# **Heaps Simplified**

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**Abstract.** The heap is a basic data structure used in a wide variety of applications, including shortest path and minimum spanning tree algorithms. In this paper we explore the design space of comparison-based, amortized-efficient heap implementations. From a consideration of dynamic single-elimination tournaments, we obtain the binomial queue, a classical heap implementation, in a simple and natural way. We give four equivalent ways of representing heaps arising from tournaments, and we obtain two new variants of binomial queues, a one-tree version and a one-pass version. We extend the one-pass version to support key decrease operations, obtaining the *rankpairing heap*, or *rp-heap*. Rank-pairing heaps combine the performance guarantees of Fibonacci heaps with simplicity approaching that of pairing heaps. Like pairing heaps, rank-pairing heaps consist of trees of arbitrary structure, but these trees are combined by rank, not by list position, and rank changes, but not structural changes, cascade during key decrease operations.

# 1 Introduction

A *meldable heap* (henceforth just a *heap*) is a data structure consisting of a set of items, each with a real-valued key, that supports the following operations:

- make-heap: return a new, empty heap.
- insert(x, H): insert item x, with predefined key, into heap H.
- find-min(H): return an item in heap H of minimum key.
- delete-min(h): delete from heap H an item of minimum key, and return it; return null if the heap is empty.
- $meld(H_1, H_2)$ : return a heap containing all the items in disjoint heaps  $H_1$  and  $H_2$ , destroying  $H_1$  and  $H_2$ .

Some applications of heaps need either or both of the following additional operations:

- decrease-key $(x, \Delta, H)$ : decrease the key of item x in heap H by amount  $\Delta > 0$ , assuming H is the unique heap containing x.
- delete(x, H): delete item x from heap H, assuming H is the unique heap containing x.

We shall assume that all keys are distinct; if they are not, we can break ties using any total order of the items. We allow only binary comparisons of keys, and we study the amortized efficiency [28] of heap operations. To obtain a bound on amortized efficiency, we assign to each configuration of the data structure a non-negative *potential*, initially zero. We define the *amortized time* of an operation to

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be its actual time plus the change in potential it causes. Then for any sequence of operations the sum of the actual times is at most the sum of the amortized times.

Since n numbers can be sorted by doing n insertions into an initially empty heap followed by n minimum deletions, the classical  $\Omega(n \log n)$  lower bound [21, p. 183] on the number of comparisons needed for sorting implies that either insertion or minimum deletion must take  $\Omega(\log n)$  amortized time, where n is the number of items currently in the heap. For simplicity in stating bounds we assume  $n \ge 2$ . We investigate simple data structures such that minimum deletion (or deletion of an arbitrary item if this operation is supported) takes  $O(\log n)$  amortized time, and each of the other supported heap operations takes O(1) amortized time. These bounds match the lower bound. (The logarithmic lower bound can be beaten by using multiway branching [12, 14].)

Many heap implementations have been proposed over the years. See e.g. [19]. We mention only those directly related to our work. The *binomial queue* of Vuillemin [29] supports all the heap operations in  $O(\log n)$  worst-case time per operation. This structure performs quite well in practice [2]. Fredman and Tarjan [11] invented the *Fibonacci heap* specifically to support key decrease operations in O(1) time, which allows efficient implementation of Dijkstra's shortest path algorithm [3, 11], Edmonds' minimum branching algorithm [6, 13], and certain minimum spanning tree algorithms [11, 13]. Fibonacci heaps support deletion of the minimum or of an arbitrary item in  $O(\log n)$  amortized time and the other heap operations in O(1) amortized time. They do not perform well in practice, however [22, 23]. As a result, a variety of alternatives to Fibonacci heaps have been proposed [4, 7, 15, 18, 19, 24], including a self-adjusting structure, the *pairing heap* [10].

Pairing heaps support all the heap operations in  $O(\log n)$  amortized time and were conjectured to support key decrease in O(1) amortized time. Despite empirical evidence supporting the conjecture [16, 22, 26], Fredman [9] showed that it is not true: pairing heaps and related data structures that do not store subtree size information require  $\Omega(\log \log n)$  amortized time per key decrease. Whether pairing heaps meet this bound is open; the best upper bound is  $O(2^{2\sqrt{\lg \lg n}})$  [25]<sup>5</sup>. Very recently El-Masry [8] proposed a more-complicated alternative to pairing heaps that does have an  $O(\log \log n)$ amortized bound per key decrease.

Fredman's result gives a time-space trade-off between the number of bits per node used to store subtree size information and the amortized time per key decrease; at least  $\lg \lg n$  bits per node are needed to obtain O(1) amortized time per key decrease. Fibonacci heaps use  $\lg \lg n + 2$  bits per node; relaxed heaps [4] use only  $\lg \lg n$  bits per node. All the structures so far proposed that achieve O(1) amortized time per key decrease do a cascade of local restructuring operations to keep the underlying trees balanced.

The bounds of Fibonacci heaps can be obtained in the worst case, but only by making the data structure more complicated: run-relaxed heaps [4] and fat heaps [18] achieve these bounds except for melding, which takes  $O(\log n)$  time worst-case; a very complicated structure of Brodal [1] achieves these bounds for all the heap operations.

Our goal is to systematically explore the design space of amortized-efficient heaps and thereby discover the simplest possible data structures. As a warm-up, we begin in Section 2 by showing that binomial queues can be obtained in a natural way by dynamizing balanced single-elimination tournaments. We give four equivalent ways of representing heaps arising from tournaments, and we give two new variants of binomial queues: a one-tree version and a one-pass version.

Our main result is in Section 3, where we extend our one-pass version of binomial queues to support key decrease. Our main insight is that it is not necessary to maintain balanced trees; all that is needed is to keep track of tree sizes, which we do by means of ranks. We call the resulting data

<sup>&</sup>lt;sup>5</sup> We denote by lg the base-two logarithm.

structure a *rank-pairing heap*, or *rp-heap*. The rp-heap achieves the bounds of Fibonacci heaps with simplicity approaching that of pairing heaps. It resembles the lazy variant of pairing heaps [10, p. 125], except that trees are combined by rank, not by list position. In an rp-heap, rank changes can cascade but not structural changes, and the trees in the heap can evolve to have arbitrary structure. We study two types of rp-heaps. One is a little simpler and uses  $\lg \lg n$  bits per node to store ranks, exactly matching Fredman's lower bound, but its analysis is complicated and yields larger constant factors; the other uses  $\lg \lg n + 1$  bits per node and has small constant factors.

In Section 4 we address the question of whether key decrease can be further simplified. We close in Section 5 with open problems.

#### 2 Tournaments as Heaps

Given a set of items with real-valued keys, one can determine the item of minimum key by running a single-elimination tournament on the items. To run such a tournament, repeatedly match pairs of items until only one item remains. To match two items, compare their keys, declare the item of smaller key the winner, and eliminate the loser. Break a tie in keys using a total order of the items.

Such a tournament provides a way to represent a heap that supports simple and efficient implementations of all the heap operations. The winner of the tournament is the item of minimum key. To insert a new item into a non-empty tournament, match it against the winner of the tournament. To meld two non-empty tournaments, match their winners. To delete the minimum from a tournament, delete its winner and run a tournament among the items that lost to the winner.

For the moment we measure efficiency by counting comparisons. Later we shall show that our bounds on comparisons also hold for running time, to within additive and multiplicative constants. Making a heap or finding the minimum takes no comparisons, an insertion or a meld takes one. The only expensive operation is minimum deletion. To make this operation efficient, we restrict matches by using *ranks*. Each item begins with a rank of zero. After a match between items of equal rank, the winner's rank increases by one. Such a match is *fair*. A match between items of unequal rank is *unfair*. After an unfair match the winner's rank does not change. Optionally, if the winner has rank less than that of the loser, its rank can increase to any value no greater than the loser's rank. One rule governs a tournament: match two items of equal rank if possible. We call such a tournament *balanced*. Except for implementation details, this is the entire description of the data structure.

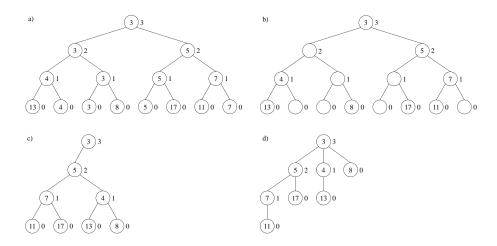
#### **Lemma 1.** A balanced tournament whose winner has rank k contains at least $2^k$ items.

*Proof.* The proof is by induction on the number of matches. The lemma holds initially. An item's rank increases to k only when the item has rank k - 1 and it beats an item of rank k - 1, or it beats an item of rank at least k. In the former case the two matched items are the winners of disjoint balanced tournaments, each containing at least  $2^{k-1}$  items. In the latter case the tournament won by the loser contains at least  $2^k$  items. After the match the combined tournament contains at least  $2^k$  items.  $\Box$ 

In stating bounds we denote by n the number of items in a tournament or heap; we assume n > 1. To bound the amortized number of comparisons per heap operation, we define the potential of a balanced tournament to be the number of unfair matches. An insertion or meld increases the potential by at most one and hence takes at most two amortized comparisons. The tournament rule and Lemma 1 guarantee that at most  $\lg n$  unfair matches occur during a minimum deletion, since an unfair match can only occur when there is at most one item per rank, and by Lemma 1 the number of ranks is at most  $\lg n + 1$ . (The minimum rank is 0; the maximum is  $|\lg n|$ .) To analyze a minimum deletion, let x

be the deleted winner. Consider the sequence of matches won by x. Each such match was either unfair or increased the rank of x. Thus if there were k such unfair matches, the total number of matches won by x was at least k and at most  $k + \lg n$ . Deletion of x eliminates these matches. Finding a new winner takes at most  $k + \lg n$  matches, of which at most  $\lg n$  are unfair. Hence the amortized number of comparisons done by the minimum deletion is at most  $k + \lg n - k + \lg n = 2 \lg n$ .

It remains to implement the data structure. The standard way to represent a tournament is by a full binary tree, whose leaves contain the items and whose internal nodes represent the matches. Each internal node contains the winner of the match and its rank after the match. The internal nodes containing an item form a path of the matches it won. The tree is *heap ordered*: the item in a node has minimum key among the items in the descendants of the node. We call this the *full representation*. (See Figure 1a.)



**Fig. 1.** Four representations of a tournament: a) full, b) half-empty, c) half-ordered, and d) heap-ordered. The number to the right of each node is its rank.

This representation uses almost twice as many nodes as necessary. As a first step in reducing the space, we remove each item from all but the highest node containing it. Our tree is now *half empty*: the root is full, and each parent has two children, one full, the other empty. Each node, whether full or empty, has a rank. This is the *half-empty representation*. (See Figure 1b.)

A half-empty heap-ordered tree represents a tournament transparently, but it uses as many nodes as the full representation. We obtain a more compact representation by eliminating the empty nodes. To do this we define the *ordered* child and the *unordered* child of a full node to be its full child and the full child of its empty sibling, respectively. The tree becomes a *half tree*: the root has an ordered child but no unordered child, every non-root node has both an ordered child and an unordered child, and any child can be missing. We give each child a rank equal to the rank of its parent in the half-empty representation, minus one. (If the parent represents a fair match, this is just the rank of the child itself in the half-empty representation, but it need not be if the parent represents an unfair match. We define the ranks in this way to avoid losing information about the ranks in the half-empty representation if there are unfair matches.) With this definition a leaf has rank zero. We adopt the convention that a missing child has rank -1 and a root has rank one larger than that of its child. (There is no need to explicitly maintain the ranks of roots.) The rank of a half tree is the rank of its root. The *ordered* and *unordered* 

*subtrees* of a parent are the subtrees rooted at its ordered and unordered children, respectively. Each subtree is *half ordered*: the item in a node has smaller key than those of all items in its ordered subtree. The items that lost to an item in a node are those in the nodes on the path starting from the ordered child of the node and descending through unordered children. This is the *half-ordered representation* of a heap. (See Figure 1c.)

In the half-ordered representation we do not need to move items among nodes as matches take place, so the items can *be* the nodes: the data structure can be *endogenous* [27]. Henceforth we shall assume an endogenous representation.

We obtain our fourth and final representation, the *heap-ordered representation*, by viewing a half tree as the binary tree representation of a tree [20, pp. 332-346]: the ordered and unordered children of a node become its first child and next sibling, respectively. The tree is heap-ordered; the children of a node are the items it defeated, most recent match first. (See Figure 1d.)

Many heap structures, including binomial queues [29], Fibonacci heaps [11], pairing heaps [10], and relaxed heaps [4], were originally presented in the heap-ordered representation. In their paper on pairing heaps, Fredman et al. [10] described the half-ordered representation and observed that it is the binary tree representation [20] of a heap-ordered tree. Later, Dutton [5] used the half-ordered representation in his *weak-heap* data structure, and Høyer [15] proposed various kinds of half-ordered balanced trees as heaps. The version of Fredman et al. [10] matches Knuth's definition [20]: the left and right children of a node are its first child and next sibling in the heap-ordered representation, respectively. The version of Dutton and Hoyer reverses left and right. To avoid confusion we have named the children based on their roles. Throughout we denote by ord(x) and unord(x) the ordered and unordered child of x, respectively.

Among the four representations we prefer the half-ordered representation because it saves space and simplifies the implementation of key decrease operations. To implement balanced tournaments in the half-ordered representation, we store with each node its rank and pointers to its ordered and unordered children. To match the roots x and y of two half trees, compare their keys, make the ordered subtree of the winner the unordered subtree of the loser, and make the new half tree rooted at the loser the ordered subtree of the winner. Set the rank of the winner's new child appropriately. (This is the rank of the winner in the full representation, minus one.) (See Figure 2.)

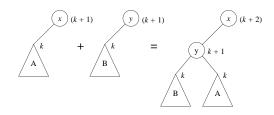


Fig. 2. A fair match between roots x and y of two half-trees. Ranks are to the right of nodes. Ranks of roots (in parentheses) need not be maintained explicitly.

A match of two roots takes O(1) time. Thus insertion and melding take O(1) time. To do a minimum deletion, form the new heap by starting at the child of the root and walking down through unordered children. Detach each node along with its ordered subtree, which together form a half tree. For each rank, keep track of a half tree of that rank. When a new half tree has the same rank as an existing one, match their roots, producing a single half tree of one higher rank. If there is already a half

tree of one higher rank, match its root and the root of the new half tree. Continue until the remaining half tree is the only one of its rank. Repeat this process for each new half tree. Once there is at most one half tree per rank, repeatedly match any two of the remaining roots until only one root remains. This method rebuilds the half tree after a minimum deletion in  $O(k + \log n)$  time, where k is the number of unfair matches won by the deleted winner.

The amortized analysis of the number of comparisons per heap operation extends to the running time and gives amortized time bounds of O(1) for insertion and melding and  $O(\log n)$  for minimum deletion.

A balanced tournament is a one-tree version of a binomial queue. In binomial queues, only fair matches are done, and a heap is represented by a set of half trees rather than by a single half tree. Fair matches guarantee that each half tree is *perfect*: the ordered subtree of the root is a perfect binary tree. During an insertion, meld, or minimum deletion, fair matches are done until there is at most one half tree per rank. Each heap operation takes  $O(\log n)$  time in the worst case. By allowing unfair matches, we are able to represent a heap by a single tree. The cost of this simplicity is a linear worst-case time bound for minimum deletion.

We obtain another version of binomial queues by avoiding unfair matches, doing fair matches only during minimum deletions, and running only a single pass of matches during a minimum deletion. The resulting data structure, which we call the *one-pass binomial queue*, has the same amortized efficiency as the one-tree version. Here are the details.

A one-pass binomial queue consists of a set of heap-ordered perfect half trees and a pointer to the root of minimum key, the *minimum node*. We represent the set of half trees by a singly-linked circular list of the roots, with the minimum node first. To insert an item into a heap, create a new, one-node half tree containing the item, add this half tree to the existing set of half trees, and update the minimum node. To meld two heaps, unite their sets of half trees and update the minimum node. To delete the minimum in a heap, take apart the half tree rooted at the minimum node by deleting it and walking down the path from its child through unordered children, making each node on the path together with its ordered subtree into a half tree. Group the half trees into a maximum number of pairs of equal rank and run the corresponding matches. (Each half tree except at most one per rank participates in one match.) Update the minimum node.

To analyze one-pass binomial queues we define the potential of a heap to be the number of half trees it contains. Lemma 1 holds for one-pass binomial queues, so no rank exceeds  $\lg n$ . A make-heap, find-min, or meld operation takes O(1) time and does not change the potential. An insertion takes O(1) time and increases the potential by one. Thus each of these operations takes O(1) amortized time. Consider a minimum deletion. Disassembling the half tree rooted at the minimum node increases the number of trees and the potential by at most  $\lg n$ . Let h be the number of half trees after the disassembly but before the round of fair matches. The total time for the minimum deletion is O(h), including the time to pair the trees by rank. There are at least  $(h - \lg n)/2 - 1$  fair matches, reducing the potential by at least this amount. If we scale the running time so that it is at most h/2, then the amortized time of the minimum deletion is  $O(\log n)$ . (Scaling the running time is equivalent to multiplying the potential by a constant factor.)

The same analysis applies if we do arbitrary additional fair matches during a minimum deletion. The extreme case is to continue doing fair matches until at most one half tree per rank remains, as in the original version of binomial queues.

## **3** Key Decrease and Arbitrary Deletion

Our next goal is to add key decrease as an O(1)-time operation. Once key decrease is supported, one can delete an arbitrary item by decreasing its key to  $-\infty$  and doing a minimum deletion.

A parameter of both key decrease and arbitrary deletion is the heap containing the given item. If the application does not provide this information and melds occur, one needs a separate data structure to maintain the partition of items into heaps. With such a data structure, the time to find the heap containing a given item is small but not O(1) [17].

Fibonacci heaps [11] were invented specifically to support key decrease efficiently, but they require two extra pointers per node (in the heap-ordered representation, to the parent and previous sibling) and they are slower in practice than other heap implementations [22, 23]. All the alternatives to Fibonacci heaps with the same amortized efficiency, including [4, 7, 15, 19, 24], have the property that decreasing a key can cause a cascade of tree restructuring, possibly delayed. But such restructuring is unnecessary, as we show by introducing a new data structure, the rank-pairing heap, or rp-heap, in which the trees have arbitrary structure and the only cascading is of rank changes.

To obtain rp-heaps, we modify one-pass binomial queues to support key decrease. To the halfordered representation we add parent pointers. We also relax the requirement that the rank of a child be exactly one less than that of its parent. Let p(x) and r(x) be the parent and rank of node x, respectively. The *rank difference* of x is r(p(x)) - r(x); the rank difference is defined only if x has a parent. We maintain the ranks so that each leaf has rank zero and the two children of a non-root have rank differences 1 and 1, or 0 and at least 1. (One of these children can be missing; the rank of a missing node is -1.) We call a half tree with ranks that obey this rule a *type-1 half tree*, and we call a set of heap-ordered type-1 half trees a *type-1 rank-pairing heap* or *rp-heap*.

Lemma 1 extends to type-1 half trees, so the maximum rank of a node in a type-1 rp-heap is  $\lg n$ , and the number of bits needed to store ranks is  $\lg \lg n$  per node, matching Fredman's lower bound.

We represent an rp-heap by a singly-linked circular list of the roots of its half trees, with the minimum node first. We implement making a heap, insertion, melding, and minimum deletion exactly as in one-pass binomial queues. Fair matches maintain the rank rule. (See Figure 2.)

We implement key decrease as follows. (See Figure 3.) To decrease the key of item x in rp-heap H by  $\Delta$ , subtract  $\Delta$  from the key of x and update the minimum node. If x is a root, stop. Otherwise, let y be the unordered child of x. Detach the subtrees rooted at x and y, and reattach the subtree rooted at y in place of the original subtree rooted at x. Add the half tree rooted at x to the set of trees representing H. There may now be a violation of the rank rule at p(y), whose new child, y, may have lower rank than x, the node it replaces. To check for a violation and restore the rank rule if necessary, let u = p(y) and repeat the following step until it stops:

Decrease rank (type 1): If u is the root, stop. Otherwise, let v and w be the children of u. Let k equal r(v) if r(v) > r(w), r(w) if r(w) > r(v), or r(w) + 1 if r(v) = r(w). If k = r(u), stop. Otherwise, let r(u) = k and u = p(u).

If u breaks the rank rule, it obeys the rule after its rank is decreased, but p(u) may not. Rank decreases propagate up along a path through the tree until all nodes obey the rule. Each successive rank decrease is by the same or a smaller amount. The definition of *decrease rank* assumes that ranks of roots are not maintained explicitly.

If we mildly restrict the fair matches done during minimum deletions, we can prove that type-1 rp-heaps match the efficiency of Fibonacci heaps, but the analysis is complicated and yields larger constant factors than one would like. Before doing this, we consider a variant obtained by relaxing

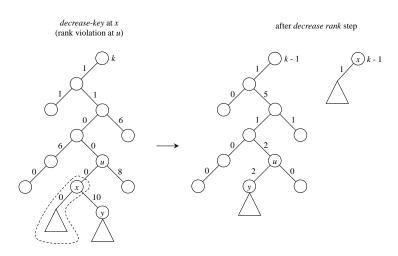


Fig. 3. Key decrease in a type-1 rp-heap.

the rank rule. Its analysis is simpler, with small constant factors, and we expect it to be efficient in practice.

The new rank rule is that each leaf has rank zero, and the two children of a non-root have rank differences 1 and 1, or 1 and 2, or 0 and at least 2. We call a half tree with ranks that obey this rule a *type-2 half tree*, and we call a set of heap-ordered type-2 half trees a *type-2 rank-pairing heap*. The implementations of the heap operations are exactly the same on type-2 rp-heaps as on type-1 rp-heaps, except for key decrease, which is the same except that the rank decrease step becomes the following:

Decrease rank (type 2): If u is the root, stop. Otherwise, let v and w be the children of u. Let k equal r(v) if r(v) > r(w) + 1, r(w) if r(w) > r(v) + 1, or  $\max\{r(v) + 1, r(w) + 1\}$  otherwise. If k = r(u), stop. Otherwise, let r(u) = k and u = p(u).

**Lemma 2.** A type-2 half tree of rank k contains at least  $\phi^k$  items, where  $\phi = (1 + \sqrt{5})/2$  is the golden ratio.

*Proof.* A type-2 half tree whose root is not a leaf can be decomposed into two type-2 half trees by detaching the ordered subtree of the root, detaching the unordered subtree of the old child of the root, and making the latter tree into the ordered subtree of the root. It follows from the rank rule that the minimum number of items  $n_k$  in a type-2 half tree of rank k satisfies the recurrence  $n_0 = 1, n_1 = 2, n_k = n_{k-1} + n_{k-2}$  for  $k \ge 2$ . This is the recurrence for the Fibonacci numbers  $F_k$ , offset by two. Hence  $n_k = F_{k+2} \ge \phi^k$  [20, p. 18].

To analyze type-2 rp-heaps, we call a node *bad* if it has an ordered child of rank difference two or more and *good* otherwise. A node of rank one or more with a missing ordered child is bad. Roots are good. Bad nodes can cause extra matches during minimum deletions; they are created only by key decrease operations.

To analyze the heap operations, we define the potential of a node of rank k to be k, k + 1, or k + 2 if it is a good child, a bad child, or a root, respectively. We define the potential of a heap to be the sum of the potentials of its nodes.

A make-heap, find-min, or meld operation takes O(1) time and does not change the potential. An insertion takes O(1) time and creates one new root of rank zero, for a potential increase of 2. Hence each of these operations takes O(1) amortized time.

Consider the deletion of a minimum node x of rank k. Deletion of x reduces the potential by k+2. Consider the effect on the potential of disassembling the half tree rooted at x. Let y be a node on the path from ord(x) descending through unordered children. There are three cases. If unord(y) has rank difference zero, then y is bad, and making it into a root cannot increase its potential. (Its rank decreases by at least 1.) If unord(y) has rank difference 1, ord(y) has rank difference at least one. In this case making y into a root can increase its potential by at most 2. If unord(y) has rank difference 2 or more, making y into a root can increase its potential by at most 3. In either of the last two cases, unord(y) can be missing. It follows that the new roots increase in potential by a total of at most 2k, and the net potential increase caused by the disassembly is at most  $k - 2 \le \log_{\phi} n - 2$  by Lemma 2.

Let h be the number of half trees in the heap after the disassembly. The entire minimum deletion takes O(h) time. A fair match of two half trees of rank j results in a new half tree of rank j + 1 whose root has a child of rank j, resulting in a net potential drop of one. There are at least  $(h - \log_{\phi} n)/2 - 1$  fair matches, reducing the potential by at least this much. The decrease in potential caused by the entire minimum deletion is thus at least  $h/2 - (3/2) \log_{\phi} n$ . If we scale the time for the minimum deletion so that it is at most  $h/2 + O(\log n)$ , then its amortized time is  $O(\log n)$ .

It remains to analyze key decrease. Consider a key decrease at a node x of rank k. If x is a root, the key decrease takes O(1) time and has no effect on the potential. Suppose x is not a root. Breaking the half tree containing x into two half trees takes O(1) time and increases the potential by at most 4: node x begins with potential at least k but once it becomes a root it has potential at most k + 3, and the old parent of x may become bad, either because its ordered child is now missing or because its new ordered child has smaller rank than its old ordered child. We scale the time so that each iteration of *decrease rank* takes time at most 1. Each iteration either decreases the rank of a node or is the last iteration, so the amortized time of an iteration other than the last is at most zero, not counting increases in potential caused by newly bad nodes. A node u other than the old parent of node x can only become bad if r(ord(u)) decreases by some amount and then r(u) decreases by less. If r(ord(u)) decreases by at least 2, the corresponding potential decrease pays for both the rank decrease step at ord(u) and for u becoming bad. If r(ord(u)) decreases by 1, the rank decrease step at u is the last. It follows that key decrease takes O(1) amortized time.

The maximum rank for a given heap size is the same for type-2 rp-heaps as for Fibonacci heaps, namely at most  $\log_{\phi} n$ . The worst-case time for a key decrease in an rp-heap (of either type) is  $\Theta(n)$ , as it is in a Fibonacci heap [11]. One can reduce the worst-case time for a key decrease to O(1) by delaying each such operation until the next minimum deletion. This requires keeping a list of possible minimum nodes that includes all the roots and all the nodes whose keys have decreased since the last minimum deletion. Making this change complicates the data structure and is likely to worsen its performance in practice.

Now we turn to the analysis of type-1 rp-heaps. We use the following terminology. A node is an i-child if it has rank difference i, and an i, j-node if its children have rank differences i and j; this definition does not distinguish between ordered and unordered children. A leaf is a 1,1-node. We call a node green if it is a leaf but not a root, or it and both of its children are 1,1-nodes; yellow if it is a 0,1 node whose 0-child is a 1, 1-node, or it is a root with no child or with a 1, 1-child; and red otherwise. Red nodes can cause extra matches during minimum deletions; they are created only by key decrease operations. Yellow nodes block the propagation of rank decreases: a rank decrease step on a yellow node is terminating. A key decrease can turn green nodes into yellow nodes but not red ones, and can turn at most one yellow node into a red node.

Our analysis requires a restriction on the fair matches. We consider two such restrictions; the first has a simpler analysis and the second has a simpler, natural implementation. First we assume that the fair matches done during a minimum deletion preferentially match as many red roots as possible. One way to guarantee this is to maintain a bucket for each rank, initially empty. Process the half trees one-at-a-time, beginning with the ones with red roots and finishing with the others. To process a half tree, insert it into the bucket for its rank if this bucket is empty; if not, do a fair match of its root with the root of the half tree in the bucket, and add the resulting half tree to the output set, leaving the bucket empty. Once all the half trees have been processed, add to the output set any tree remaining in a bucket. This process assures that there is at most one match per rank between a red root and a yellow root.

We define the potential of a node of rank k to be k, k + 2, or k + 4 if it is a green or yellow child, a yellow root, or it is red, respectively. The potential of a heap is the sum of the potentials of its nodes.

The analysis of make-heap, find-min, meld, and insert is exactly the same as for type-2 rp-heaps: each such operation takes O(1) amortized time; only insertions increase the potential, by two units per insertion.

Consider the deletion of a minimum node x of rank k. Deletion of x reduces the potential by at least k + 2. Disassembly of the half tree rooted at x produces a set of half trees, one per node on the path from ord(x) descending through unordered children. The nodes on this path become roots; they are the only nodes whose potentials can change. Except for at most 2 units of potential, we shall charge each such increase to a rank between 0 and k - 1, at most 4 units per rank. The total charge is then at most 4k + 2, so the potential increase caused by the disassembly is at most 4k + 2 - (k + 2) = 3k.

Let y be a node on the path. We consider five cases, one of which has two subcases. If y is red and ord(y) is not a 0-child, then its potential does not increase when it becomes a root; indeed, if y becomes yellow, its potential drops by at least 2. If y is red and ord(y) is a 0-child, then the potential of y increases by 1 if it becomes a red root and drops by 1 if it becomes a yellow root: its rank increases by 1. If y is yellow and ord(y) is a 0-child, the potential of y increases by 3 when it becomes a root: its rank increases by 1 and it stays yellow. If y is green, its potential increases by 2 when it becomes a root: it becomes yellow. In each of these cases we charge any increase in potential to the rank of y.

If y is yellow and unord(y) is a 0-child, the potential of y increases by 4 if it becomes a red root or by 2 if it becomes a yellow root. From y walk down the path of unordered children until coming to a node z that is not green or is the last node on the path. Node z is a 1,1-node, so it is not yellow. If it is red, its potential does not increase when it becomes a root, since its rank does not change. We charge the potential increase of y to the rank of z. If z is green, it is a leaf. We charge 2 units of the potential increase of y to rank zero; any remainder (2 units) is uncharged. The total charge to rank zero is 4, 2 for y and 2 for z. As claimed, each rank is charged at most 4 units and at most 4 units are uncharged. We conclude that the potential increase caused by the disassembly is at most  $3k \leq 3 \lg n$ .

Let h be the number of half trees in the heap after the disassembly. The entire minimum deletion takes O(h) time. There are at least  $(h - \lg n)/2 - 1$  fair matches. Each match of two red roots reduces the potential by 1 since the winner becomes yellow; so does each match of two yellow roots. A match between a yellow root and a red root can increase the potential by at most 1, but there are at most  $\lg n$  such matches. Thus the minimum deletion reduces the potential by at least  $(h - \lg n)/2 - 5 \lg n - 1 = h/2 - (11/2) \lg n - 1$ . If we scale the time for the minimum deletion so that it is at most h/2, then the amortized time is  $O(\log n)$ .

Finally, consider decreasing the key of node x. The only nodes whose potential can increase are x and its ancestors. If unord(x) is a 0-child, there is only one rank decrease step, and x becoming a root increases its potential by at most 4. If unord(x) is not a 0-child, x becoming a root increases its potential by at most 3: its rank may increase by 1, but it cannot become red. (It may already be red.)

Let u be a proper ancestor of x that becomes red as a result of the key decrease. Node u cannot be green before the key decrease, it must be yellow. If it is yellow, its rank cannot decrease unless it is a root. Thus it is the last node subject to a key decrease step; its potential increases by at most 4. Each key decrease step except the last decreases the rank of a node and hence the potential by at least 1. If we scale the actual time of a rank decrease step so it is at most 1, then the amortized time of such a step is at most zero unless it is the last step. Thus key decrease takes O(1) amortized time.

A more natural way to do matches during a minimum deletion is to preferentially pair half trees produced by the disassembly. To do this, use the same bucketing scheme as in the method that preferentially pairs red roots, but process the half trees produced by the disassembly first, in the order they are produced, followed by the remaining half trees. This method avoids the need to check root colors, at an extra cost of at most 2 units of potential per key decrease.

In order to analyze the method, we need a more elaborate potential function. We call a root *fresh* if it has just been created by a disassembly, its rank is at least 1, and it has not yet participated in a match. A fresh root becomes *stale* once it wins a match or the minimum deletion is finished. Thus all roots are stale except in the middle of minimum deletions. We define the potential of a node of rank k to be k, k + 2, k + 4, or k + 6 if it is a green or yellow child, a stale yellow root, a red child or a fresh root, or a stale red root, respectively. This gives fresh yellow roots and stale red roots two extra units of potential.

These definitions have the following effect. The winner of a fair match is a stale yellow root. Such a match reduces the potential by at least 1 unless it matches a fresh red root against a stale yellow root, in which case it increases the potential by at most 1. After the matches between fresh roots, each remaining fresh red root is responsible for up to 2 units of extra potential, either because it is matched against a stale yellow root or it becomes stale without being matched. There is at most one such root per positive rank. We charge the extra 2 units against the rank.

The analysis of make-heap, find-min, meld, and insert does not change. The analysis of key decrease is the same except that the potential can increase by an additional 2 units because a stale red root has 2 extra units of potential; at most one node needs 2 extra units, the node whose key decreases or the root of the original tree containing it, but not both.

Consider the deletion of a minimum node x of rank k. In the disassembly process we need to account for 2 extra units of potential for each fresh yellow root. This extra potential does not materially affect the analysis unless the new root y was previously yellow and ord(y) was a 0-child. In this case y must be the last fresh node of its rank, which means that there is no fresh red root of this rank remaining after the matches between fresh roots. We charge the extra 2 units against the rank. The analysis of the matching process proceeds as before; the constant factors are exactly the same. The extra 2 units of potential per rank in the new analysis correspond to the at most  $\lg n$  matches between red and yellow roots in the old analysis. We conclude that minimum deletion takes  $O(\log n)$  amortized time.

# 4 Simpler Key Decreases?

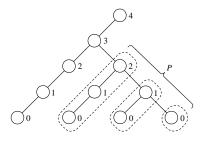
It is natural to ask whether there is an even simpler way to decrease keys while retaining the amortized efficiency of Fibonacci heaps. We give two answers: "no" and "maybe". We answer "no" by showing that two possible methods fail. The first method allows arbitrarily negative but bounded positive rank differences. With such a rank rule, the rank decrease process following a key decrease need examine only ancestors of the node whose key decreases, not their siblings. Such a method can take  $\Omega(\log n)$  time per key decrease, however, as the following counterexample shows. Let *b* be the maximum allowed rank difference. Choose *k* arbitrarily. By means of a suitable sequence of insertions and minimum deletions, build a heap that contains a perfect half tree of each rank from 0 through bk + 1. Let x be the root of the half tree of rank bk + 1. Consider the path of unordered children descending from ord(x). Decrease the key of each node on this path whose rank is not divisible by b. Each such key decrease takes O(1) time and does not violate the rank rule, so no ranks change. Now the path consists of k + 1 nodes, each with rank difference b except the topmost. Decrease the keys of these nodes, smallest rank to largest. Each such key decrease will cause a cascade of rank decreases all the way to the topmost node on the path. The total time for these k + 1 key decreases is  $\Omega(k^2)$ . After all the key decreases the heap contains three perfect half trees of rank zero and two of each rank from 1 through bk. A minimum deletion (of one of the roots of rank zero) followed by an insertion makes the heap again into a set of perfect half trees, one of each rank from 0 through bk + 1. Each execution of this cycle does  $O(\log n)$  key decreases, one minimum deletion, and one insertion, and takes  $\Omega(\log^2 n)$  time.

The second, even simpler method spends only O(1) time worst-case on each key decrease, thus avoiding arbitrary cascading. In this case by doing enough operations one can build a half tree of each possible rank, up to a rank that is  $\omega(\log n)$ . Once this is done, repeatedly doing an insertion followed by a minimum deletion (of the just-inserted item) will result in each minimum deletion taking  $\omega(\log n)$  time. Here are the details. Suppose each key decrease changes the ranks of nodes at most d pointers away from the node whose key decreases, where d is fixed. Choose k arbitrarily. By means of a suitable sequence of insertions and minimum deletions, build a heap that contains a perfect half tree of each rank from 0 through k. On each node of distance d + 2 or greater from the root, in decreasing order by distance, do a key decrease with  $\Delta = \infty$  followed by a minimum deletion. No roots can be affected by any of these operations, so the heap still consists of one half tree of each rank, but each heap contains only  $2^{d+1}$  nodes, so there are  $\lfloor n/2^{d+1} \rfloor$  half trees. Now repeat the cycle of an insertion followed by a minimum deletion. Each such cycle takes  $\Omega(n/2^{d+1})$  time. The choice of "d + 2" in this construction guarantees that no key decrease can reach the child of a root, which implicitly stores the rank of the root.

This construction works even if we add extra pointers to the half trees, as in Fibonacci heaps. In a half tree, the *ordered ancestor* of a node x is the parent of the nearest ancestor of x (including x itself) that is an ordered child. The ordered ancestor corresponds to the parent in the heap-ordered representation. Suppose we augment half trees with ordered ancestor pointers. Even for such an augmented structure, the latter construction gives a bad example, except that the size of a constructed half tree of rank k is  $O(k^{d+1})$  instead of  $O(2^{d+1})$ , and each cycle of an insertion followed by a minimum deletion takes  $\Omega(n^{1/(d+2)})$  time.

One limitation of this construction is that building the initial set of half trees takes a number of operations exponential in the size of the heap on which repeated insertions and minimum deletions are done. Thus it is not a counterexample to the following question: is there a fixed d such that if each key decrease is followed by at most d rank decrease steps (say of type 1), then the amortized time is O(1) per insert, meld, and key decrease, and  $O(\log m)$  per deletion, where m is the total number of insertions? A related question is whether Fibonacci heaps without cascading cuts have these bounds. We conjecture that the answer is yes for some positive d, perhaps even d = 1. The following counterexample shows that the answer is no for d = 0; that is, for the method in which a key decrease changes no ranks except for the implicit ranks of roots. For arbitrary k, build a half tree of each rank from 0 through k, each consisting of a root and a path of ordered children, inductively as follows. Given such half trees of ranks 0 through k - 1, insert an item less than all those in the heap and then do k cycles, each consisting of an insertion followed by a minimum deletion that deletes the just-inserted item. The result will be one half tree of rank k consisting of the root, a path of ordered children descending from the root, a path P of unordered children descending from the ordered children childr

of the root, and a path of ordered children descending from each node of P; every child has rank difference 1. (See Figure 4.) Do a rank decrease on each node of P. Now there is a collection of half trees of rank 0 through k except for k - 1, each a path. Repeat this process on the set of half trees up to rank k - 2, resulting in a set of half trees of ranks 0 through k with k - 2 missing. Continue in this way until only rank 0 is missing, and then do a single insertion. Now there is a half tree of each rank, 0 through k. The total number of heap operations required to increase the maximum rank from k - 1 to k is  $O(k^2)$ , so in m heap operations one can build a set of half trees of each possible rank up to a rank that is  $\Omega(m^{1/3})$ . Each successive cycle of an insertion followed by a minimum deletion takes  $\Omega(m^{1/3})$  time.



**Fig. 4.** A half tree of rank k = 4 buildable in  $O(k^3)$  operations if key decreases do not change ranks. Key decreases on the unordered children detach the circled subtrees.

# 5 Remarks

We have presented a new data structure, the rank-pairing heap, that combines the performance guarantees of Fibonacci heaps with simplicity approaching that of pairing heaps. Our preliminary experiments suggest that rank-pairing heaps may be competitive with pairing heaps in practice, and we intend to do more-thorough experiments. Several theoretical questions remain: Can the restriction on fair matches in type-1 rp-heaps be removed? Can the constant factors in the analysis of rp-heaps be reduced? How is efficiency affected if only a constant number of rank decrease steps are done after a key decrease? Is there a nice one-tree version of rp-heaps analogous to balanced tournaments? Can Brodal's worst-case-efficient heap implementation be simplified?

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### References

- G. Brodal. Worst-case efficient priority queues. In SODA, pages 52–58. Society for Industrial and Applied Mathematics Philadelphia, PA, USA, 1996.
- 2. M. R. Brown. Implementation and analysis of binomial queue algorithms. *SIAM J. on Comput.*, pages 298–319, 1978.

- 3. E. Dijkstra. A note on two problems in connexion with graphs. *Numerische Mathematik*, 1(1):269–271, 1959.
- J. R. Driscoll, H. N. Gabow, R. Shrairman, and R. E. Tarjan. Relaxed heaps: an alternative to Fibonacci heaps with applications to parallel computation. *Comm. of the ACM*, 31(11):1343–1354, 1988.
- R. Dutton. The weak-heap data structure. Technical report, Technical Report CS-TR-92-09, University of Central Florida, Orlando, FL 32816, 1992.
- 6. Edmonds,, J. Optimum branchings. J. Res. Nat. Bur. Standards, B71:233-240, 1967.
- 7. A. Elmasry. Violation heaps: A better substitute for fibonacci heaps. CoRR, abs/0812.2851, 2008.
- 8. A. Elmasry. Pairing heaps with  $O(\log \log n)$  decrease cost. In SODA, pages 471–476. Society for Industrial and Applied Mathematics, 2009.
- 9. M. L. Fredman. On the efficiency of pairing heaps and related data structures. J. ACM, 46(4):473-501, 1999.
- M. L. Fredman, R. Sedgewick, D. D. Sleator, and R. E. Tarjan. The pairing heap: A new form of self-adjusting heap. *Algorithmica*, 1(1):111–129, 1986.
- M. L. Fredman and R. E. Tarjan. Fibonacci heaps and their uses in improved network optimization algorithms. J. of the ACM, 34(3):596–615, 1987.
- M. L. Fredman and D. E. Willard. Trans-dichotomous algorithms for minimum spanning trees and shortest paths. J. of Comput. and Sys. Sci., pages 533–551, 1994.
- 13. H. N. Gabow, Z. Galil, T. H. Spencer, and R. E. Tarjan. Efficient algorithms for finding minimum spanning trees in undirected and directed graphs. *Combinatorica*, 6(2):109–122, 1986.
- 14. Y. Han and M. Thorup. Integer sorting in  $O(n\sqrt{\log \log n})$  expected time and linear space. In *FOCS*, pages 135–144, 2002.
- 15. P. Høyer. A general technique for implementation of efficient priority queues. In ISTCS, pages 57-66, 1995.
- D. W. Jones. An empirical comparison of priority-queue and event-set implementations. *Communications of the ACM*, 29(4):300–311.
- 17. H. Kaplan, N. Shafrir, and R. E. Tarjan. Meldable heaps and boolean union-find. In *STOC*, pages 573–582, 2002.
- H. Kaplan and R. E. Tarjan. New heap data structures. Technical report, Technical Report TR-597-99, Department of Computer Science, Princeton University, 1999.
- 19. H. Kaplan and R. E. Tarjan. Thin heaps, thick heaps. ACM Trans. Algorithms, 4(1):1–14, 2008.
- 20. D. E. Knuth. *The Art of Computer Programming, Volume 1: Fundamental Algorithms*. Addison-Wesley, 1973.
- 21. D. E. Knuth. The Art of Computer Programming, Volume 3: Sorting and Searching. Addison-Wesley, 1973.
- 22. A. M. Liao. Three priority queue applications revisited. Algorithmica, 7:415-427, 1992.
- 23. B. Moret and H. Shapiro. An empirical analysis of algorithms for constructing a minimum spanning tree. *DIMACS Series in Disc. Math. and Theor. Comput. Sci.*, 15:99–117, 1994.
- 24. G. L. Peterson. A balanced tree scheme for meldable heaps with updates. Technical Report GIT-ICS-87-23, School of Informatics and Computer Science, Georgia Institute of Technology, 1987.
- 25. S. Pettie. Towards a final analysis of pairing heaps. pages 174–183. IEEE Computer Society, 2005.
- 26. J. T. Stasko and J. S. Vitter. Pairing heaps: experiments and analysis. Commun. ACM, 30(3):234-249, 1987.
- 27. R. E. Tarjan. Data Structures and Network Algorithms. SIAM, 1983.
- 28. R. E. Tarjan. Amortized computational complexity. SIAM J. on Algebraic and Disc. Methods, 6:306, 1985.
- 29. J. Vuillemin. A data structure for manipulating priority queues. Commun. ACM, 21(4):309-315, 1978.