6.851: Advanced Data Structures

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1 Overview

In this lecture, we look at various data structures for static trees, in which, given a static tree, we perform some preprocessing and then answer queries on the data structure. The three problems we look at in this lecture are range minimum queries (RMQ), least common ancestors (LCA), and level ancestors (LA); we will support all these queries in constant time per operation and linear space.

1.1 Range Minimum Query (RMQ)

In the range minimum query problem, we are given (and we preprocess) an array A of n numbers. In a query, the goal is to find the minimum element in a range spanned by A[i] and A[j]:

$$RMQ(i, j) = (arg)min\{A[i], A[i+1], \dots, A[j]\}$$
$$= k, \text{ where } i \le k \le j \text{ and } A[k] \text{ is minimized}$$

We care not only about the value of the minimum element, but also about the index k of the minimum element between A[i] and A[j]; given the index, it is easy to look up the actual value of the minimum element, so it is a more interesting problem to find the index of the minimum element between A[i] and A[j].

The range minimum query problem is not a tree problem, but it closely related to a tree problem (least common ancestor).

1.2 Lowest Common Ancestor (LCA)

In the lowest common ancestor problem, we want to preprocess a rooted tree T with n nodes. In a query, we are given two nodes x and y and the goal is to find their lowest common ancestor in T:



1.3 Level Ancestor (LA)

Finally, we will also solve the level ancestor problem, in which we are again given a rooted tree T. Given a node x and an integer k, the query goal is to find the k^{th} ancestor of node x:

$$LA(x,k) = parent^k(x)$$

Of course, k cannot be larger than the depth of x.



All of these problems will be solved in the word RAM model, though the use of model is not as essential as it has been in the integer data structures we have discussed over the previous lectures. Although lowest common ancestor and level ancestor seem like similar problems, fairly different techniques are necessary to solve them as far as anyone knows. The range minimum query problem, however, is basically identical to that of finding the lowest common ancestor.

2 Reductions between RMQ and LCA

2.1 Cartesian Trees: Reduction from RMQ to LCA

A Cartesian tree is a nice reduction mechanism from an array A to a binary tree T, dating back to a 1984 paper by Gabow, Bentley, and Tarjan [1], and provides an equivalence between RMQ and LCA.

To construct a Cartesian tree, we begin with the minimum element of the array A, which we can call A[i]. This element becomes the root of the Cartesian tree T. Then the left subtree of T is a Cartesian tree on all elements to the left of A[i] (which we can write as A[< i]), and the right subtree of T is likewise a Cartesian tree on the elements A[> i].

An example is shown below for the array A = [8, 7, 2, 8, 6, 9, 4, 5]. The minimum of the array is 2, which gets promoted to the root. This decomposes the problem into two halves, one for the left subarray [8,7] and one for the right subarray [8,6,9,4,5]. 7 is the minimum element in the left subarray and becomes the left child of the root; 4 is the minimum element of the right subarray and is the right child of the root. This procedure continues until we get the binary tree in the diagram below.



The resulting tree T is a min heap. More interestingly, the result range minimum query for a given range in the array A is the lowest common ancestor of those two endpoints in the corresponding Cartesian tree T.

In the case of ties between multiple equal minimum elements, two options are:

- 1. Break ties arbitrarily, picking one of the equal elements as the minimum.
- 2. Consider all of the equal elements to be one "node", making a non-binary tree.

Because the second option is slightly messier, in this class we will break ties arbitrarily, though either option will not affect the answer.

2.1.1 Construction in linear time

Construction of the Cartesian tree according to the naive recursive algorithm will take at least $n \lg n$ time, and may even take quadratic time. In fact, Cartesian trees can be computed in linear time, using a method that is basically the same as the method seen in the last lecture for building a compressed trie in linear time.

Walk through the array from left to right, inserting each element into the tree by walking up the right spine of the tree (starting from the leaf), and inserting the element in the appropriate place. Because we are building the tree from left to right, each inserted element will by definition be the rightmost element of the tree created so far.

For example, if we have a tree for which the subarray [2,4,5] has been inserted, and the next element is 3, then insertion has the following result:



We walk up the right spine of the tree starting at 5, and continue until we reach 2 (the root), which is the first element smaller than 3. The edge between 2 and 4 is replaced with a new edge from 2 to 3, and the previous right subtree of 2 becomes the left subtree of 3.

Each such insertion takes constant amortized time; although sometimes paths may be long, each insertion only touches nodes along the right spine of the tree, and any node along the right spine that has been touched ends up in the left subtree of the inserted node. Any node along the right spine is touched at most once, and we can charge the expensive inserts to the decrease in length of the right spine.

Therefore, construction of Cartesian trees can be done in linear time, even in the comparison model.

2.2 Reduction from LCA to RMQ

We can also reduce in the other direction, reducing from LCA to RMQ by reconstructing an array A when we are given a binary tree T. To do this, we do an in-order traversal of the nodes in the tree. However, we must have numbers to use as the values of the array; to this end, we label each node with its depth in the tree.



This sequence behaves exactly like the original array A = [8, 7, 2, 8, 6, 9, 4, 5], from which this tree was constructed. The result for RMQ(i, j) on the resulting array A is the same as calling LCA(i, j) on the input tree for the corresponding nodes.

2.3 RMQ universe reduction

An interesting consequence of the reductions between RMQ and LCA is that they allow us to do universe reductions for RMQ problems. There are no guarantees on the bounds of any numbers in the original array given for range minimum queries, and in general the elements may be in any arbitrary ordered universe. By chaining the two reductions above, first by building a Cartesian tree from the elements and then by converting back from the tree to an array of depths, we can reduce the range to the set of integers $\{0, 1, \ldots, n-1\}$.

This universe reduction is handy; the algorithms described above assume a comparison model, but after the pair of reductions we can now assume that all of the inputs are small integers, which allows us to solve things in constant time in the word RAM model.

3 Constant time LCA and RMQ

3.1 Results

The LCA and RMQ problems can both be optimally solved with constant query time and linear storage space. The first known technique is documented in a 1984 paper by Harel and Tarjan, [2]. In lecture we looked at an algorithm based on a 2004 paper by Bender and Colton, [3].

3.2 Reduction from LCA to ± 1 RMQ

The algorithm by Bender and Colton solves a special case of the RMQ problem where adjacent values differ by either +1 or -1 called ± 1 RMQ.

First, we will take a look at a reduction from LCA to ± 1 RMQ. The earlier reduction does not work as differences might have absolute values larger than 1. For our new approach, we perform an Eulerian tour based on the in-order traversal and write down every visit in our LCA array. At every step of the tour we either go down a level or up a level, so the difference between two adjacent values in our array is either +1 or -1.



Since every edge is visited twice, we have more entries in the array than in the original in-order traversal, but still only O(N). To answer LCA(x, y), calculate RMQ(in-order(x), in-order(y)) where in-order(x) is the in-order occurrence of the node x in the array. These occurrences can be stored while creating the array. Observe that this new array can also be created by filling in the gaps in the array from the original algorithm. Thus the minimum between any two original values does not change, and this new algorithm also works.

3.3 Constant time, $n \lg n$ space RMQ

A simple datastructure than can answer RMQ queries in constant time but uses only $n \lg n$ space can be created by precomputing the RMQ for all intervals with lengths that are powers of 2. There are a total of $O(n \lg n)$ such intervals, as there are $\lg n$ intervals with lengths that are powers of 2 no longer than n, with n possible start locations.

We claim that any queried interval is the (non-disjoint) union of two power of 2 intervals. Say the query has length k. Then the query can be covered by the two intervals of length $2^{\lfloor \lg k \rfloor}$ that touch the beginning and ending of the query. The query can be answered by taking the min of the two precomputed answers. Observe that this works because we can take the min of an element twice

without any problems. Furthermore, this also enables us to store the location of the minimum element.

3.4 Indirection to remove log factors

We have seen indirection used to remove log factors of time, but here we will apply indirection to achieve a O(n) space bound. Divide the array into bottom groups of size $\frac{1}{2} \lg n$ (this specific constant will be used later). Then, store a parent array of size $2n/\lg n$ that stores the min of every group.

Now a query can be answered by finding the RMQ of a sequence in the parent array, and at most two RMQ queries in bottom groups. Note that we might have to answer two sided queries in a bottom group for sufficiently small queries. For the parent array we can use the $n \lg n$ space algorithm as the logarithms cancel out.

3.5 RMQ on very small arrays

The only remaining question is how to solve the RMQ problem on arrays of size $n' = \frac{1}{2} \lg n$. The idea is to use lookup tables, since the total number of different possible arrays is very small.

Observe that we only need to look at the relative values in a group to find the location of the minimum element. This means that we can shift all the values so that the first element in the group is 0. Then, once we know the location, we can look in the original array to find the value of the minimum element.

Now we will use the fact the array elements differ by either +1 or -1. After shifting the first value to 0, there are now only $2^{\frac{1}{2} \lg n} = \sqrt{n}$ different possible bottom groups since every group is completely defined by its n' long sequence of +1 and -1s. This total of \sqrt{n} is far smaller than the actual number of groups!

In fact, it is small enough so that we can store a lookup table for any of the n'^2 possible queries for any of the \sqrt{n} groups in $O(\sqrt{n}(\frac{1}{2} \lg n)^2 \lg \lg n)$ bits, which easily fits in O(n) space. Now every group can store a pointer into the lookup table, and all queries can be answered in constant time with linear space for the parent array, bottom groups, and tables.

3.6 Generalized RMQ

We have LCA using ± 1 RMQ in linear space and constant time. Since general RMQ can be reduced LCA using universe reduction, we also have a linear space, constant time RMQ algorithm.

4 Level Ancestor Queries (LAQ)

First we introduce notation. Let h(v) be the height of a node v in a tree. Given a node v and level l, LAQ(v, l) is the ancestor a of v such that h(a) - h(v) = l. Today we will study a variety of data structures with various preprocessing and query times which answer LAQ(v, l) queries. For a data structure which requires f(n) query time and g(n) preprocessing time, we will denote its running

time as $\langle g(n), f(n) \rangle$. The following algorithms are taken from the set found in a paper from Bender and Farach-Colton[4].

4.1 Algorithm A: $\langle O(n^2), O(1) \rangle$

Basic idea is to use a look-up table with one axis corresponding to nodes and the other to levels. Fill in the table using dynamic programming by increasing level. This is the *brute force* approach.

4.2 Algorithm B: $\langle O(n \log n), O(\log n) \rangle$

The basic idea is to use **jump pointers**. These are pointers at a node which reference one of the node's ancestors. For each node create jump pointers to ancestors at levels $1, 2, 4, \ldots, 2^k$. Queries are answered by repeatedly jumping from node to node, each time jumping more than half of the remaining levels between the current ancestor and goal ancestor. So worst-case number of jumps is $O(\log n)$. Preprocessing is done by filling in jump pointers using dynamic programming.

4.3 Algorithm C: $\langle O(n), O(\sqrt{n}) \rangle$

The basic idea is to use a **longest path decomposition** where a tree is split recursively by removing the longest path it contains and iterating on the remaining connected subtrees. Each path removed is stored as an array in top-to-bottom path order, and each array has a pointer from its first element (the root of the path) to it parent in the tree (an element of the path-array from the previous recursive level). A query is answered by moving upwards in this *tree of arrays*, traversing each array in O(1) time. In the worst case the longest path decomposition may result in longest paths of sizes $k, k - 1, \ldots, 2, 1$ each of which has only one child, resulting in a tree of arrays with height $O(\sqrt{n})$. Building the decomposition can be done in linear by by precomputing node heights once and reusing them to find the longest paths quickly.

4.4 Algorithm D: $\langle O(n), O(\log n) \rangle$

The basic idea is to use **ladder decomposition**. The idea is similar to longest path decomposition, but each path is extended by a factor of two backwards (up the tree past the root of the longest path). If the extended path reaches the root, it stops. From the ladder property, we know that node v lies on a longest path of size at least h(v). As a result, one does at most $O(\log n)$ ladder jumps before reaching the root, so queries are done in $O(\log n)$ time. Preprocessing is done similarly to Algorithm C.

4.5 Algorithm E: $\langle O(n \log n), O(1) \rangle$

The idea is to combine jump pointers (Algorithm B) and ladders (Algorithm D). Each query will use one jump pointer and one ladder to reach the desired node. First a jump is performed to get at least halfway to the ancestor. The node jumped to is contained in a ladder which also contains the goal ancestor.

4.6 Algorithm F: $\langle O(n), O(1) \rangle$

An algorithm developed by Dietz[5] also solves LCA queries in $\langle O(n), O(1) \rangle$ but is more complicated. Here we combine Algorithm E with a reduction in the number of nodes for which jump pointers are calculated. The motivation is that if one knows the level ancestor of v at level l, one knows the level ancestor of a descendant of v at level l'. So we compute jump pointers only for leaves, guaranteeing every node has a descendant in this set. So far, preprocessing time is $O(n + L \log n)$ where L is the number of leaves. Unforunately, for an arbitrary tree, L = O(n).

4.6.1 Building a tree with $O(\frac{n}{\log n})$ leaves

Split the tree structure into two components: a **macro-tree** at the root, and a set of **micro-trees** (of maximal size $\frac{1}{4} \log n$) rooted at the leaves of the macro-tree. Consider a depth-first search, keeping track of the orientation of of the *i*th edge, using 0 for downwards and 1 for upwards. A micro-tree can be described by a binary sequence, e.g. W = 001001011 where for a tree of size n, |W| = 2n - 1. So an upper bound the number of micro-trees possible is $2^{2n-1} = 2^{2(\frac{1}{4} \log n)-1} = O(\sqrt{n})$. But this is a loose upper bound, as not all binary sequences are possible, e.g. $00000\ldots$ A valid micro-tree sequences has an equal number of zeros and ones and any prefix of a valid sequence as at least as many zeros as ones.

4.6.2 Use macro/micro-tree for a $\langle O(n), O(1) \rangle$ solution to LAQ

We will use macro/micro-trees to build a tree with $O(\frac{n}{\log n})$ leaves and compute jump pointers only for its leaves (O(n) time). We also compute all microtrees and their look-up tables (see Algorithm A) in $O(\sqrt{n}\log n)$ time. So total preprocessing time is O(n). A query LAQ(v,l) is performed in the following way: If v is in the macro-tree, jump to the leaf descendant of v, then jump from the leaf and climb a ladder. If v is in a micro-tree and LAQ(v,l) is in the micro-tree, use the look-up table for the leaf. If v is in a micro-tree and LAQ(v,l) is not in the micro-tree, then jump to the leaf descendant of v, then jump from the leaf and climb a ladder.

References

- H. Gabow, J. Bentley, R. Tarjan. Scaling and Related Techniques for Geometry Problems. In STOC '84: Proc. 16th ACM Symp. Theory of Computing, pages 135-143, 1984.
- [2] Dov Harel, Robert Endre Tarjan, Fast Algorithms for Finding Nearest Common Ancestors, SIAM J. Comput. 13(2): 338-355 (1984)
- [3] Michael A. Bender, Martin Farach-Colton, The LCA Problem Revisited, LATIN 2000: 88-94
- [4] M. Bender, M. Farach-Colton. The Level Ancestor Problem simplified. Lecture Notes in Computer Science. 321: 5-12. 2004.
- [5] P. Dietz. Finding level-ancestors in dynamic trees. Algorithms and Data Structures, 2nd Workshop WADS '91 Proceedings. Lecture Notes in Computer Science 519: 32-40. 1991.

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