

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 $A(n, m) \geq \lfloor \log n \rfloor$ ¹, and their rather com-

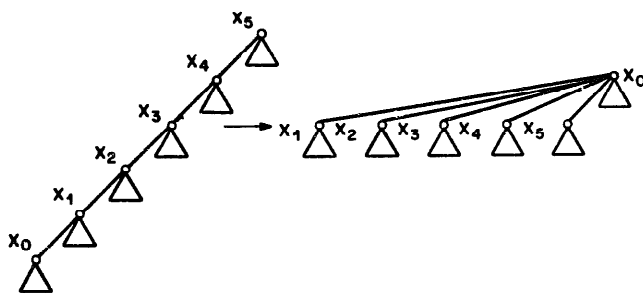


Fig. 1. Path reversal (triangles denote subtrees).

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¹ 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 Φ_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 Φ_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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References

- [1] M.L. Fredman, R. Sedgewick, D.D. Sleator, and R.E. Tarjan, The pairing heap: a new form of self-adjusting heap, *Algorithmica* 1 (1986) 111-129.
- [2] D. Ginat and A. Udaya Shankar, Correctness proof and amortization analysis of a log N distributed mutual exclusion algorithm, Tech. Rept. CS-TR-2038, Department of Computer Science, University of Maryland, 1988.
- [3] D.D. Sleator and R.E. Tarjan, Self-adjusting binary search trees, *J. Assoc. Comput. Mach.* 32 (1985) 652-686.
- [4] R.E. Tarjan, Amortized computational complexity, *SIAM J. Alg. Disc. Meth.* 6 (1985) 306-318.
- [5] R.E. Tarjan and J. van Leeuwen, Worst-case analysis of set union algorithms, *J. Assoc. Comput. Mach.* 31 (1984) 245-281.
- [6] M. Trehel and M. Naimi, A distributed algorithm for mutual exclusion based on data structures and fault tolerance, In: *Proc. Sixth Ann. Internat. Phoenix Conf. on Computers and Communication*, Scottsdale, AZ (February 1987) 35-39.