CS 161, Lecture 6 Scribe: Sanjana Rajan (2015)

The plan for this lecture is to finish the analysis of Randomized Quicksort, to show that the sorting algorithms we have seen so far are in some sense optimal, and to give a very different sorting algorithm that can sort in linear time when the numbers are not too big.

## 1 QuickSort Review

Here, we finish the analysis of QuickSort. For more details about QuickSort, including the notation used in these notes, see the notes for Lecture 5. Recall the claims about the runtime of QuickSort proved in the previous lecture.

**Claim 1.** The runtime of QuickSort is O(n+X) where X is the total number of comparisons made by the algorithm.

Claim 2. Let  $z_1 < z_2 < ... < z_n$  refer to numbers in the input array A, but in sorted order. Let  $p_{ij}$  be the probability that  $z_i$  and  $z_j$  are compared at any point of the algorithm. Then, the expected number of comparisons made by QuickSort is

$$E[X] = \sum_{i=1}^{n-1} \sum_{j=i+1}^{n} p_{ij}.$$

Last lecture, we used these theorems to argue that if  $p_{ij} = \frac{2}{j-i+1}$ , then the runtime of RandQuickSort is  $O(n \log n)$ . Thus, to finish our proof of the run time of RandQuickSort, we need to analyze the probabilities  $p_{ij}$  and to show that they are indeed  $p_{ij} = \frac{2}{j-i+1}$ .

Claim 3. For any i, j with i < j:  $p_{ij} = \frac{2}{j-i+1}$ .

*Proof.* We first note that in RandQuickSort, the only time that two elements are compared is when one of them is selected as a pivot. Consider the set  $Z = \{z_i, z_{i+1}, \ldots, z_j\}$ , that is, the elements between  $z_i$  and  $z_j$  in the sorted order. Let RandQuickSort(A, p, r) be the *last call* of RandQuickSort for which all elements of Z are in A(p:r). Thus, in this call, the pivot must be some  $z_k \in Z$  (as otherwise all of Z will be in A <or A >and RandQuickSort(A, p, r) wouldn't be the last call for which  $Z \subset A(p:r)$ ).

Let  $z_k$  be the pivot. We will consider two cases - either  $z_k$  is one of the endpoints,  $z_i$  or  $z_j$ , or  $z_k$  is some other element in Z.

Case 1: Either  $z_k = z_i$  or  $z_k = z_j$ 

In this case,  $z_i$  and  $z_j$  are clearly compared within this call.

Case 2:  $z_k$  is neither  $z_i$  nor  $z_j$ 

In this case,  $z_k$  is one of  $\{z_{i+1}, \ldots, z_{j-1}\}$ . Then  $z_k$  is compared to  $z_i$  and  $z_j$ . Since  $z_i < z_k$  and  $z_k < z_j$ ,  $z_i$  will be in  $A_{\leq}$  and  $z_j$  will be in  $A_{\geq}$ , so  $z_i$  and  $z_j$  will never be compared.

Thus,  $p_{ij} = \Pr(z_i \text{ and } z_j \text{ are compared}) = \Pr(\text{pivot is } z_i \text{ or } z_j | \text{ pivot is in } Z)$ . Because the pivot was picked uniformly at random,  $\Pr(\text{pivot is } z_i \text{ or } z_j | \text{ pivot is in } Z) = \frac{2}{|Z|} = \frac{2}{j-i+1}$ , completing this proof.

## 2 Lower Bounds for Comparison Sorting Algorithms

We call a sorting algorithm a *comparison-based* sorting algorithm if the sorted order is determined solely through pairwise comparisons of elements, and not the actual values that are being compared. The two most well-known comparison sorting algorithms are MergeSort and QuickSort, but there are many others which also share this property.

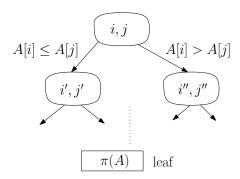


Figure 1: The general framework for a decision tree. In this case, i is compared to j at the root, which could lead to i' being compared to j' (if  $A[i] \leq A[j]$ ) or i'' being compared to j'' (if A[i] > A[j]). The leaves of the tree are permutations of the entries of A.

In this section, we will show that comparison-based sorting algorithms all have inherent limitations. We'll show that any deterministic comparison sorting algorithm must perform  $\Omega(n \log n)$  comparisons to sort n elements. That is, for any sorting algorithm that doesn't look at the actual values being compared, the algorithm's running time is lower bounded by  $\Omega(n \log n)$ .

To show this lower bound, we will completely disregard any operations besides comparisons. Then we can model all comparison sorting algorithms with a class of binary trees called *decision trees*.

Each node of the decision tree corresponding to a comparison-based algorithm represents a comparison that the algorithm makes between two elements of the array. The comparison stored at the root is the first comparison performed by the algorithm. Each node of the tree contains two (ordered) indices i, j of the array, meaning that A(i) and A(j) are two be compared. There are two possible cases - either  $A(i) \leq A(j)$  or A(i) > A(j). The left child of the node corresponds to the next comparison that the algorithm would make if  $A(i) \leq A(j)$ , and the right child corresponds to the next comparison whenever A(i) > A(j). This way any internal node of the decision tree has exactly two children. At each leaf of the decision tree (the termination of the algorithm), there is a permutation of A representing the final sorted order based on the root to leaf path.

Each run of the comparison-based sorting algorithm on an input A corresponds to a path from the root to some leaf in the decision tree. The number of total comparisons made on input A is the length of the corresponding root to leaf path. Thus, the worst-case number of total comparisons for a comparison sorting algorithm will be the height of its decision tree. We use this to prove a lower-bound on the number of comparisons made by an arbitrary comparison sorting algorithm.

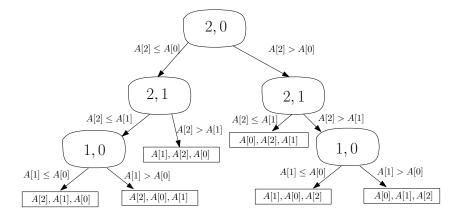


Figure 2: A decision tree for Quicksort running on an array of length 3.

**Theorem 2.1.** Any comparison sorting algorithm must perform  $\Omega(n \log n)$  comparisons to sort n elements.

*Proof.* As stated earlier, the worst-case number of total comparisons for a comparison sorting algorithm will be the height h of its corresponding decision tree. Thus, we will prove this theorem by showing that  $h \ge \Omega(n \log n)$ .

We first note that every permutation of the n indices of the input must appear at one of the leaves of the decision tree; otherwise, there would exist some possible input that would not be correctly sorted. There are n! permutations of these indices, so the number of leaves  $\ell$  is at least n!. Additionally, because the tree is binary, a decision tree of height h will have no more than  $2^h$  leaves. Thus,  $n! \leq \ell \leq 2^h$  which implies that  $h \geq \log(n!)$ .

By Stirling's approximation,  $n! \ge \Omega((\frac{n}{e})^n)$ . Therefore,  $\log(n!) \ge \Omega(\log n^n - \log e^n) = \Omega(n\log(n))$ . Thus, we have  $h > \Omega(n\log n)$ , completing the proof.

## 3 CountingSort

We now look at an integer sorting algorithm called CountingSort, which requires some additional assumptions but has a much improved runtime. The main point is that this algorithm actually uses the values stored the input array to sort instead of purely comparisons. Because of this, for many inputs it can run in linear time.

## 3.1 Algorithm Description and Intuition

Input: Integer array A of length n where all elements are in  $\{0,...k\}$  Output: Sorted integer array B of length n

Given some element x in A, we now consider at which indices it will appear in the sorted array B. Let  $C_i$  be the number of elements that are less than or equal to i. Then, because the number of elements less than x is equivalent to the number of elements less than or equal to x-1, we know that the first x in B will be in position  $C_{x-1}$ . The last occurrence of x will be in position  $C_x - 1$ . Thus, to sort A, it suffices to compute  $C_x$  for every  $x \in \{0, ..., k\}$ . The pseudocode for CountingSort is below.

```
Algorithm 1: CountingSort(A, k)
                                              // |A| = n
If |A| = 1, return A;
B \leftarrow \text{array of length } n;
C \leftarrow \text{array of length } k \text{ initialized to the array of all 0s};
for j from 0 to n-1 do
  C[A[j]] \leftarrow C[A[j]] + 1;
                                                                        Find number of occurrences of each element
for j from 1 to k do
  C[j] \leftarrow C[j] + C[j-1];
                                                                                     Now, C[j] = C_j as defined earlier
for j = n - 1 \rightarrow 0 do
     x \leftarrow A[j];
     B[C[x]-1] \leftarrow x;
     C[x] \leftarrow C[x] - 1;
return B;
```

The correctness proof of the algorithm is as follows. The first for loop computes the number of occurrences of each x in A, storing the number in C[x]. Then the second for loop computes for each x, the number of elements  $\leq x$ . It does this inductively. As the base case,  $C_0$  is exactly the number of 0s in A, so it equals C[0]. Suppose that C[i-1] contains the number of elements  $\leq i-1$ . Then the number of elements  $\leq i$  is C[i-1]+ the number of occurrences of i. Thus, one can compute  $C_i$  by summing C[i-1] and C[i]. Finally, the last loop places each element x in its correct position, via the argument we made earlier.

**Claim 4.** CountingSort sorts n integer elements between 0 and k in O(n+k) time.

*Proof.* This runtime follows easily from the above pseudocode. The first for loop from iterates over n elements and performs constant-time operations on each, so it can be completed in time O(n). Through similar reasoning, the second for loop can be completed in time O(k), and the third for loop can be completed in time O(n). Thus, the total runtime will be O(n+k).

One nice additional property of CountingSort is that the occurrences of some element x will appear in the same order in the sorted array B as they did in the original array A. This property is known as stability and is critical to CountingSort's application to RadixSort, another non-comparison sorting algorithm. It may also be useful if you are sorting objects by some key, but want to maintain the order of the objects who share a key.