

## A TIGHT AMORTIZED BOUND FOR PATH REVERSAL

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Path reversal is a form of path compression used in a disjoint set union algorithm and a mutual exclusion algorithm. We derive a tight upper bound on the amortized cost of path reversal.

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Let  $T$  be a rooted tree. A *path reversal* at a node  $x$  in  $T$  is performed by traversing the path from  $x$  to the tree root  $r$  and making  $x$  the parent of each node on the path other than  $x$ . Thus  $x$  becomes the new tree root. (See Fig. 1). The *cost* of the reversal is the number of edges on the path reversed. Path reversal is a variant of the standard path compression algorithm for maintaining disjoint sets under union [5]. It has also been used in a novel mutual exclusion algorithm [2,6].

Suppose that a sequence of  $m$  reversals is performed on an arbitrary initial  $n$ -node tree. What is the total cost of the sequence? Let  $T(n, m)$  be the

worst-case cost of such a sequence, and let  $A(n, m) = T(n, m)/m$ . We are most interested in the value of  $A(n, m)$  for fixed  $n$  as  $m$  grows. As discussed by Tarjan and Van Leeuwen [5], binomial trees provide a class of examples showing that  $A(n, m) \geq \lfloor \log n \rfloor$ <sup>1</sup>, and their rather com-

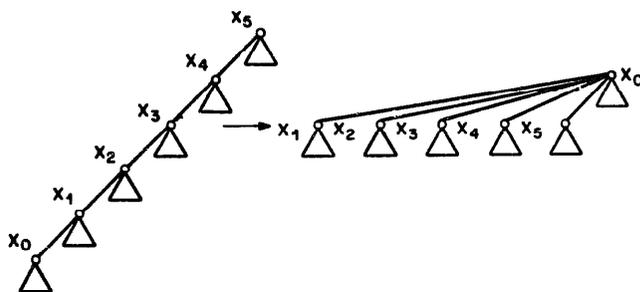


Fig. 1. Path reversal (triangles denote subtrees).

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<sup>1</sup> All logarithms in this paper are base 2.

plicated and their rather complicated analysis gives an upper bound of

$$A(n, m) = O\left(\log n + \frac{n \log n}{m}\right).$$

Ginat and Shankar [2] prove that

$$A(n, m) \leq 2 \log n + \frac{n \log n}{m}.$$

We shall prove that

$$A(n, m) \leq \log n + \frac{n \log n}{2m}.$$

In the special case that the initial tree consists of a root with  $n - 1$  children, which is the case in the mutual exclusion algorithm, the bound is

$$A(n, m) \leq \log n.$$

To obtain the bound, we apply the *potential function* method of amortized analysis (see [4]). Let the *size*  $s(x)$  of a node  $x$  in  $T$  be the number of descendants of  $x$ , including  $x$  itself. Let the *potential* of  $T$  be

$$\Phi(T) = \frac{1}{2} \sum_{x \in T} \log s(x).$$

Define the *amortized cost* of a path reversal over a path of  $k$  edges to be  $k - \Phi(T) + \Phi(T')$ , where  $T$  and  $T'$  are the trees before and after the reversal, respectively. For any sequence of  $m$  reversals, we have

$$\sum_{i=1}^m a_i = \sum_{i=1}^m (t_i - \Phi_{i-1} + \Phi_i) = \sum_{i=1}^m t_i - \Phi_0 + \Phi_m,$$

where  $a_i$ ,  $t_i$ , and  $\Phi_i$  are the amortized cost of the  $i$ th reversal, the actual cost of the  $i$ th reversal, and the potential after the  $i$ th reversal respectively, and  $\Phi_0$  is the potential of the initial tree. Since  $\Phi_0 \leq \frac{1}{2}n \log n$  and  $\Phi_m \geq \frac{1}{2} \log n$ , this inequality yields

$$\sum_{i=1}^m t_i \leq \sum_{i=1}^m a_i + \frac{1}{2}(n - 1) \log n,$$

which in turn implies

$$A(n, m) \leq \frac{1}{m} \sum_{i=1}^m a_i + \frac{n \log n}{2m}.$$

We shall prove that the amortized cost of any reversal is at most  $\log n$ , thereby showing that

$$A(n, m) \leq \log n + \frac{n \log n}{2m}.$$

When the initial tree consists of a root with  $n - 1$  children, the bound drops to  $A(n, m) \leq \log n$ , since then  $\Phi_0 \leq \Phi_m$ , and the extra additive term drops out.

Let  $x_0, x_1, x_2, \dots, x_k$  be a path that is reversed, and let  $A$  be the amortized cost of the reversal. For  $0 \leq i \leq k$ , let  $s_i$  be the size of  $x_i$  before the reversal. The size of  $x_0$  after the reversal is  $s_k$  and the size of  $x_i$  after the reversal, for  $1 \leq i \leq k$ , is  $s_i - s_{i-1}$ . We can thus write  $A$  as

$$\begin{aligned} A &= k - \sum_{i=0}^k \frac{1}{2} \log s_i + \frac{1}{2} \log s_k \\ &\quad + \sum_{i=1}^k \frac{1}{2} \log(s_i - s_{i-1}) \\ &= k + \frac{1}{2} \sum_{i=0}^{k-1} (\log(s_{i+1} - s_i) - \log s_i) \\ &= k + \frac{1}{2} \sum_{i=0}^{k-1} \log((s_{i+1} - s_i)/s_i). \end{aligned}$$

For  $0 \leq i \leq k - 1$ , let  $\alpha_i = s_{i+1}/s_i$ . Note that  $(s_{i+1} - s_i)/s_i = \alpha_i - 1$ . We have

$$\begin{aligned} A &= k + \frac{1}{2} \sum_{i=0}^{k-1} \log(\alpha_i - 1) \\ &= \sum_{i=0}^{k-1} \left(1 + \frac{1}{2} \log(\alpha_i - 1)\right). \end{aligned}$$

We now make use of the following inequality, which will be verified below: for all  $\alpha > 1$ ,  $1 + \frac{1}{2} \log(\alpha - 1) \leq \log \alpha$ . From this inequality we obtain

$$\begin{aligned} A &\leq \sum_{i=0}^{k-1} \log \alpha_i \\ &= \sum_{i=0}^{k-1} \log(s_{i+1}/s_i) = \sum_{i=0}^{k-1} (\log s_{i+1} - \log s_i) \\ &= \log s_k - \log s_0 \\ &\leq \log n, \end{aligned}$$

since  $s_k = n$  and  $s_0 \geq 1$ .

This completes the amortized analysis. We verify the needed inequality by the following chain of reasoning:

$$\begin{aligned}
 0 &\leq (\alpha - 2)^2 \\
 &\Rightarrow 0 \leq \alpha^2 - 4\alpha + 4 \\
 &\Rightarrow 4(\alpha - 1) \leq \alpha^2 \\
 &\Rightarrow \log(4(\alpha - 1)) \leq \log(\alpha^2) \\
 &\Rightarrow 2 + \log(\alpha - 1) \leq 2 \log \alpha \\
 &\Rightarrow 1 + \frac{1}{2} \log(\alpha - 1) \leq \log \alpha.
 \end{aligned}$$

We conclude some remarks. The definition of the potential function used here has been borrowed from Sleator and Tarjan's analysis of splay trees [3]; it has also been used to analyze pairing heaps [1]. As in the case of splay trees, the upper bound can be generalized in the following way. Assign to each tree node  $x$  a fixed but arbitrary positive weight  $w(x)$ . Define the *total weight* of  $x$ ,  $tw(x)$ , to be the sum of the weights of all descendants of  $x$ , including  $x$  itself. Define the potential of the tree  $T$  to be

$$\Phi(T) = \frac{1}{2} \sum_{x \in T} \log tw(x).$$

A straightforward extension of the above analysis shows that the total cost of a sequence of  $m$  reversals is at most

$$\sum_{i=1}^m \log(W/w_i) + \Phi_0 - \Phi_m,$$

where  $w_i$  is the weight of the node  $x_i$  at which the  $i$ th reversal starts and  $W$  is the sum of all the node weights.

Choosing  $w(x) = 1$  for all  $x \in T$  gives our original result. Choosing  $w(x) = f(x) + 1$ , where  $f(x)$  is the number of times a reversal begins at  $x$ , gives an upper bound for the total time of all reversals of

$$\sum_{i=1}^m \log\left(\frac{n+m}{f(x_i)}\right) + \frac{1}{2} \sum_{x \in T} \log\left(\frac{n+m}{f(x)}\right).$$

It is striking that the "sum of logarithms" potential function serves to analyze three different data structures. We are at a loss to explain this phenomenon; whereas there is a clear connection between splay trees and pairing heaps (see [1]), no such connection between trees with path reversal and the other two data structures is apparent. In the case of path reversal, the sum of logarithms potential function gives a bound that is exact to within an additive term depending only on the initial and final trees. It would be extremely interesting and useful to have a systematic method for deriving appropriate potential functions. The three examples of splaying, pairing, and reversal offer a setting in which to search for such a method.

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