3, 28, 0, 31, 0\}$ . The median of R is (3+31)/2 = 17. For  $\lambda = 17$ , 1 cut is required, so  $\lambda_2$  is reset to 17. But there are no cells with all values at most 17, so none gets discarded. The third selection produces the same result. On the next iteration of the outer repeat loop, the three cells are quartered into 12 cells, as shown in Fig. 2.2(a). Set R = $\{59, 45, 44, 23, 28, 20, 17, 0, 42, 25, 27, 3, 11, 0, 0, 0, 31, 22, 16, 0, 15, 0, 0, 0\}. \ \ The \ median$ of this set is (16+17)/2 = 16.5. For  $\lambda = 16.5$ , 1 cut is required, so  $\lambda_2$  is reset to 16.5. There are five cells with all values at least 16.5, and these are discarded. The remaining cells have  $R = \{17, 0, 27, 3, 11, 0, 0, 0, 16, 0, 15, 0, 0, 0\}$ . The median of this set is  $0 \le \lambda_1$ , so that the two cells with all values no larger than 0 are discarded. The remaining cells have  $R = \{17, 0, 27, 3, 11, 0, 16, 0, 15, 0\}$ . The median of this set is (3+11)/2 = $7 \le \lambda_1$ . No cells can be discarded, leaving the cells as pictured in Fig. 2.2(b). On the next iteration of the repeat loop, these cells are quartered, giving 20 cells of dimension  $1 \times 1$ . Then  $R = \{17, 6, 11, 0, 27, 12, 18, 3, 11, 9, 2, 0, 16, 1, 15, 0, 15, 7, 8, 0\}$ . The median is (8+9)/2 = 8.5. For  $\lambda = 8.5$ , 3 cuts are required, so  $\lambda_1$  is reset to 8.5. Discarding the half of the values and selecting the median gives 15. For  $\lambda = 15$ , 2 cuts are required, so  $\lambda_2$  is reset to 15. Discarding all but  $\{9, 11, 11, 12\}$  and selecting the median gives 11. For  $\lambda = 11$ , 3 cuts are required, so  $\lambda_1$  is reset to 11. All values but 12 are discarded. On the next iteration of the outer repeat-loop, this value is tested and found to require 3 cuts, so  $\lambda_1$  is reset to 12. With all values at this point discarded, MSEARCH will terminate with  $\lambda^* = \lambda_1 = 12$ .

We next describe a simple approach to the max-min problem on a tree. We first define an edge-path-partition of a tree rooted at a vertex of degree 1. Partition the edges of the tree into paths, where a vertex is an endpoint of a path if and only if it is of degree not equal to 2 with respect to the tree. Call any path in an edge-path-partition that contains a leaf in the tree a leaf-path. As an example, consider the vertex-weighted tree in Fig. 2.3(a). The edge-path-partition for T is shown in Fig. 2.3(b). There are 7 paths in the partition, as shown. Four of these paths are leaf-paths.

We now proceed with the approach for the tree. Initialize  $\lambda_1$  to 0 and  $\lambda_2$  to  $\infty$ . Initialize T to be the tree rooted at a vertex of degree 1. Repeat the following. Form an edge-path-partition of T. For each leaf-path  $P_j$  in the edge-path-partition, form sorted matrix  $M(P_j)$ . Run MSEARCH on this collection of matrices, with stopping count 0. We use the straightforward feasibility test described above. If T contains just one leaf, halt with  $\lambda^* = \lambda_1$ . (For the min-max problem,  $\lambda^* = \lambda_2$ .) If T contains more than one leaf, do the following. For each leaf-path  $P_j$ , infer the cuts in  $P_j$  such that each component in turn going up in  $P_j$ , except the highest component on  $P_j$ , has total weight as small as possible but greater than  $\lambda_1$ . (For the min-max problem, infer the cuts on  $P_j$  such that each component in turn going up  $P_j$  has total weight

as large as possible but less than  $\lambda_2$ .) Delete all vertices beneath the top vertex of each leaf-path  $P_j$ , and add to the weight of the top vertex in  $P_j$  the weight of the other vertices in the highest component of  $P_j$ . This leaves a smaller tree in which all leaf-paths in the original tree have been deleted. The smaller tree has at most half of the number of leaves of T. Reset T to be the smaller tree, and k to be the number of cuts remaining to be made in this tree. This concludes the description of the repeat loop, and with it the algorithm.

As an example we consider the max-min problem on the tree shown in Fig. 2.3(a), with k=3. After a call to MSEARCH given the matrices corresponding to the leafpaths,  $\lambda_1=9$  and  $\lambda_2=15$ . One cut can be placed between the vertices of weight 15 and 6. The weights of the two leaves of weight 4 are added to the weight of their parent, giving it weight 13. The weights of all descendants of the vertex with weight 2, except the weight of 15, are added to the vertex of weight 2, giving it a weight also of 13. Then all edges in the leaf-paths are deleted, and k is reset to 2. The edge-pathpartition of the resulting tree is shown in Fig. 2.4(a). There are two leaf-paths in this partition, with vertex weights 13, 4, and 3 on one, and 13 and 3 on the other. After a call to MSEARCH given the matrices corresponding to these leaf-paths,  $\lambda_1=9$  and  $\lambda_2=13$ . Two cuts can be placed, above each of the vertices of weight 13. The weight of the vertex of weight 4 is then added to its parent, giving it weight 7. Then all edges in the leaf-paths are deleted, and k is reset to 0. The edge-path-partition of the resulting tree is shown in Fig. 2.4(b). After a call to MSEARCH given the matrices corresponding to the leaf-paths,  $\lambda_1=12$  and  $\lambda_2=13$ . Thus  $\lambda^*=12$ .

For correctness of the tree algorithm, note that  $\lambda_1$  always corresponds to a feasible value, and that the cuts are inferred on leaf-paths assuming (correctly) that any subsequent value  $\lambda$  to be tested will have  $\lambda > \lambda_1$ . Also note that we choose the edge-path-partition to be a partition of edges rather than vertices, so that the inference of

cuts will be correct in the min-max version of the problem too.

We analyze the time as follows. By Theorem 2.1, each call to MSEARCH will produce  $O(\log n)$  values, and the time to produce these, exclusive of feasibility testing, will be  $O(\sum_{j=1}^{N} n_j) = O(n)$ . Since each feasibility test takes O(n) time, the test values generated by running MSEARCH can be tested in a total of  $O(n \log n)$  time. Since the number of leaves is at least halved on each such iteration, the number of iterations needed until all cuts have been made is  $O(\log n)$ . Thus the total time to partition the tree by this method is  $O(n(\log n)^2)$ . This beats the time of  $O(n(\log n)^3)$  for Megiddo's algorithm, and matches the time of  $O(n(\log n)^2)$  for Cole's algorithm. We show how to do better in section 4.

# 3. Partitioning a Path

In this section we present three algorithms to perform parametric search on a path of n vertices. Each algorithm is an improvement on the previous, and we present them one at a time to assist the reader in understanding our approach. The first algorithm uses two phases of search to achieve  $O(n \log \log n)$  time. The first phase gathers information with which subsequent feasibility tests can be performed in o(n) time. The second phase then completes the parametric search using this faster feasibility testing. The second algorithm gathers information on more than one phase, and takes  $O(n \log^{-} n)$  time.<sup>2</sup> The third algorithm uses a variety of refinements to reduce the time down to O(n). Our discussion focuses on the max-min problem; at the end of the section we identify the changes necessary for the min-max problem. Throughout this section we assume that the vertices of any path or subpath are indexed in increasing order from the start to the end of the path.

We describe several key ideas for fast feasibility testing in a path. We partition

<sup>&</sup>lt;sup>2</sup>The function  $\log^{n} k$  is the iterated logarithm of k, defined by  $\log^{n} 1 = \log^{n} 2 = 1$  and  $\log^{n} k = 1 + \log^{n} [\log k]$  for k > 2.

the vertices of the path P into subpaths as follows. For a given integer r, define an r-partition of a path P as a partition of the vertices of P into subpaths  $P_j$ , where each subpath except the last contains r vertices, and the last contains at most r vertices. For each subpath  $P_j$ , let  $f_j$  and  $t_j$  be the indices of the first and last vertices, resp., in  $P_j$ . Parametric search is used to generate a data structure that allows us to determine the number of cuts in most subpaths quickly. Let  $\lambda_1$  and  $\lambda_2$  be the parametric search bounds on  $\lambda^*$ . Define an active value to be any candidate value  $\lambda$  that arises from a subpath and satisfies  $\lambda_1 < \lambda < \lambda_2$ . For each vertex  $v_l$  on any subpath  $P_j$  that has no active values, two quantities ncut(l) and rmdr(l) can be determined. The quantity ncut(l) is the maximum number of cuts on the subpath from  $v_l$  to  $t_j$  such that each connected component in this subpath, with the exception of the last, has weight at least  $\lambda_2$ . (Since we know that  $\lambda_1$  is feasible, we set up  $ncut(\cdot)$  to test values  $\lambda > \lambda_1$  for feasibility. Since all candidate values arising from  $P_j$  have been discarded, the next value on  $P_j$  larger than  $\lambda_1$  is at least  $\lambda_2$ .) For convenience of description, we assume that a cut may be placed after  $t_j$ , producing a component of weight 0. (We compensate for this extra cut by subtracting 1 from the number of cuts.) The quantity rmdr(l) is the maximum weight possible for the last component, given that the number of edges cut in the subpath from  $v_i$  to  $t_j$  is ncut(i). Given the  $ncut(\cdot)$  and  $rmdr(\cdot)$  values, each feasibility test FTEST1 in the second phase runs in o(n) time.

We now describe algorithm PATH1. The first phase of PATH1 is as follows. Initialize  $\lambda_1$  to 0 and  $\lambda_2$  to  $\infty$ . Let f(n) be the largest power of 2 no larger than  $\log n$ . Form an f(n)-partition of path P. For each subpath  $P_j$ , form sorted matrix  $M(P_j)$ , giving a set  $\mathcal{M}_1$  of  $\lceil n/f(n) \rceil$  sorted matrices, each (except possibly the last) of dimension  $f(n) \times f(n)$ . Call algorithm MSEARCH with the set  $\mathcal{M}_1$  of matrices and a stopping count of  $n/(f(n))^2$ , using feasibility test FTEST0. At most  $n/(f(n))^2$  active values remain from  $\mathcal{M}_1$ . For each subpath  $P_j$  that contains no active values, compute the functions  $ncut(\cdot)$  and  $rmdr(\cdot)$ , using algorithm DIGEST1, to be described shortly.

The second phase of PATH1 is as follows. Let M(P) be the  $n \times n$  sorted matrix representing all sums on the path P. Run algorithm MSEARCH on the set  $\{M(P)\}$  with stopping count 0, using feasibility test FTEST1. At the termination of MSEARCH,  $\lambda^{\bullet} = \lambda_1$ . This completes the description of PATH1.

We illustrate PATH1 on path P in Fig. 2.1, with k = 3. (This is the same example that we discussed near the end of the previous section.) For path P, n=8 and thus f(n) = 2. The set  $\mathcal{M}_1$  is shown in Fig. 3.1. Procedure MSEARCH is called on  $\mathcal{M}_1$  with stopping count  $n/(f(n)^2)=2$ . These are first quartered, giving sixteen  $1 \times 1$  cells, and then  $R = \{17, 6, 11, 0, 11, 9, 2, 0, 16, 1, 15, 0, 15, 7, 8, 0\}$ . The median of these is (7+8)/2 = 7.5. For  $\lambda = 7.5$ , 3 cuts are required, so  $\lambda_1$  is reset to 7.5, and appropriate values are discarded, leaving  $R = \{17, 11, 11, 9, 16, 15, 15, 8\}$ . The median of these is (11+15)/2=13. For  $\lambda=13$ , 2 cuts are required, so  $\lambda_2$  is reset to 13, and appropriate values are discarded, leaving  $R = \{11, 11, 9, 8\}$ . The median of these is (9+11)/2 = 10. For  $\lambda = 10$ , 3 cuts are required, so  $\lambda_1$  is reset to 10, and 9 and 8 are discarded, leaving  $R = \{11, 11\}$ . This completes one iteration of the outer repeat-loop of MSEARCH. Since the number of values remaining is less than or equal to the stopping count, MSEARCH terminates with  $\lambda_1 = 10$  and  $\lambda_2 = 13$ . There are two subpaths with no active values, namely the subpath with the 5-th and 6-th vertices of P, and the one with the 7-th and 8-th vertices of P. Computing ncut and rmdr values gives ncut(5) = 1, rmdr(5) = 0, ncut(6) = 1, rmdr(6) = 0, ncut(7) = 1, rmdr(7) = 0, ncut(8) = 0, rmdr(8) = 8. The execution of the second phase of PATH1 follows the example near the end of the preceding section, except that FTEST1 is used, and whenever a test value  $\lambda$  does not satisfy  $\lambda_1 < \lambda < \lambda_2$ , the test need not be performed.

We next describe procedure DIGEST1 for the max-min problem, which computes

ncut(l) and rmdr(l) for all vertices  $v_l$  in subpath  $P_j$ . The approach is simple dynamic programming, for which we give the dynamic programming equations. Let  $w(u,v) = A_v - A_{u-1}$ , the sum of the weights of vertices from vertex u to vertex v. Let l range from  $t_j + 1$  down to  $f_j$ . If  $w(l,t_j) < \lambda_2$ , then ncut(l) = 0 and  $rmdr(l) = w(l,t_j)$ . Otherwise, ncut(l) = 1 + ncut(h+1) and rmdr(l) = rmdr(h+1), where h is the smallest index such that  $w(l,h) \geq \lambda_2$ . The dynamic programming equations can be implemented using a pass over the subpath from the last vertex to the first with pointers l and h monotonically nonincreasing.

Lemma 3.1. Let  $P_j$  be a subpath that has no corresponding active values. Then DIGEST1 computes ncut(l) and rmdr(l) for all vertices  $v_l$  in  $P_j$  in linear time. Proof. For each of r values of h, a constant amount of work is done. The remaining work can be apportioned as constant for each of O(r) values of l.  $\square$ 

We next describe feasibility test *FTEST1* that can be performed using the *ncut* and rmdr values. The key idea is to use binary search to find the first cut in each subpath, and then, if the subpath has no active values, to index into the data structures to find the number of cuts remaining in the subpath, and the size of the last component. Let  $\lambda$  be the value to be tested, with  $\lambda_1 < \lambda < \lambda_2$ .

```
proc FTEST1 (path P, integer k, real \lambda)

numcut \leftarrow -1; remainder \leftarrow 0

for each subpath P_j starting with the first do

if remainder + w(f_j, t_j) < \lambda

then remainder \leftarrow remainder + w(f_j, t_j)

else Binary search to find the smallest l such that w(f_j, v_l) + remainder \geq \lambda.

if v_l is t_j

then numcut \leftarrow numcut + 1; remainder \leftarrow 0

else if there are no active values for P_j

then numcut \leftarrow numcut + 1 + ncut(l+1); remainder \leftarrow rmdr(l+1)

else

Scan P_j starting with v_{l+1} to compute k' and r', the values
```

```
of ncut(l+1) and rmdr(l+1), resp., as though \lambda_2 = \lambda. numcut \leftarrow numcut + k'; \ remainder \leftarrow r' endif endif endif endfor if numcut \geq k then \lambda is feasible (\lambda \leq \lambda^*) else \lambda is not feasible (\lambda > \lambda^*) endproc
```

Lemma 3.2. Let P be a path of n vertices partitioned into  $\lceil n/r \rceil$  subpaths, each of size at most r. Let all but at most  $n/r^2$  subpaths have the functions  $ncut(\cdot)$  and  $rmdr(\cdot)$  computed for all vertices on them. Algorithm FTEST1 determines the feasibility of a test value in  $O((n/r)\log r)$  time.

**Proof.** There are  $\lceil n/r \rceil$  subpaths to be examined. Consider the examination of subpath  $P_j$ . The search for  $v_l$  uses  $O(\log r)$  time. If there are no active values for  $P_j$ , then all other operations performed in examining  $P_j$  take constant time. There are O(n/r) such paths, which take  $O((n/r)\log r)$  time in total. If there are active values for  $P_j$ , then examining the path takes O(r) time. But there are only  $O(n/r^2)$  subpaths with active values, which take O(n/r) time in total.  $\square$ 

Theorem 3.3. Algorithm PATH1 solves the max-min k-partitioning problem on a path of n vertices in  $O(n \log \log n)$  time.

Proof. By Theorem 2.1, the number of test values produced on the first call to MSEARCH is  $O(\log \log n)$ , and these are produced in O(n) time. Since each feasibility test takes O(n) time, the total time for feasibility testing is  $O(n \log \log n)$ . By Lemma 3.1 the total time to compute the ncut and rmdr values for  $O(n/\log n)$  subpaths is O(n). By Theorem 2.1, the second call to MSEARCH produces  $O(\log n)$  test values in O(n) time. By Lemma 3.2, each call to feasibility test FTEST1 takes  $O(n \log \log n/\log n)$ , or  $O(n \log \log n)$  for all tests.  $\square$ 

Applying the above strategy with more than two phases produces our second algorithm, PATH2, which runs in  $O(n \log^{n} n)$  time. The idea is to start with the straightforward feasibility test, and bootstrap our computation by repeatedly constructing data structures for ever faster feasibility tests. In each phase, we generate these data structures as we search among sets of sorted matrices that represent larger and larger subpaths. Initially all values are active, so that  $ncut(\cdot)$  and  $rmdr(\cdot)$  are uninitialized, and FTEST1 will perform initially much like FTEST0. We call PATH2 only when  $n \geq 16$ .

```
Proc PATH2 (path P, integer k)

Let f(n) be the largest power of 2 no larger than (\log n)^2.

Let lev be the smallest nonnegative integer such that f^{lev}(n) = 16.

\lambda_1 \leftarrow 0; \lambda_2 \leftarrow \infty

for each value of i from lev down to 0 do

r_i \leftarrow f^i(n)

Form an r_i-partition of P.

For each subpath P_j, form a sorted matrix for P_j.

Let \mathcal{M}_i be the resulting set of sorted matrices.

Call MSEARCH on \mathcal{M}_i, with stopping count n/r_i^2, using FTEST1.

Identify the at most n/r_i^2 active values between \lambda_1 and \lambda_2.

For each subpath P_j with no active values, call DIGEST1.

endfor

\lambda^* \leftarrow \lambda_1
endproc
```

Theorem 3.4. Algorithm PATH2 solves the max-min k-partitioning problem on a path of n vertices in  $O(n \log^2 n)$  time.

Proof. There are lev + 1 phases, where lev is  $O(\log^* n)$ . Each call to MSEARCH with matrices of size  $r_i \times r_i$  uses O(n) time and generates  $O(\log r_i)$  test values. Performing all feasibility tests on phase lev uses O(n) time, and on phase i < lev uses  $O((n/r_{i+1})\log r_{i+1}\log r_i)$  time. Since  $r_{i+1}$  is  $O((\log r_i)^2)$ , the total time for all feasibility tests is O(n). The time to compute  $ncut(\cdot)$  and  $rmdr(\cdot)$  values at the end of the phase is also O(n). Thus the time for each of the  $O(\log^* n)$  phases is O(n).  $\square$ 

We now discuss our third algorithm PATH3, which uses linear time. In PATH2, O(n) time per phase is spent on each of two activities: computing  $ncut(\cdot)$  and  $rmdr(\cdot)$  values, and producing test values. We modify PATH2 to use O(n) time in total on these activities. First, we use the  $ncut(\cdot)$  and  $rmdr(\cdot)$  data structures already generated to generate the  $ncut(\cdot)$  and  $rmdr(\cdot)$  data structures on the next phase. Second, we call MSEARCH on sets of trimmed submatrices to produce our search values more efficiently.

We first describe our routine DIGEST2, which updates the arrays containing  $ncut(\cdot)$  and  $rmdr(\cdot)$  more efficiently than DIGEST1. When MSEARCH terminates on phase i, all of the at most  $n/r_i^2$  active values fall between  $\lambda_1$  and  $\lambda_2$ . For each subpath  $P_j$  with no active values, we want to determine ncut and rmdr values. We shall avoid computing information that we do not need. We shall compute ncut(l) and rmdr(l) for each vertex  $v_l$  with  $w(f_j, l) \leq \lambda_2$ . To assist in computing these data structures, we shall also keep track of next(l), the index of the vertex  $v_q$  such that  $rmdr(l) = w(q, t_j)$ . Before the searching phases begin, initialize ncut(l) to be 0 and next(l) to be l for each vertex  $v_l$ . We do not maintain  $rmdr(\cdot)$  explicitly for any vertex  $v_l$  such that ncut(l) = 0.

Let  $P_j$  be a subpath in the  $r_i$ -partition. We now describe how to generate the appropriate values for  $P_j$ , given the ncut, rmdr and next values for the subpaths of  $P_j$  that are in the  $r_{i+1}$ -partition. Our approach is to identify those vertices of  $P_j$  that are within  $\lambda^*$  of  $f_j$  and whose ncut value will be increased on the phase. These vertices are inserted into a queue. We then sweep through  $P_j$ , examining the vertices in the queue, and inserting additional vertices in the queue if they follow a cut inferred by a vertex in the queue. We save the vertices examined on a stack, so that ncut, rmdr and next values can be computed for the vertices examined, in reverse order, once the sweep through  $P_j$  is complete. To make the sweep efficient, the entries in the queue

should be in increasing order. However, when we sweep through a subpath of  $P_j$ , the entries generated for the queue are in either increasing order, or in rotated increasing order (i.e., some rotation of the whole list gives a list in increasing order). Why the entries to the queue are generated in this order is explained in the proof of Lemma 3.5. The queue entries should be rearranged periodically, so that they are considered in sorted order.

We now restate the above strategy in detail. Recall that  $f_j$  and  $t_j$  are the indices of the first and last vertices, resp., in  $P_j$ .

```
proc DIGEST2 (subpath P_j) /* P_j contains no active values */
if w(f_i, t_i) \geq \lambda_2
then
     Initialize a queue with a mark.
     for each vertex v_l in turn while w(f_j, l-1) < \lambda_2 and w(next(l), t_j) \geq \lambda_2 do
          Insert l into the queue.
     end
     Initialize a stack to be empty.
     while the queue contains more than just the mark do
          if the mark is at the front of the queue
          then
               Put the queue in sorted order, with the mark at the end.
               Set q to be the largest value in the queue.
          endif
          Extract l from the queue.
          if next(l) = l
          then while w(l, q-1) < \lambda_2 do q \leftarrow q+1 endwhile
          else q \leftarrow next(l)
          endif
          Push the pair (l,q) onto the stack.
          if w(q,t_j) < \lambda_2 then rmdr(q) \leftarrow w(q,t_j) endif
          If q is not in the queue and w(q,t_j) \geq \lambda_2, then insert q into the queue.
     endwhile
     while the stack is not empty do
          Pop a pair (l,q) from the stack.
          rmdr(l) \leftarrow rmdr(q); next(l) \leftarrow next(q).
          if ncut(l) = 0 then ncut(l) \leftarrow 1 + ncut(q) else ncut(l) \leftarrow ncut(l) + ncut(q)
      endwhile
endif
```

#### endproc

**Lemma 3.5.** Over all phases, the time for all calls to DIGEST2 is O(n).

**Proof.** First, we bound the time to scan by increments of 1 for a vertex  $v_q$  given vertex  $v_l$  when ncut(l) = 0. A vertex  $v_l$  is in the queue only if the total weight to the end of the path is at least  $\lambda^*$ . After the scan, ncut(l) > 0 and  $next(l) \ge q$ , so that the vertices scanned when ncut(l) = 0 are not scanned again. Thus in the linear scans over vertices when ncut(l) = 0, each vertex  $v_q$  is scanned at most once. Thus the total charge for this is O(n) over all phases.

Next, we bound the time to handle vertices inserted into the queue after the queue initialization as follows. Any vertex is inserted into the queue after the queue initialization at most once. This follows since the  $next(\cdot)$  function is set to point beyond this vertex for the next phase. Thus the total cost of handling such vertices is O(n) over all phases.

Finally, we bound the cost of handling a vertex inserted into the queue during queue initialization on a phase. Consider a vertex  $v_l$  inserted into the queue during queue initialization. On the first phase when this happens, charge 1 to the vertex. Over all phases each vertex can be so charged at most once, yielding a total cost of O(n) overall. Since  $v_l$  is inserted into the queue during queue initialization, there is at least one cut placed on the subpath containing  $v_l$  in this phase. Consider any subsequent phase i on which the vertex  $v_l$  is inserted into the queue during queue initialization. Let  $P_{j_l}$  be the subpath of  $P_j$  that contains  $v_l$ . No subpath of  $P_j$  following  $P_{j_l}$  contains a vertex inserted into the queue during queue initialization on phase i, since the total weight of vertices on  $P_{j_l}$  is at least  $\lambda^*$ . Furthermore, the subpaths of  $P_j$  (if any) preceding  $P_{j_l}$  have total weight less than  $\lambda^*$ , since  $v_l$  was inserted into the queue during queue initialization. Hence this is the first phase that vertices on such subpaths (if any) are inserted into the queue during queue initialization. Thus at

most  $r_{i+1}$  vertices, all from subpath  $P_{j_i}$ , are the only vertices from the  $r_i$  vertices of subpath  $P_j$  that are inserted into the queue during queue initialization for a second or greater time on phase i. Thus the total charge for phase i is  $(n/r_i)r_{i+1}$  for handling these vertices. Over all phases this latter cost is O(n).

We bound the time for all queue rearrangements as follows. If l and l' are two entries in the portion of the queue after the mark, then ncut(l) and ncut(l') differ by at most one. In fact as one scans the entries in the queue after the mark, then the ncut value stays constant while the rmdr value is nonincreasing, until at some index (if it exists), the ncut value decreases by 1, the rmdr value increases, and then for the remaining entries the ncut value stays constant while the rmdr value is once again nonincreasing. Thus the queue at the time of reorganization is a rotated sorted list. It takes time proportional to its length to reorganize a rotated sorted list. Because of the manner in which marks are handled, a vertex participates in a queue reorganization once for each time it is inserted into the queue. By the above arguments, the total number of insertions into the queue over all phases in O(n). Thus the time for all queue reorganizations is O(n).  $\square$ 

We next discuss how to generate sets of submatrices that are supplied as arguments to MSEARCH, so that the total time of MSEARCH over all phases is O(n). If a value in a sorted matrix is known to be too small to be  $\lambda^*$ , then we would like not to include it in the set supplied to MSEARCH. If a value is known to be too large, then on subsequent phases, it and any values that can be deduced by matrix position to be larger than it should be ignored.

On phase i, we consider a set  $\mathcal{M}_i^T$  of submatrices of M(P). To identify small and large values, we consider a set of thin matrices generated as follows. For each submatrix in  $\mathcal{M}_i^T$ , we take the first row as a thin matrix, and the first column as a thin matrix. We call MSEARCH on this resulting set  $\mathcal{T}_i$  of thin matrices, with stopping

count  $n/r_i$ . When MSEARCH returns, only  $\lfloor n/r_i \rfloor$  values in the thin matrices are still active. The other values are thus characterized as too small to be  $\lambda^*$  or too large to be  $\lambda^*$ .

We trim (logically) each submatrix in  $\mathcal{M}_{i}^{\mathcal{I}}$  to remove any row whose first element is too small, and any column whose first element is too small. We then call MSEARCH to search in the set  $\mathcal{M}_{i}^{\mathcal{I}}$  of resulting trimmed submatrices. We shall also include the set  $\mathcal{A}_{i}$  of elements remaining as active from the previous phase. We explain shortly why these are included.

We cannot trim a row or column whose first element is too large, since this might result in  $\lambda^*$  being discarded. Thus we cannot avoid leaving these rows or columns in. We make sure however that these large values are not considered in subsequent phases. On subsequent phases we could delete the corresponding rows and columns from M(P) and then compact M(P), but this appears to be too expensive to do. Instead, we extract from M(P) a set  $\mathcal{M}_{i-1}$  of submatrices to be considered on phase i-1. None of these submatrices contain values that are in the same row or column as a large value, or to the left and above, or to the right and below, a large value. There are  $O(n/r_i)$  such submatrices, which are not in general square. While subpaths do not change, each is represented by a set of submatrices that do not in general contain all sums associated with the subpath. Also, values that are still active must not be lost from consideration, since they might be in a row or column that is deleted. To prevent losing such values, any value still active at the end of phase i is brought along to the next phase, and the set  $\mathcal{A}_{i-1}$  of these values needs to be included in some of the calls to MSEARCH.

We note that there is a natural order on submatrices in  $\mathcal{M}_i$ . Let M' and M'' be distinct submatrices of  $\mathcal{M}_i$ . Any row of M(P) does not contain elements of both M' and M'', and similarly for columns. Without loss of generality assume that an element

from M' is in a higher-indexed row of M(P) than any element from M''. Then every element of M'' is in a column of M(P) of higher index than any element in M'. We assume that the submatrices of  $\mathcal{M}_i$  are maintained in order from lower left of M(P) to upper right of M(P). At the beginning of phase i the elements of the matrices in  $\mathcal{M}_i$  are intersected with the elements in the matrices of the set  $\mathcal{M}_i^P$  of matrices for the subpaths in the  $r_i$ -partition of P to give a set of elements stored in submatrices  $\mathcal{M}_i^T$ . (This intersection should not be carried out element by element. Instead, it is sufficient to identify each maximal submatrix of M(P) that is a submatrix of a matrix in  $\mathcal{M}_i$  and also a submatrix of a matrix in  $\mathcal{M}_i^P$ . We denote this operation by the symbol  $\mathbb{Q}$ .) We now summarize our algorithm PATH3, which we call only when  $n \geq 16$ .

```
proc PATH3 (path P, integer k)
Let f(n) be the largest power of 2 no larger than (\log n)^2.
Let lev be the smallest nonnegative integer such that f^{lev}(n) = 16.
\mathcal{M}_{lev} \leftarrow \{M(P)\}
\mathcal{A}_{lev} \leftarrow \emptyset
for each vertex v_l do ncut(l) \leftarrow 0; next(l) \leftarrow l endfor
\lambda_1 \leftarrow 0; \ \lambda_2 \leftarrow \infty.
for each value of i from lev down to 0 do
      r_i \leftarrow f^i(n)
      Form the r_i-partition of P, and the set \mathcal{M}_i^{\mathcal{P}} of submatrices of M(P).
      \mathcal{M}_{i}^{\mathcal{I}} \leftarrow \mathcal{M}_{i} \cap \mathcal{M}_{i}^{\mathcal{P}}.
       Form the set T_i of thin matrices for \mathcal{M}_i^T, by taking the first row
            and the first column from each matrix in \mathcal{M}_{i}^{2}.
       Call MSEARCH on T_i, with stopping count n/r_i.
       Initialize the set of trimmed submatrices \mathcal{M}_i^T to be empty.
       for each submatrix M' in \mathcal{M}_i^T do
            Insert a corresponding matrix M'' into \mathcal{M}_{i}^{T}, where M'' is M' with
                  every row or column that is headed by a small value deleted.
       endfor
       Call MSEARCH on \mathcal{M}_i^T \cup \mathcal{A}_i with stopping count n/r_i^2.
       Let A_{i-1} be the elements remaining upon the return from MSEARCH.
       Initialize \mathcal{M}_{i-1} to be \mathcal{M}_i.
       for each submatrix M' in \mathcal{M}_{i-1} that contains a large value do
            Delete any element in the same row or column as a large value.
```

from M' is in a higher-indexed row of M(P) than any element from M''. Then every element of M'' is in a column of M(P) of higher index than any element in M'. We assume that the submatrices of  $\mathcal{M}_i$  are maintained in order from lower left of M(P) to upper right of M(P). At the beginning of phase i the elements of the matrices in  $\mathcal{M}_i$  are intersected with the elements in the matrices of the set  $\mathcal{M}_i^p$  of matrices for the subpaths in the  $r_i$ -partition of P to give a set of elements stored in submatrices  $\mathcal{M}_i^{\mathcal{I}}$ . (This intersection should not be carried out element by element. Instead, it is sufficient to identify each maximal submatrix of M(P) that is a submatrix of a matrix in  $\mathcal{M}_i$  and also a submatrix of a matrix in  $\mathcal{M}_i^p$ . We denote this operation by the symbol  $\mathbb{Q}$ .) We now summarize our algorithm PATH3, which we call only when  $n \geq 16$ .

```
proc PATH3 (path P, integer k)
Let f(n) be the largest power of 2 no larger than (\log n)^2.
Let lev be the smallest nonnegative integer such that f^{lev}(n) = 16.
\mathcal{M}_{lev} \leftarrow \{M(P)\}
\mathcal{A}_{lev} \leftarrow \emptyset
for each vertex v_l do ncut(l) \leftarrow 0; next(l) \leftarrow l endfor
\lambda_1 \leftarrow 0; \ \lambda_2 \leftarrow \infty.
for each value of i from lev down to 0 do
      r_i \leftarrow f'(n)
      Form the r_i-partition of P, and the set \mathcal{M}_i^P of submatrices of M(P).
      \mathcal{M}_{i}^{I} \leftarrow \mathcal{M}_{i} \cap \mathcal{M}_{i}^{p}.
      Form the set T_i of thin matrices for \mathcal{M}_i^T, by taking the first row
            and the first column from each matrix in \mathcal{M}_{i}^{\mathcal{I}}.
       Call MSEARCH on T_i, with stopping count n/r_i.
       Initialize the set of trimmed submatrices \mathcal{M}_{i}^{T} to be empty.
      for each submatrix M' in \mathcal{M}_i^{\mathcal{I}} do
            Insert a corresponding matrix M'' into \mathcal{M}_{i}^{T}, where M'' is M' with
                  every row or column that is headed by a small value deleted.
       endfor
       Call MSEARCH on \mathcal{M}_i^T \cup \mathcal{A}_i with stopping count n/r_i^2.
       Let A_{i-1} be the elements remaining upon the return from MSEARCH.
       Initialize \mathcal{M}_{i-1} to be \mathcal{M}_i.
       for each submatrix M' in \mathcal{M}_{i-1} that contains a large value do
            Delete any element in the same row or column as a large value.
```

Delete any element that is to the left and above a large value, or to the right and below a large value. endfor

For each subpath  $P_j$ , call DIGEST2.

endfor

 $\lambda^* \leftarrow \lambda_1$  endproc

We illustrate our approach in Figures 3.2 and 3.3. Suppose that n=16, so that M(P) is a  $16 \times 16$  matrix. We show this matrix symbolically, with an asterisk in each element position. We start with  $\mathcal{M}_{lev}$  containing the one matrix M(P). This is shown in Fig. 3.2(a) by circling each asterisk. For the sake of illustration, suppose that  $(\log n)^2$  were equal to 4. (Clearly, this is not true, but choosing a value of n which is a power of 2 and for which f(n) < n makes the example a bit big.) Then our path is partitioned into 4 subpaths. The four submatrices in  $\mathcal{M}_{lev}^{\mathcal{P}}$ , corresponding to the four subpaths are shown in bold in Fig. 3.2(b). The elementwise intersection of  $\mathcal{M}_{lev}$  and  $\mathcal{M}_{lev}^{\mathcal{P}}$  gives a set  $\mathcal{M}_{lev}^{\mathcal{I}}$  that is the same as  $\mathcal{M}_{lev}^{\mathcal{P}}$ . The set of elements from the members of  $\mathcal{T}_{lev}$ , consisting of the first row and first column of each matrix in  $\mathcal{M}_{lev}^{\mathcal{I}}$  is shown in Fig. 3.2(c). The characterization of all but at most n/4 of these elements is shown in Fig. 3.2(d). The elements in the thin matrices that are not characterized as large (L) or small (S) are indicated by a question mark. The set  $\mathcal{M}_{lev}^{\mathcal{I}}$  of trimmed submatrices is shown in Fig. 3.3(b).

**Theorem 3.6.** Algorithm *PATH3* solves the max-min k-partitioning problem on a path of n vertices in O(n) time.

**Proof.** We analyze the time for manipulating and searching matrices as follows. We start phase i with a set  $\mathcal{M}_i$  of  $O(n/r_{i+1})$  submatrices of M(P) and a set  $\mathcal{A}_i$  of  $O(n/r_{i+1}^2)$  elements. Since there are  $O(n/r_{i+1})$  submatrices in  $\mathcal{M}_i$ , the time to

form  $\mathcal{M}_{i}^{\mathcal{I}}$  is  $O(n/r_{i+1})$ . There are  $O(n/r_{i+1})$  of these submatrices. Thus there are  $O(n/r_{i+1})$  thin matrices, and the time to form the set  $T_i$  of thin matrices is  $O(n/r_{i+1})$ . By Theorem 2.1, the call to MSEARCH on  $T_i$ , uses  $O((n/r_{i+1})\log r_i)$  time to produce  $O(\log r_i)$  search values. Given the values remaining from this search, it takes  $O(n/r_{i+1})$  time to generate the trimmed submatrices  $\mathcal{M}_i^T$ . Let the l-th trimmed submatrix have dimension  $n_l \times m_l$ . Let large(i) be the number of large values produced by searching in the set of thin matrices on phase i. Since no two such large values can appear in the same row or in the same column of M(P),  $\sum_{i} large(i) \leq 2n$ . A bound on  $\sum_{l} n_{l} + m_{l}$  is  $2n/r_{i} + large(i)$ . The most time is used in producing search values when as many trimmed matrices as possible are of dimensions  $r_i \times r_i$ . The number of these is  $O((n/r_i + large(i))/r_i)$ . Thus by Theorem 2.1, the call to MSEARCH on  $\mathcal{M}_i^T \cup \mathcal{A}_i$  uses  $O((n/r_i + large(i)) + n/r_{i+1})$  time to produce  $O(\log r_i)$  search values. The time to form  $\mathcal{M}_{i-1}$  using the large values resulting from the call to MSEARCH on the thin matrices is  $O(n/r_{i+1})$ . Thus the time to manipulate matrices and produce search values on phase i is  $O((n/r_{i+1}) \log r_i + large(i))$ , which is  $O(n/\log r_i + large(i))$ . This sums to O(n) over all phases. By Lemma 3.2, the time to test a value for feasibility is  $O((n/r_{i+1})\log r_{i+1})$ . Thus all feasibility tests on phase i uses time that is  $O((n/r_{i+1})\log r_{i+1}\log r_i)$ , which is  $O((n/\log r_i)\log\log r_i)$ . This sums to O(n) over all phases.

Finally, by Lemma 3.5 the time to generate the functions ncut and rmdr over all phases is O(n).  $\square$ 

We briefly survey the differences needed to solve the min-max problem. Let ncut(l) be the minimum number of cuts on the subpath in  $P_j$  from  $v_l$  to  $t_j$  such that each connected component in this subpath, has weight at most  $\lambda_1$ , and rmdr(l) is the minimum weight possible for the last component, given that the number of edges cut in  $P_j$  is ncut(l). Procedure DIGEST1 can similarly be described by a set of dynamic

programming equations: If  $w(l,t_j) < \lambda_1$ , then ncut(l) = 0 and  $rmdr(l) = w(l,t_j)$ . Otherwise, ncut(l) = 1 + ncut(h+1) and rmdr(l) = rmdr(h+1), where h is the largest index such that  $w(l,h) \leq \lambda_1$ . Procedure FTEST1 is similar except that l is the largest index such that  $w(f_j,v_l) + remainder \leq \lambda$ , path  $P_j$  is scanned as though  $\lambda_1 = \lambda$ , 1 is subtracted after completion of the for-loop if remainder = 0, and  $\lambda$  is feasible if and only if  $numcut \leq k$ . Algorithms PATH1, PATH2, and PATH3 are similar, except that at termination,  $\lambda^{\bullet} = \lambda_2$ . Procedure DIGEST2 is similar, except that any index l inserted into the queue satisfies  $w(f_j, l) \leq \lambda_1$  and  $w(next(l), t_j) \geq \lambda_1$ , low(j) is the largest index such that  $w(f_j, low(j)) \leq \lambda_1$ , high(j) is the smallest index such that  $w(high(j), t_j) \leq \lambda_1$ , q is incremented while  $w(l, q) \leq \lambda_1$ , and the other tests should have  $\lambda_1$  rather than  $\lambda_2$ .

Theorem 3.7. The min-max k-partitioning problem can be solved on a path of n vertices in O(n) time.

Proof. The above changes will not affect the asymptotic running time of algorithm *PATH3*. □

### 4. Partitioning a Tree

In this section we present three algorithms to perform parametric search on a tree of n vertices. The structure of this section is similar to that of the previous section, but concentrates on the additional ideas needed to deal with a tree. The first algorithm uses two phases of search to achieve  $O(n(\log \log n)^2)$  time. The first phase gathers information with which subsequent feasibility tests can be performed in o(n) time. The second phase then completes the parametric search using this faster feasibility testing. The second algorithm gathers information on more than one phase, and takes  $O(n \log^2 n)$  time. The third algorithm uses a variety of refinements to reduce the time down to O(n). Throughout this section we assume that the tree

is rooted, that paths in the tree run from descendants to ancestors and are indexed in this order within any such designated path.

We describe several key ideas for fast feasibility testing in a tree. A feasibility test will not be efficient if there are too many paths in the edge-path-partition. Thus we may need to prune the tree to reduce the number of leaves, and hence the number of paths in the edge-path-partition of the resulting tree. Define a vertex-path-partition of a tree T as a set of paths of tree T formed as follows. Find an edge-path-partition of T, and delete the top vertex from each path except the path containing the root of T. For each path, let the bottom vertex of the path with respect to the tree be its first vertex. For a tree T and desired subpath length r we define the path-tree  $T_c$  as follows. Form a vertex-path-partition of T, and form an T-partition of each path in the vertex-path-partition. Let  $T_c$  be a tree in which each node represents a subpath formed during the T-partition, and node T0 is a parent of node T1 in the subpath for T2 is a child of the lowest vertex in the subpath for T3.

As an example, consider the vertex-weighted tree T in Fig. 4.1(a). (This is the same tree that was used to illustrate the edge-path-partition in Fig. 2.1.) The vertex-path-partition for T is shown in Fig. 4.1(b). The r-partition for each path in the vertex-path-partition is shown in Fig. 4.2(a), with r=2. Note that there is only one path containing more than 2 vertices in the vertex-path-partition, in the lower right corner of Fig. 4.1(b). This path is split accordingly. The path-tree  $T_c$  for T with r=2 is shown in Fig. 4.2(b). Here some of the nodes (those representing subpaths that contain more than one vertex) are depicted as large "blobs" so that the correspondence to Fig. 4.2(a) is clearer.

We now describe our first algorithm, TREE1, for partitioning a tree. The first phase reduces the problem of partitioning a tree to the problem of partitioning a tree

with fewer than  $2n/(\log n)^2$  leaves, and then generates a data structure for a fast feasibility test. Note that when there are at least  $2n/(\log n)^2$  leaves, at least half of the leaf-paths are of length at most  $(\log n)^2$ . Initializing  $T_c$  to T indicates that we initially view each subpath in T' to be of length 1.

```
proc TREE1 (tree T, integer k)
\lambda_1 \leftarrow 0; \ \lambda_2 \leftarrow \infty
T' \leftarrow T; k' \leftarrow k; T_c \leftarrow T
/* phase 1 */
r \leftarrow \lfloor (\log n)^2 \rfloor
Form an edge-path-partition of T'.
while there are at least 2n/r leaves in T' do
     Let \mathcal{L} be the set of sorted matrices for leaf-paths of length at most r.
     Let n' be the number of vertices on these leaf-paths.
     Call MSEARCH on \mathcal{L} with stopping count n'/(2r), using FTEST2 for T_c and k.
     for each leaf-path with no active values do
          Identify the cuts on the path, and subtract their number from k'.
          Add to the top vertex the weight of the other vertices in its component.
          Remove the leaf-path from T' and modify the edge-path-partition of T'.
      endfor
endwhile
Form the vertex-path-partition of T',
      and form an r-partition of each path in the vertex-path-partition.
Let M be the set of sorted matrices for the resulting subpaths.
Call MSEARCH on \mathcal{M}, with stopping count n/r^2, using FTEST2 for T_c and k.
For each subpath P_j in T' with no active values, call DIGEST1.
Reset T to a copy of T', reset k to be k', and update T_c for T using r.
/* phase 2 */
Form an edge-path-partition of T'.
while there is more than one leaf in T' do
      Let \mathcal{M} be the set of sorted matrices for paths in the edge-path-partition.
      Call MSEARCH on \mathcal{M} with stopping count 0, using FTEST2 for T_c and k.
      for each leaf-path do
          Identify the cuts on the path, and subtract their number from k'.
           Add to the top vertex the weight of the other vertices in its component.
           Remove the leaf-path from T' and modify the edge-path-partition of T'.
      endfor
 endwhile
 \lambda^* \leftarrow \lambda_1
 endproc
```

We next describe the sublinear feasibility test FTEST2 for  $T_c$  and k. As before we assume that  $\lambda_1 < \lambda < \lambda_2$ . Perform a traversal of  $T_c$ , starting at the root. At node v, representing subpath  $P_j$ , do the following. Set remainder(v) to 0 and set numcut(v) to -1 if v is the root and to 0 otherwise. Then recursively explore each child w of v. For each child w, add remainder(w) to remainder(v) and add numcut(w) to numcut(v). Then handle the subpath represented by v by executing the body of the for-loop in FTEST1, but using remainder(v) in place of remainder. This completes the description of the examination of a subpath. When the traversal returns to the root of  $T_c$  all subpaths have been examined. If  $numcut(root) \geq k$ , then  $\lambda$  is feasible  $(\lambda \leq \lambda^*)$ , else  $\lambda$  is not feasible  $(\lambda > \lambda^*)$ . This completes the description of FTEST2. Note that when all subpaths contain active values, then FTEST2 reduces in general to the straightforward linear-time feasibility test.

Lemma 4.1. Let T be a tree of n vertices and at most n/r leaves, for some integer r. Let T be partitioned into at most  $2\lceil n/r \rceil$  subpaths, each of size at most r. Let all but at most  $n/r^2$  subpaths have no active values and have the functions  $ncut(\cdot)$  and  $rmdr(\cdot)$  computed for all vertices on them. Algorithm FTEST2 correctly determines the feasibility of a test value in  $O((n/r)\log r)$  time.

Proof. Since there are at most n/r leaves, there is a partition into at most  $2\lceil n/r\rceil$  subpaths. As in the proof of Lemma 3.2, the examination of a subpath  $P_j$  with no active values will use  $O(\log r)$  time. This totals  $O((n/r)\log r)$  time for the  $\Theta(n/r)$  such paths. If there are active values for  $P_j$ , then examining the path will take O(r) time. For all of the  $O(n/r^2)$  such subpaths this totals O(n/r) time.  $\square$ 

Theorem 4.2. Algorithm *TREE1* solves the max-min k-partitioning problem on a tree of n vertices in  $O(n(\log \log n)^2)$  time.

Proof. We first consider the generation of tree T'. Each call to MSEARCH

will take O(n') time, produce  $O(\log \log n)$  search values. Thus the total time to produce search values over all iterations will be O(n). It will cost O(n) time to test each search value. Let p be the number of leaf-paths before MSEARCH is called to prune the tree. Since  $p \geq 2n/r$ , the number of leaf-paths of length at most r is at most p/2. The number of active values at the termination of MSEARCH is at most  $n'/(2r) \leq n/(2r) \leq p/4$ . Thus there are at least p/2 - p/4 = p/4 leaf-paths with no active values on them. Since the removal of as few as two such paths can cause an interior vertex to become a leaf, the tree resulting at the end of an iteration will have at most 7/8 of the leaves of the tree at the beginning of the iteration. The number of iterations needed to reduce the tree to one with at most 2n/r leaves will be  $O(\log r) = O(\log \log n)$ . Thus the total time to generate T' will be  $O(n(\log \log n)^2)$ .

Once T has been reset to T',  $T_c$  can be generated in O(n) time. By Lemma 4.1, any subsequent feasibility test will use  $O((n/r)\log r) = O(n\log\log n/(\log n)^2)$  time. Each vertex in  $T_c$  will participate in the formation of a sorted matrix for just one call to MSEARCH. Thus the total time to produce test values in all calls to MSEARCH during the second phase is O(n), and each call will produce  $O(\log n)$  test values. Since the number of leaves is at least halved on each iteration of the while loop, the number of calls to MSEARCH is  $O(\log n)$ . Thus the number of values tested is at most  $O((\log n)^2)$ . Since these can each be tested in  $O(n\log\log n/(\log n)^2)$  time, the total time for feasibility testing in the second phase is  $O(n\log\log n)$ .  $\square$ 

We next apply the above strategy recursively to yield an  $O(n \log^* n)$ -time algorithm TREE2. The idea is to start with the tree and a straightforward feasibility test, and bootstrap the computation by alternately pruning the tree to reduce the number of leaves (and hence the number of paths in an edge-path-partition), and then constructing data structures for an ever faster feasibility tests. Initially all values are active, so that functions  $ncut(\cdot)$  and  $rmdr(\cdot)$  are uninitialized. We call TREE2 when

```
proc TREE2 (tree T, integer k)

Let f(n) be the largest power of 2 no larger than (\log n)^3.

Let lev be the smallest nonnegative integer such that f^{lev}(n) = 2^9.

\lambda_1 \leftarrow 0; \ \lambda_2 \leftarrow \infty

T' \leftarrow T; \ k' \leftarrow k; \ T_c \leftarrow T

for each value of i from lev down to 0 do

Perform phase 1 of TREE1, replacing \lfloor (\log n)^2 \rfloor by f^i(n),

r by r_i, \mathcal{L} by \mathcal{L}_i, n' by n_i, and \mathcal{M} by \mathcal{M}_i.

endfor

\lambda^{\bullet} \leftarrow \lambda_1
endproc
```

Theorem 4.3. Algorithm TREE2 solves the max-min k-partitioning problem on a path of n vertices in  $O(n \log^{n} n)$  time.

**Proof.** For correctness, note that when i = 0,  $r_i = n$  and the while-loop (taken from *TREE1*) will not terminate until T' is a simple path. Then the stopping count on MSEARCH in  $\mathcal{M}_0$  will be 1/n, so that no active values will remain at the end of phase 0.

We analyze the time as follows. There will be lev+1 phases, where lev is  $O(\log^* n)$ . One call to MSEARCH with matrices of size at most  $r_i \times r_i$  will use O(n') time and generate  $O(\log r_i)$  test values, so that all such calls in phase i will use O(n) time and produce  $O((\log r_i)^2)$  test values. The time to compute  $ncut(\cdot)$  and  $rmdr(\cdot)$  values using DIGEST1 at the end of the phase will also be O(n). All other activities will use O(n) time over all phases. In phase lev, performing all feasibility tests will use O(n) time. Performing all feasibility tests on any other phase i will use  $O((n/r_{i+1})\log r_{i+1}(\log r_i)^2)$  time. Since  $r_{i+1}$  is  $O((\log r_i)^3)$ , this time is O(n) in total. The time to recompute  $ncut(\cdot)$  and  $rmdr(\cdot)$  values for vertices whose weight has changed is  $O((n/r_{i+1})\log r_{i+1})$ , since  $O(n/r_{i+1})$  nodes will have their weights change, and each recomputation involves a binary search on a subpath of length at most  $r_{i+1}$ .

We now discuss algorithm TREE3, which runs in linear time. In TREE2, O(n) time per phase is spent on each of three activities: computing  $ncut(\cdot)$  and  $rmdr(\cdot)$  values, producing search values, and feasibility testing. We shall modify TREE2 to use O(n) time in total on these activities. In place of DIGEST1 use DIGEST2. Since we do not know which paths will be removed, we lay out the tree in a certain way and then copy paths when it is determined that they should be "glued" together. As in PATH3, we call MSEARCH on sets of trimmed submatrices to produce our search values more efficiently. Constituting these matrices is trickier, because we do not know ahead of time by how much certain vertices will have their weight increased. This will force us to represent the paths in a special way.

As in TREE2, we will modify the current copy T' of the tree as we do feasibility testing on an old copy T of the tree. In TREE2 we copy T' into T at the end of a phase, but this straightforward copying is too expensive for TREE3. Instead, we maintain two complete sets of data structures, one for T and one for T'. We make changes to the structures for T' as we proceed on a phase, and we make the identical changes to T at the end of the phase.

As TREE3 progresses, the tree is pruned, and some of the edges in the edgepath-partition are deleted, while others are concatenated together. We rely in several places on a path in a vertex-path-partition being represented in consecutive locations of an array. To facilitate these activities, store the weights of the vertices of the tree initially in an array in the following order. Order the children of each vertex from left to right by nondecreasing height in the tree. Then list the vertices in postorder. This takes O(n) time to set up, since the heights of all vertices can be found in O(n)time, and then pairs containing parent and height can be ordered lexicographically in O(n) time. Arrays that are parallel to the above weight array will be used to store ncut, rmdr, and next values, as well as additional information used to compute the weight of specified subpaths. We discuss this additional information in due course.

When paths are removed from the tree, the storage of the tree within the array is then reorganized as follows. Let P be a leaf-path in the edge-path-partition of the current tree, and let t be the top vertex of P. We shall remove P-t. Let P' be the path whose bottom vertex is t. If t will have only one child remaining after removal of P-t, and this child was originally not the rightmost child of t, do the following. Let P'' be the other path whose top vertex is t. Copy the vertices of P''-t in order so that the top vertex of this path is in the location preceding the location of t in the array. Modify the weight of t by adding the remainder left from removing P-t. Also, ncut, rmdr and next values for all copied vertices should also be copied into the new locations in their arrays. The additional information should be copied and modified, as we discuss shortly.

Note that the bottom vertex of P'' may have been the parent of several vertices. When P'' - t is moved, the children of its bottom vertex (and subtrees rooted at them) are not copied. It is simple to store in a location formerly assigned to the bottom vertex of P'' the current location of this vertex, and to also store a pointer back. If the path containing P'' is copied, then this pointer can be reset easily. When only one child of the bottom vertex of P'' remains, the corresponding path can be copied to in front of P''. We claim that the total time to perform all rearrangements will be O(n). The time to copy each vertex and copy and adjust its accompanying information is constant. Because of the way in which the tree is stored in the array, at most one vertex will be copied from any array location.

We now discuss the additional information that needs to be maintained so that we can compute the weight of any specified subpath quickly. A first attempt might be to initialize the value  $A_v$  for every vertex v to be the sum of the weights of all descendants

of v (including itself). Note that for any path in the edge-path-partition, if w is an ancestor of v, w is not the top of the path, and v is not the bottom of the path, then  $A_w - A_{v-1}$  will be the sum of all vertex weights from v up to w. Including the bottom or top vertex in the specified subpath involves adding the weight of whichever or both of these vertices into a difference of the same form. When the weight of a vertex t is changed, the value  $A_t$  can be changed, and the  $A_v$  values for each vertex v in a copied path can be changed. The time for changing values can be charged to that of copying the path. However, path P'' - t should not be copied if the top vertex of P'' - t was originally the rightmost child of t. In this case there is no copying to which the cost of changing  $A_v$  values could be charged. We next describe a mechanism that overcomes this problem.

Let a path in the edge-path-partition be called *light* if its weight is less than  $\lambda_2$ , and heavy otherwise. We handle light and heavy paths differently. For a light path, we keep track of the total weight in the path. For light paths, if the total weight of vertices in P is at most  $\lambda_1$ , then no value induced by P is in contention for being  $\lambda^*$ . However, there may be many light paths whose weights are greater than  $\lambda_1$ , and thus their weights, and the weights of certain of their subpaths, could be candidates for being  $\lambda^*$ . On phase i, there could be as many as  $O(n/r_i)$  after the tree has been pruned. We reduce this number as follows. Form the set of weights of such paths. Then run MSEARCH on this set with stopping count  $n/r_i^2$ . This will reduce the number of light paths of weight greater than  $\lambda_1$  to  $O(n/r_i^2)$ . These are then not included in the call to MSEARCH that allows better data structures to be constructed on phase i. Thus we tolerate  $O(n/r_i^2)$  light paths that will continue to have active values.

We next discuss the representation of a heavy path. Each heavy path P is represented as a sequence of overlapping subpaths, each of weight at least  $\lambda_2$ . Each vertex

of P is in at most two overlapping subpaths. Each overlapping subpath, except the first, overlaps the previous overlapping subpath on vertices of total weight at least  $\lambda_2$ , and each overlapping subpath, except the last, overlaps the following overlapping subpath on vertices of total weight at least  $\lambda_2$ . Thus any sequence of vertices of weight at most  $\lambda_2$  that is contained in the path is contained in one of its overlapping subpaths. For each overlapping subpath, we shall maintain  $A_v$  values for every vertex v except the top vertex of the overlapping subpath. A vertex v that is in two overlapping subpaths is allowed to have different  $A_v$  values for each overlapping subpath (though the  $A_v$  values within any given overlapping subpath are consistent).

Initially no path is heavy, since initially  $\lambda_2 = \infty$ . A heavy path P can be created in one of two ways.

- 1. Path P was light until the value of  $\lambda_2$  was reduced by MSEARCH.
- Path P is formed by the concatenation of two or more paths.

If a heavy path P is created in the first way, then represent P by two overlapping subpaths that are both copies of P, designating one as the first overlapping subpath and the other as the second. Set the  $A_v$  values of both of the overlapping subpaths so that  $A_w - A_{v-1}$  is the correct value for any v and w, where w is not the top vertex, v is not the bottom vertex, and v is a descendant of w. If P is the concatenation of paths, all of which were light, then do the same thing.

Otherwise, path P is formed by the concatenation of paths, with at least one of them being heavy. Do the following to generate the representation for P. While there is a light path P' to be concatenated to a heavy path P'', combine the two as follows. If P' precedes P'', then extend the first overlapping subpath of P'' to incorporate P', adjusting the  $A_{\nu}$  values for the vertices of P'. If P' follows P'', then extend the last overlapping subpath of P'' and adjust the  $A_{\nu}$  values for the vertices of P'. This

completes the description of the concatenation of light with heavy paths. While there is more than one heavy path, concatenate adjacent heavy paths P' and P'' as follows. Assume that P' precedes P''. Combine the last overlapping subpath of P' with the first of P'', changing all the  $A_{\nu}$  values of the first subpath of P''. Note that a vertex can have its  $A_{\nu}$  value changed at most twice before it is in overlapping subpaths that are neither the first nor the last.

We use three arrays to hold the  $A_{v}$  values of the overlapping subpaths. When the last overlapping subpath of P' is combined with the first overlapping subpath of P'', either both are already in the same array, or there is an array that is empty in the corresponding positions of both subpaths, or the array containing one of them can accommodate the other. To keep track of which overlapping subpaths vertices are in, give each overlapping subpath a unique index, and maintain three additional arrays. For each vertex, the corresponding location in the additional array will contain the index of the overlapping subpath.

As an example, consider the portion of the tree shown in Fig. 4.3. The weights of the vertices are listed in postorder in the first array. Suppose that TREE3 has been running for a while, so that currently  $\lambda_1 = 4$  and  $\lambda_2 = 16$ . Of the five paths in the edge-path-partition that are shown in the figure, all but the path consisting of vertices with weights 13,1,1 are heavy, and are thus represented in the remaining 2 pairs of arrays. Path 14,3,4 is represented by the two overlapping subpaths of index 1 and 2, path 4,8,7 is represented by the two overlapping subpaths of index 3 and 4, path 4,11,1 is represented by the two overlapping subpaths of index 5 and 6, and path 1,10,6 is represented by the two overlapping subpaths of index 7 and 8. In each case we use the overlapping subpath of smaller index as the first of the two overlapping subpaths. Values of the form  $A_{\nu}$  are set for each vertex in an overlapping subpath except the top vertex, and are set for convenience to be the sum of the weights of all

vertices in the subtree rooted at v. Any value in the arrays that is unknown because we have not provided a complete example is marked with an asterisk, and any array location whose value need not yet be set is marked with a dash. We have not shown the third pair of arrays, because they are not yet needed.

We continue the example by supposing that  $\lambda=14$  is tested for feasibility. Suppose that  $\lambda$  is not feasible, so that  $\lambda_2$  is set to 14. The the path 13,1,1 becomes a heavy path, and is represented by overlapping subpaths of index 9 and 10. Then suppose that  $\lambda=7$  is tested for feasibility. Suppose that  $\lambda$  is feasible, so that  $\lambda_1$  is set to 7. Then a cut can be inferred in the leaf-path 14,3,4, between 14 and 3. The two edges of the leaf-path can be removed, and the remaining weight of 3 can be added to the weight of the top vertex, giving weight 7. Then paths 7,8,7 and 7,11,1 are concatenated together, by combining overlapping subpaths 4 and 5 into overlapping subpath 11. Note that overlapping subpath 11 must be formed in the third pair of arrays. The resulting path has overlapping subpaths 3, 11, and 6. The state of the tree and the arrays at this point is shown in Fig. 4.4.

We continue the example further by testing  $\lambda = 13$  for feasibility. Suppose that  $\lambda$  is not feasible, so that  $\lambda_2$  is set to 13. Then a cut can be inferred in the leaf-path 13,1,1, between 13 and 1. The two edges of the leaf-path can be removed, and the remaining weight of 1 can be added to the weight of the top vertex, giving weight 2. Then paths 7,8,7,11,2 and 2,10,6 are concatenated together, by combining overlapping subpaths 6 and 7 into overlapping subpath 12. Note that overlapping subpath 12 must be put in the first pair of arrays, extending forward from overlapping subpath 7. The resulting path has overlapping subpaths 3, 11, 12, and 8. The state of the tree and the arrays at this point is shown in Fig. 4.5.

To compute the weight of vertices from v up to w, where v and  $w \neq v$  are in the same heavy path, do the following. Index into the array structure to find the at most

two overlapping subpaths containing v and the at most two overlapping subpaths containing w-1. Then test if one of the overlapping subpaths containing v also contains w. If so, compute the desired value as follows. If w is not the top vertex of this overlapping subpath, and v is not the bottom vertex, then compute  $A_w - A_{v-1}$ . If w is the top vertex and v is the bottom vertex, then add the weights of w and v to  $A_{w-1} - A_v$ . The other two cases are handled similarly. If no overlapping subpath contains both v and w, then the weight from v to w is too large, and it is sufficient to use  $\lambda_2$  as the resulting value.

Given such a representation of a path, we discuss the formation of sorted matrices upon which to search. For each overlapping subpath, a sorted matrix will be induced. Note that since each vertex is in at most two overlapping subpaths, the total dimensions of these matrices will be only twice the dimensions of a single matrix representing the path.

We next discuss how to perform efficient feasibility testing, given our representation of paths. In a feasibility test, we would like to search any path P in time proportional to the logarithm of its length. This can be done if we maintain a second representation of the path as a red-black balanced search tree [T]. In this tree there will be a node x for every vertex u in P. Node x will contain two additional fields, wt(x) and ct(x). Field wt(x) will contain the sum of weights of all vertices whose nodes are in the subtree rooted at x, and field ct(x) will equal the number of nodes in the subtree rooted at x. With this tree it is easy to search for a vertex v in P such that v is the first vertex in P such that the sum of the weights to v is at least a certain value, and to determine as the same time the position of v in P. The search will take time proportional to the logarithm of the length of P. When two paths P' and P'' need to be concatenated together, the corresponding search trees can be merged in time proportional to the logarithm of the combined path length.

We now summarize our algorithm TREE3, which we call only when  $n \geq 2^9$ .

```
proc TREE3 (tree T, integer k)
Let f(n) be the largest power of 2 no larger than (\log n)^3.
Let lev be the smallest nonnegative integer such that f^{lev}(n) = 2^9.
\mathcal{M}_{lev} \leftarrow \emptyset
\mathcal{A}_{lev} \leftarrow \emptyset
\lambda_1 \leftarrow 0; \ \lambda_2 \leftarrow \infty
T' \leftarrow T; k' \leftarrow k; T_c \leftarrow T
for each value of i from lev down to 0 do
     r_i \leftarrow f^i(n)
      Form an edge-path-partition of T'.
      Perform the while-loop of TREE2 to prune T'. (Refer to TREE1.)
      Update the representations of the paths of T' that changed.
      Call MSEARCH on the set of weights of light paths,
           with stopping count n/r_i^2, using FTEST2 for T_c and k.
      Augment \mathcal{M}_i to reflect all changes to the set of overlapping subpaths.
      Perform the body of the for-loop for a phase from PATH3,
           except that rather than FTEST1, FTEST2 is used (for T_c and k),
           and DIGEST2, rather than DIGEST1, is applied to subpaths in T'.
      Update T to reflect the changes to T', reset k to be k', and update T_c for T.
endfor
\lambda^* \leftarrow \lambda_1
endproc
```

We start phase i with a set  $\mathcal{M}_i$  of  $O(n/r_{i+1})$  submatrices of matrices representing all paths in T, except for  $O(n/r_{i+1}^2)$  light paths, and a set  $\mathcal{A}_i$  of  $O(n/r_{i+1}^2)$  elements. The value  $\lambda^*$  is either  $\lambda_1$ , or is one of the elements of  $\mathcal{A}_i$ , or is in one of the submatrices of  $\mathcal{M}_i$ , or is a sum in one of the light paths.

Theorem 4.4. Algorithm TREE3 solves the max-min k-partitioning problem on a tree of n vertices in O(n) time.

Proof. We analyze the time for TREE3 as follows. Phase i starts with a set  $\mathcal{M}_i$  of  $(n/r_{i+1})$  matrices representing all paths, except for  $(n/r_{i+1}^2)$  light paths, and a set  $\mathcal{A}_i$  of  $(n/r_{i+1}^2)$  elements. Pruning the tree will use O(n') time to prune away  $\Theta(n')$  vertices. This will be a total of O(n) over all phases. As discussed earlier, copying

paths in the array that represents the tree will be accounted as a constant charge for each array location, which will be a total of O(n) over all phases. Concatenating two paths by merging their balanced tree representations will cost time proportional to the logarithm of the length of the longer path. Since there will be  $O(n/r_{i+1})$  concatenations, the worst case time for these will be  $O((n/r_{i+1})\log\log r_{i+1})$  on phase i. Modifying the path tree  $T_c$  and identifying paths will take  $O(n/r_{i+1})$  time. Calling MSEARCH on the set of weights of light paths will take  $O(n/r_i)$  time to produce  $O(\log r_i)$  search values. The total time to test these will be  $O((n/r_{i+1})\log r_i\log r_{i+1})$ , which is  $O((n/r_i)\log\log r_i)$ . Augmenting  $\mathcal{P}_i$  and  $\mathcal{M}_i$  will cost constant time for each vertex appearing for the first or second time in such a path. The remaining operations are analyzed as in the proof of Theorem 3.6. Thus the total time for TREE3 over all phases is O(n).  $\square$ 

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## References

- [AP] E. M. Arkin and C. H. Papadimitriou. On the complexity of circulations. J. Algorithms, 7:134-145, 1986.
- [BP] R. I. Becker and Y. Perl. Shifting algorithms tree partitioning with general weighting functions. J. Algorithms, 4:101-120, 1983.
- [BPS] R. I. Becker, Y. Perl, and S. R. Schach. A shifting algorithm for min-max tree partitioning. J. ACM, 29:58-67, 1982.
- [BFPRT] M. Blum, R. W. Floyd, V. R. Pratt, R. L. Rivest, and R. E. Tarjan. Time bounds for selection. J. Comput. Syst. Sci., 7:448-461, 1972.
- [CT] R. Chandrasekaran and A. Tamir. Polynomially bounded algorithms for locating p-centers on a tree. Math. Prog., 22:304-315, 1982.

- [CEIY] M. Chrobak, D. Eppstein, G. F. Italiano, and M. Yung. Efficient sequential and parallel algorithms for computing recovery points in trees and paths. In Proceedings of the 2nd ACM-SIAM Symposium on Discrete Algorithms, pages 158-167, 1991.
- [CK] M. J. Chung and M. S. Krishnamoorthy. Algorithms for placing recovery points. Inf. Proc. Lett., 28:177-181, 1988.
- [C1] R. Cole. Partitioning point sets in arbitrary dimensions. Theor. Comput. Sci., 49:239-265, 1987.
- [C2] R. Cole. Slowing down sorting networks to obtain faster sorting algorithms. J. ACM, 34:200-208, 1987.
- [CSSS] R. Cole, J. S. Salowe, W. L. Steiger, and E. Szemeredi. An optimal-time algorithm for slope selection. SIAM J. Comput., 18:792-810, 1989.
- [F1] G. N. Frederickson. Optimal algorithms for tree partitioning. In Proc. 2nd ACM-SIAM Symp. on Discrete Algorithms, pages 168-177, San Francisco, January 1991.
- [F2] G. N. Frederickson. Parametric search and locating supply centers in trees. In Proc. 2nd Workshop on Algorithms and Data Structures, pages 299-319, Ottawa, Canada, August 1991. Springer-Verlag, LNCS 519.
- [F3] G. N. Frederickson. Optimal parametric search algorithms in trees II: p-center problems and checkpointing. 1992.
- [FJ1] G. N. Frederickson and D. B. Johnson. Finding kth paths and p-centers by generating and searching good data structures. J. Algorithms, 4:61-80, 1983.
- [FJ2] G. N. Frederickson and D. B. Johnson. Generalized selection and ranking: sorted matrices. SIAM J. on Computing, 13:14~30, 1984.
- [Go1] A. J. Goldman. Optimal center location in simple networks. *Transport.* Sci., 5:212-233, 1971.
- [Go2] A. J. Goldman. Minimax location of a facility in an undirected tree graph. Transport. Sci., 6:407-418, 1972.

- [Gu] D. Gusfield. Parametric combinatorial computing and a problem of program module distribution. J.ACM, 30:551-563, 1983.
- [Hk] S. L. Hakimi. Optimum locations of switching centers and the absolute centers and medians of a graph. Operations Res., 12:450-459, 1964.
- [Hn1] G. Y. Handler. Minimax location of a facility in an undirected tree graph. Transport. Sci., 7:287-293, 1973.
- [Hn2] G. Y. Handler. Finding two-centers of a tree: the continuous case. Transport. Sci., 12:93-106, 1978.
- [KH] O. Kariv and S. L. Hakimi. An algorithmic approach to network location problems. SIAM J. Appl. Math., 37:513-538, 1979.
- [KM] S. Kundu and J. Misra. A linear tree partitioning algorithm. SIAM J. on Computing, 6:151-154, 1977.
- [M] N. Megiddo. Applying parallel computation algorithms in the design of serial algorithms. J. ACM, 30:852-865, 1983.
- [MTZC] N. Megiddo, A. Tamir, E. Zemel, and R. Chandrasekaran. An  $O(n \log^2 n)$  algorithm for the kth longest path in a tree with applications to location problems. SIAM J. Comput., 10:328-337, 1981.
- [PS] Y. Perl and S. R. Schach. Max-min tree partitioning. J. ACM, 28:5-15, 1981.
- [PV] Y. Perl and U. Vishkin. Efficient implementation of a shifting algorithm. Disc. Appl. Math., 12:71-80, 1985.
- [R] M. Reichling. On the detection of a common intersection of k-convex polyhedra. Lect. Notes Comput. Sci., 333:180-187, 1988.
- [S] J. S. Salowe. L-infinity interdistance selection by parametric search. Inf. Proc. Lett., 30:9-14, 1989.
- [T] R. E. Tarjan. Updating a balanced search tree in O(1) rotations. Info. Proc. Lett., 16:253-257, 1983.

[Z] E. Zemel. A linear time randomizing algorithm for searching ranking functions. Algorithmica, 2:81-90, 1987.

	6	11	9	2	1	15	7	8
P	0-	<del>-</del> 0-	0-	<del>-</del> o-	<del>-</del> 0-	<del>-</del> 0-	<del>-</del> 0	<del></del> 0

		$X_2$									
		0	6	17	26	28	29	44	51		
	59	59	53	42	33	31	30	15	8		
$X_1$	51	51	45	34	25	23	22	7	0		
	44	44	38	27	18	16	15	0	0		
	29	29	23	12	3	1	0	0	0		
	28	29 28	22	11	2	0	0	0	0		
	26	26	20	9	0	0	0		0		
	17	17 6	11	0	0	0	0	0	0		
	6	6	0	0	0	0	0	0	0		
		M(P)									

Figure 2.1 Path P, vectors  $X_1$  and  $X_2$ , and matrix M(P).

	59	53	42	33	31	30	15	8
1 1 1 1	51	45	34	25	23	22	7	0
1 1 1	44	38	27	18	16	15	0	0
1 1	29	23	12	3	1	0	0	0
1 1	28	22	11	2				
1	26	20	9	0	 			
	17	11	0	0	i ! ! !			
1	6	0	0	0	1 			

Figure 2.2(a) Cells after quartering the second iteration of the outer repeat loop.

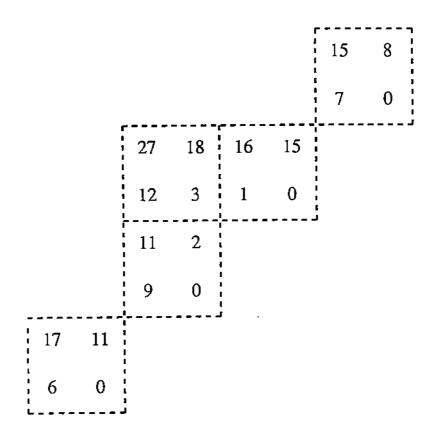


Figure 2.2(b) Cells at the end of the second iteration of the outer repeat loop.

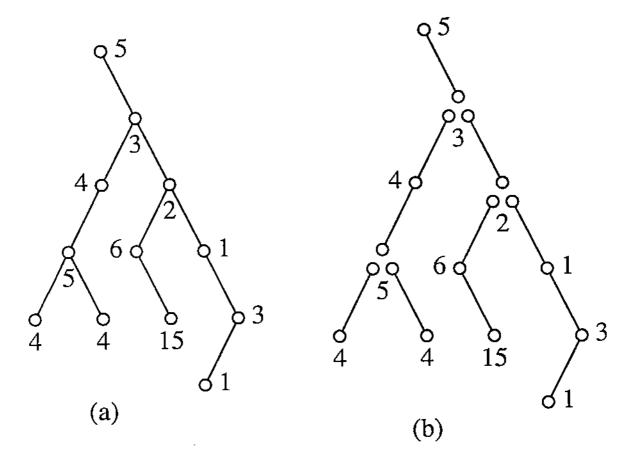


Figure 2.3 (a) A vertex-weighted tree T. (b) The edge-path-partition for T.

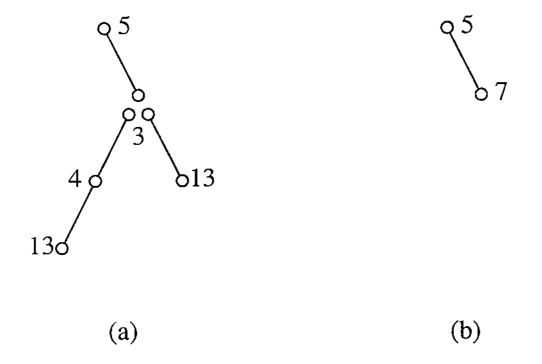


Figure 2.4 (a) Partition after leaf-paths deleted. (b) Partition after further deletion.

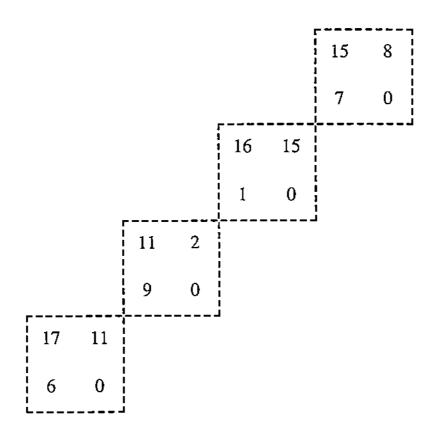


Figure 3.1 Set  $M_1$  of submatrices for the subpaths.

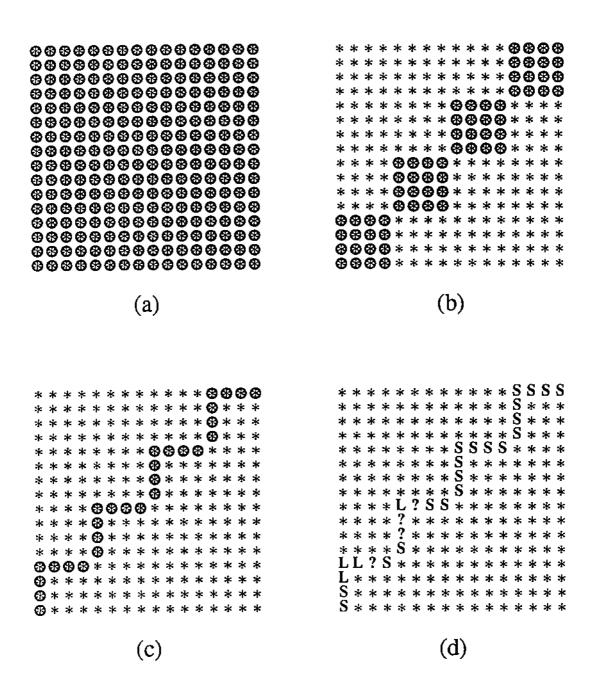


Figure 3.2 (a) Matrix M(P) in bold. (b) Submatrices  $M_{lev}^{P}$  for the subpaths. (c) Thin matrices  $T_{lev}$ . (d) Characterization of elements in thin matrices.

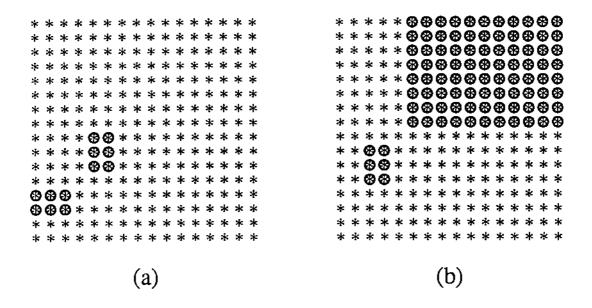


Figure 3.3 (a) Trimmed submatrices  $\mathbf{M}_{lev}^{\mathbf{T}}$ . (b) Submatrices  $\mathbf{M}_{lev-1}$ .

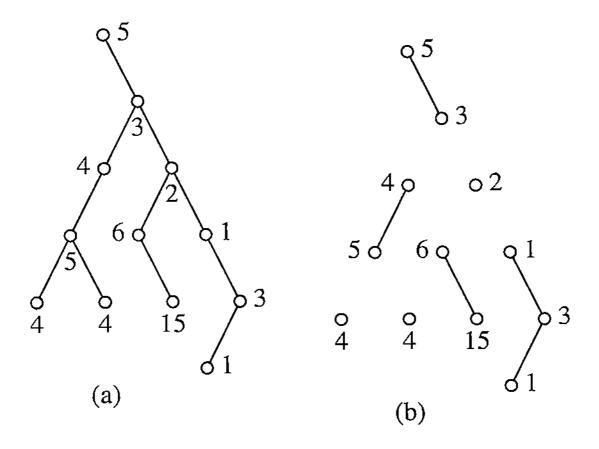


Figure 4.1 (a) A vertex-weighted tree T. (b) The vertex-path-partition for T.

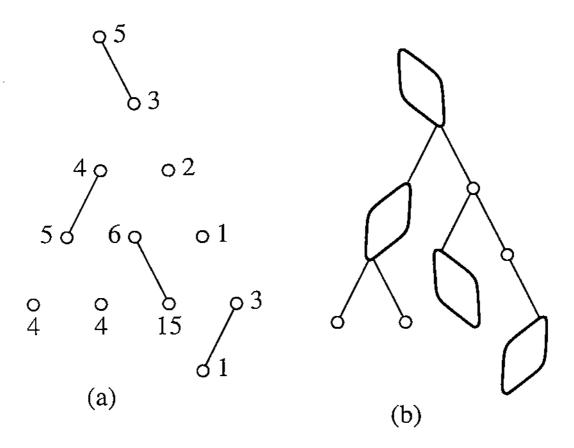


Figure 4.2 (a) The 2-partitions of the vertex-path-partition for T. (b) The path-tree  $T_c$  for T with r=2.

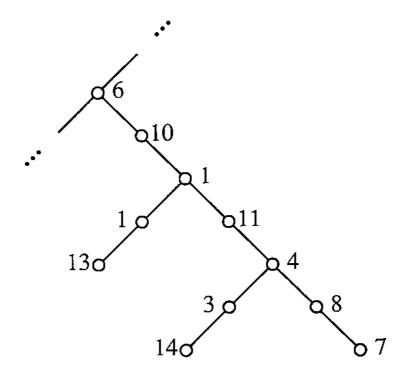
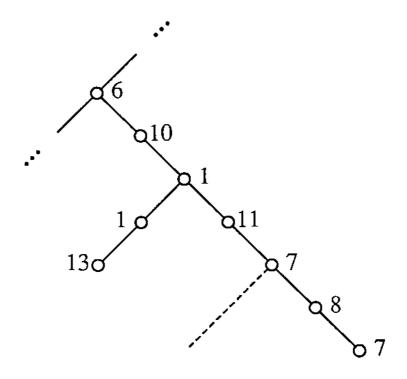


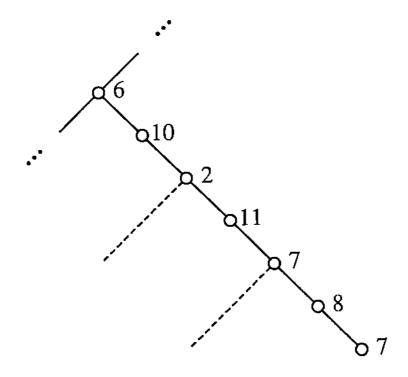


Figure 4.3 A portion of a tree and some of its associated arrays.



 13	1	-	-	7	8	7	11	1	10	6	• • •
 9	9	-	-	3	3	-	_	7	7	*	
 13	14	-	-	7	15	-	-	62	72	*	
 10	10	-	-	-	-	6	6	8	8	*	
 13	14	-	-	-	-	36	47	62	72	*	
 -	-	-	-	11	11	11	11	-	-	*	• • •
 _	_	_	_	7	15	22	33	_	_	*	

Figure 4.4 A portion of the tree later and some of its associated arrays.



• • •	-	-	-	-	7	8	7	11	2	10	6	• • •
	-	_	-	-	3	3	12	12	12	12	*	
	-	-	-	-	7	15	36	47	49	59	*	• • •
	-	-	-	-	-	-	-	-	8	8	*	
	-	_	-	-	-	-	-	-	62	72	*	
	-	-	-	-	11	11	11	11	-	-	*	
	_	_	_	_	7	15	22	33	_	-	*	

Figure 4.5 A portion of a tree yet later and some of its associated arrays.