

# Design and Analysis of Algorithms, Honors

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The other extreme could be posting the source code of the program. This would be a complete description of an algorithm, because even the computer and execute it. However, source code includes details that **only the computer needs**, such as types, etc. So, when talking about algorithms, it makes sense to use a middleground, known as **pseudocode**.

**Definition 1. Pseudocode** is a descriptive, step by step set of instructions that detail the structure of a program. English is allowed, and the level of detail depends on the context and the audience.

## Lecture 1: Intro and Review

### 1 Intro to Algorithms

#### 1.1 Properties of Algorithms

What kind of properties can an algorithm even have?

**Property.** Algorithms can be categorized by speed, memory, readability (simple to understand), accuracy (approximation quality), and requirements for the input.

The focus of this class will be on **speed**. We will also talk about accuracy, and algorithms will be mostly readable (but not always). We need some language that will allow us to describe algorithms.

#### 1.2 Describing Algorithms

**Example.** Let's take for example selection sort.

**Explanation.** Using plain english, selection sort is: repeatedly finding the smallest element and appending it to the output.

Algorithms can be described at different levels of detail. One extreme is very informally, like the sentence above. This type helps convey the **key concepts**, but can be rather vague, ambiguous, and hard to understand.

**Algorithm 1** Pseudocode for Selection Sort

---

**Any array of elements** *input*

**A sorted array** *output*

```
1: for  $i \leftarrow 1 \dots n$  do
2:   Find  $j \geq i$  with smallest  $arr[j]$ 
3:   Swap  $arr[i]$  and  $arr[j]$ 
4: end for
```

---

More detailed descriptions can also be given with pseudocode.

#### 1.3 Runtime of Algorithms

Many factors can impact the runtime of an algorithm. For example, Selection Sort, when ran on the professor's laptop had a runtime of  $0.017n^2 + 0.669n + 0.114$ ms. This runtime will vary depending on the context, things like hardware, the programming language, and temperature.

Because we cannot determine these constants for every computer, we will just **ignore them** for analyzing algorithms.

**Definition 2. Big-O Notation** is the notation used to analyze runtimes. It essentially ignores unknown constant factors. More formally, it states that  $f(n) = O(g(n))$  if and only if there exists  $n_0$  and  $c > 0$  such that  $f(n) \leq c \cdot g(n) \forall n \geq n_0$ .

**Lemma 1.**  $\lim_{n \rightarrow \infty} \frac{f(n)}{g(n)} < \infty \Rightarrow f(n) = O(g(n))$ .

We can find the runtime of algorithms by counting the number of operations.

**Example.** Selection sort runs in  $O(n^2)$ .

**Explanation.** We have

$$\sum_{i=1}^n \sum_{j=i}^n 1 = \frac{n(n-1)}{2} = O(n^2).$$

Note that the second equality comes from taking the limit.

**Definition 3.** Omega notation denotes the relation  $\geq$ .

**Lemma 2.**  $\lim_{n \rightarrow \infty} \frac{g(n)}{f(n)} \Rightarrow f(n) = \Omega(g(n))$ .

**Definition 4.** Little-O notation denotes the strict relation  $<$ .

**Lemma 3.**  $\lim_{n \rightarrow \infty} \frac{f(n)}{g(n)} = 0 \Rightarrow f(n) = o(g(n))$ .

**Definition 5.** Little Omega notation denotes the strict relation  $>$ .

**Lemma 4.**  $\lim_{n \rightarrow \infty} \frac{g(n)}{f(n)} \Rightarrow f(n) = \omega(g(n))$ .

**Definition 6.** Theta notation denotes the equality relation  $=$ .

**Lemma 5.**  $f(n) = O(g(n)) \wedge f(n) = \Omega(g(n)) \Rightarrow f(n) = \Theta(g(n))$ .

## Lecture 2: Divide and Conquer

### 2 Divide and Conquer

The main idea of divide and conquer is to split the problem into smaller subproblems, and solve those subproblems by doing the same. Finally, we combine the solutions together to solve the main problem.

**Example.** One example of a divide and conquer is Merge Sort.

#### Algorithm 2 Merge Sort

*A, an array input*

**Sorted array output**

```

1: if  $n = 1$  then
2:   return  $A$ 
3: end if
4: left = MergeSort( $A[0 \dots \lceil \frac{n}{2} \rceil - 1]$ )
5: right = MergeSort( $A[\lceil \frac{n}{2} \rceil \dots n - 1]$ )
6: return Merge(left, right)

```

#### Algorithm 3 Merge

*L, R, two sorted arrays input*

**Merged sorted array output**

```

1:  $l = 0, r = 0, i = 0$ 
2: while  $l < a$  and  $r < b$  do
3:   if  $L[l] < R[r]$  then
4:     output[i] =  $L[l]$ 
5:      $l += 1$ 
6:   else
7:     output[i] =  $R[r]$ 
8:      $r += 1$ 
9:   end if
10:   $i += 1$ 
11: end while
12: if  $l = a$  then
13:   output.append( $R[r \dots b]$ )
14: else
15:   output.append( $L[l \dots a]$ )
16: end if
17: return state

```

However, how do we prove that such an algorithm is correct?

**Lemma 6.** Merge( $L, R$ ) correctly merges  $L$  and  $R$  such that the output is sorted if  $L$  and  $R$  are sorted.

**Proof.** We wish to show that after the  $i$ -th (1 indexed) iteration of the while loop, output[1... $i$ ] is sorted. We will proceed with induction on  $i$ .

**Base case:**  $i = 0$ . In this case, nothing has occurred yet, and nothing is in the output, so the output is sorted.

**Inductive step:** By the inductive hypothesis, output[0... $i - 1$ ] is sorted. Let  $l, r$  be the value of variables  $l, r$  at the start of the  $i$ -th iteration. Note that output[ $i - 1$ ] is

either

$$\left\{ \begin{array}{l} L[l-1] = \min(L[l-1], R[r]) \\ \quad \leq \min(L[l], R[r]) \\ \quad = \text{output}[i] \text{ if prev. was } L \\ R[r-1] = \min(L[l], R[r-1]) \\ \quad \leq \min(L[l], R[r]) \\ \quad = \text{output}[i] \text{ if prev. was } R \end{array} \right.$$

which means that  $\text{output}[i-1] \leq \text{output}[i]$ , which means that  $\text{output}[0 \dots i]$  is also sorted. After the while loop,  $\text{output}[i-1]$  is either

$$\left\{ \begin{array}{l} L[a] \leq R[r] \\ \quad \Rightarrow \text{out.append}(R[r \dots b]) \text{ sorted} \\ R[b] \leq L[l] \\ \quad \Rightarrow \text{out.append}(R[l \dots a]) \text{ sorted} \end{array} \right.$$

Therefore, the output is a sorted array. ■

**Theorem 1.** Merge sort is correct.

**Proof.** We will process with induction on  $n$ .

**Base case:**  $n = 1$ . Then, the array  $A$  is just one element, such that  $A$  is sorted. Merge-Sort returns  $A$ , so it returns a sorted array.

**Inductive step:** We assume that merge sort works for all arrays with size  $< n$ . Let  $A$  be an array of length  $n$ . Note that  $A[0 \dots \lceil \frac{n}{2} \rceil - 1]$  and  $A[\lceil \frac{n}{2} \rceil \dots n - 1]$  are both shorter arrays of length  $< n$ . Therefore, by the inductive hypothesis, we know these two subarrays must be sorted. Since merge is correct, the output must also be sorted. ■

Now, let us argue about the time complexity of merge sort.

**Theorem 2.** Merge sort has time complexity  $n \log n$ .

**Proof.** Let  $T(n)$  be the time complexity for sorting an  $n$ -length array. Note that  $T(n) = 2 \cdot T(\frac{n}{2}) + O(n)$ . This is because merge is linear. We have the base case of  $T(1) = 1$ .

Let's look at the recursion tree. At the  $i$ -th level, there are  $2^i$  calls to merge sort, and the input has length  $\frac{n}{2^i}$ . Each merge takes  $O(\frac{n}{2^i})$

time, which implies that all merges performed at level  $i$  take  $2^i \cdot O(\frac{n}{2^i}) = O(n)$  time.

Because we have  $\log_2(n)$  levels, and each level takes  $O(n)$  time, then merge sort takes  $O(n \log n)$  time. ■

**Theorem 3.** In the general case, with  $a$  recursive calls and each subproblem of length  $\frac{n}{b}$ , and  $O(n^c)$  time to combine solutions, we have  $T(n) = a \cdot T(\frac{n}{b}) + O(n^c)$ .

**Proof.** Then, the total time for level  $i$  is  $O(a^i \cdot (\frac{n}{b^i})^c)$ , and the total time is  $O(\sum_{i=0}^{n^{\log_b(n)}} a^i (\frac{n}{b^i})^c)$ , which has three cases.

$$\text{Total Time} = \begin{cases} O(n^{\log_b(a)}), & \text{if } a > b^c \\ O(n^c \log(n)), & \text{if } a = b^c \\ O(n^c), & \text{if } a < b^c \end{cases}$$

## Lecture 3: Divide and Conquer 2

One general approach that could be used for divide and conquer is, first, split the input into equal parts, assume you have solved the subparts, and then answer the question: how do we merge solutions? Note that sometimes we can improve the complexity of merge as well.

**Example.** Given an array  $A[0 \dots n-1]$ , find the  $\max A[i] - A[j]$  such that  $i < j$ .

**Explanation.** The naive iterative solution is iterating over all pairs  $i, j$  and checking the difference. Otherwise, we can use a divide and conquer solution given below.

```
def max_dif(A):
    if n == 1:
        return float('-inf')
    left = max_dif(A[:len(A)/2])
    right = max_dif(A[len(A)/2 + 1:])
    return max(left, right, max(left) - max(right))
```

Then, the complexity of this is  $T(n) = 2 \cdot T(\frac{n}{2}) + O(n) = O(n \log n)$ . Note that we can optimize this solution by optimizing the linear time merge to constant time. Instead of returning only the solution, we can return as well the maximum and minimum element of each of the arrays.

```
def max_dif(l, r, A):
    if l == r:
        return float('-inf'), A[l], A[l]
    m = (l + r) // 2
    left, min_l, max_l = max_dif(A[l:m])
```

```

right, min_r, max_r = max_dif(A[m + 1:r
])
return max(left, right, max_l - min_r),
min(min_l, min_r), max(max_l,
max_r)

```

## 2.1 Karatsuba Multiplication

Multiplication is not actually  $O(n)$  - it is  $O(n^2)$ . This is because addition takes a lot longer as numbers  $n$  grow bigger. This is the algorithm used by the python interpreter.

Consider a number  $n$  bits long. Then, we can write  $A = 2^{\frac{n}{2}} \cdot A_1 + A_2$  and  $B = 2^{\frac{n}{2}} \cdot B_1 + B_2$ . Then,

$$A \cdot B = 2^n A_1 \cdot B_1 + 2^{\frac{n}{2}} (A_1 B_2 + B_1 A_2) + A_2 B_2.$$

This multiplication only contains 4 multiplications of length (bits)  $\frac{n}{2}$ , which is faster (?). Using the Master Theorem to calculate, we only get an algorithm of  $O(n^2)$  time.

We can continue to optimize by instead multiplying with the fact that

$$(A_1 + A_2) \cdot (B_1 + B_2) = A_1 B_2 + A_1 B_2 + B_1 A_2 + A_2 B_2.$$

Let  $x$  be this value. Let  $y = A_1 \cdot B_1$  and  $z = A_2 \cdot B_2$ . Then, we can instead return

$$2^n y + 2^{\frac{n}{2}} (x - y - z) + z.$$

Then, by the master theorem with merge (addition and subtraction) of  $O(n)$ , this algorithm has a runtime of  $O(n^{\log_2(3)})$ .

## Lecture 4: Divide and Conquer 3

Note that arithmetic will take  $O(1)$  unless numbers blow up (grow exponentially).

Today we will cover a different version of divide and conquer. One example of this version is binary search.

```

def bin_search(arr, k):
    if n == 1:
        if arr[0] == k:
            return arr[0]
        else:
            raise Exception("")
    if a[n/2] < k:
        return bin_search(arr[n/2+1:],
            k)
    else:
        return bin_search(arr[:n/2], k)

```

By the master theorem, we have  $a = 1, b = 2, c = 0$  such that binary search runs in  $O(\log n)$  time. Another similar algorithm to binary search in which we only recurse into one spot is finding the  $k$ -th largest element in array, with a quickselect-type algorithm.

```

def k_smallest(arr, k):
    p = A[rand()]
    smaller = []
    larger = []
    for i in range(n):
        if arr[i] <= p:
            smaller.append(arr[i])
        else:
            larger.append(arr[i])
    if smaller == k:
        return p
    if smaller < k:
        return k_smallest(larger, k - len(
            smaller))
    if smaller > k:
        return k_smallest(smaller, k)

```

If we can guarantee a good pivot  $\frac{3}{10}n \leq \text{smaller} \leq \frac{7}{10}n$ , then the number of recursions is at most  $\log_{\frac{10}{7}}(n) = O(\log n)$ . We can guarantee these goods pivots with the median of medians algorithm.

First, we split our array  $A$  into  $\frac{n}{5}$  groups, each of size 5. We can then sort each of these elements in "constant" time. Then, the median of each group is in the middle index of each 5. Now, we return the median of the array containing all medians. We can do this with recursion (divide and conquer), by calling `ksmallest` on it again.

We claim that the median of medians is a good  $p$ . Imagine the upper left rectangular part of the median array. This is less than or equal to the median of medians. Same with the lower right rectangular part. Note that the number of elements in the rectangular parts are both  $\frac{3}{10}$ . Therefore,  $p$  is at least the  $\frac{3}{10}$ -th element (same with the upper side).

Note that the median of medians algorithm takes constant time because sorting each of the 5 arrays is constant, so the sorting is linear. Then, selecting the median of the medians is also linear time, by a recursive call. We have a runtime of

$$T(n) + T\left(\frac{n}{5}\right) + O(n) + T\left(\frac{7}{10}n\right) = O(n).$$

Note that somehow, if we split into 3 groups, then the runtime is  $n \log n$ . However, splitting into any odd number greater than 3 works.

## Lecture 5: Review for Exam 1

**Theorem 4.** Median of medians of  $O(n)$ .

**Proof.** We wish to show that

$$T(n) = T\left(\frac{n}{5}\right) + O(n) + T\left(\frac{7}{10}n\right) \leq C \cdot n.$$

We proceed by induction. We know that

$$T(n) \leq T\left(\frac{n}{5}\right) + T\left(\frac{7}{10}n\right) + d \cdot n.$$

by definition of  $O(n)$ .

Assume  $T(n') \leq C \cdot n'$  for all  $n' < n$ . Therefore,

$$T(n) \leq C \cdot \frac{n}{5} + C \cdot \frac{7n}{10} + d \cdot n.$$

Simplifying, we have

$$T(n) \leq C \cdot n \cdot \left( \frac{2}{10} + \frac{7}{10} \right) + d \cdot n.$$

We pick  $C = d \cdot 10$  such that

$$T(n) \leq C \cdot n.$$

as desired. Note that for the trivial base case, we know that this algorithm for  $n = 1$  takes constant time. ■

## Lecture 6: Graphs, DFS, Topological Order

### 3 Graphs

Insert every class' graph definitions here.

Two common representations of graphs are adjacency matrices, adjacency lists, and edge lists.

**Definition 7.** An **adjacency matrix** is a matrix where

$$A_{ij} = \begin{cases} 1 & \text{if there is an edge between } i, j \\ 0 & \text{otherwise} \end{cases}.$$

However, an adjacency list uses too much space, so an adjacency list can be used, which is at most  $O(V + E)$  space. This list stores only direct neighbors.

#### 3.1 DFS

Yeah, it's just DFS. At a current node, go to the next node. After you've finished exploring everything from that node, come back to this node and go to next node after. Keep a visited array to not visit vertices more than once.

**Lemma 7.**  $\text{DFS}(v)$  will visit every vertex that can be reached from  $v$  via some path.

**Proof.** By contradiction. Assume that there is a path  $v, v_1, v_2, \dots, v_k$ . Assume  $v_k$  is not visited by the call to  $\text{DFS}(v)$ . Let  $v_i$  be the last vertex that did get visited. That means  $\text{DFS}(v_i)$  must have been called. However, that means the edge between  $v_i$  and  $v_{i+1}$  must have been looked

at. The only way then that  $\text{DFS}(v_{i+1})$  doesn't get called, is if it is already visited. However, that means that  $\text{DFS}(v_{i+1})$  must have been called. This is a contradiction, as then  $v_{i+1}$  must be the last visited vertex. ■

**Definition 8.** A graph  $G$  is a **directed acyclic graph** if it is a directed graph without cycles.

You can think of directed acyclic graphs as dependency graphs. What happens if we have a cycle? Bad things. How can we find these cycles? We can run DFS, and if we explore a vertex that is already in the current path, then we know there is a cycle. We can use backtracking to extract this cycle, if necessary.

#### 3.2 Topological Sort

We can order the vertices of our graph topologically, so that all dependencies are processed before the nodes that require them. This is just a dfs where we only recurse once all vertex indegrees have been processed already.

## Lecture 7: BFS, Dijkstra

### 3.3 BFS

Note that BFS is just DFS, but instead of processing the most recent node explored, we process the latest node explored. To find shortest paths, we can track the parent of each node, and go backwards.

**Theorem 5.** BFS finds the shortest path.

**Proof.** Let  $v_i$  be the vertices with distance  $i$  to the start point of BFS. We wish to show that BFS visits all vertices from  $v_i$  before visiting a vertex from  $v_{i+1}$ . We proceed by induction on  $i$ .

**Base case:**  $i = 0$ . Then,  $v_0$  is just the start point which is visited first.

**Step case:**  $i \rightarrow i + 1$ . Let's look at BFS right after we finished visiting all vertices  $v_i$ . Note that no vertices of distance  $v_{i+1}$  have been visited, which implies that no vertices of distance  $v_{i+2}$  has been enqueued, and all vertices from  $v_{i+1}$  are currently in the queue. Since we use a FIFO queue all of  $v_{i+1}$  must be visited before a vertex from  $v_{i+2}$ . Furthermore, the edge we use to traverse to  $v_{i+1}$  requires that an edge  $(u, v_{i+1})$  such that  $u \in v_i$ . ■

However, what happens if we want to find the shortest path in a graph where there are edge weights?

### 3.4 Dijkstra

Instead of a first in first out queue, we use a priority queue. This allows us to process vertices in a way such that the cost to travel to that vertex is less. This is called Dijkstra's, and takes  $O(|E| \log |E|)$  time.

**Theorem 6.** Dijkstra is optimal.

**Proof.** Let  $d(v)$  be the length of the shortest path from  $s$  to  $v$  for all  $v \in V$ , where  $s$  is the start vertex. Assume the algorithm stores an incorrect value in the distance array. Let  $v_1, v_2, \dots, v_n$  be the vertices in the order in which they were visited by Dijkstra. Let  $v_i$  be the first vertex where  $d(v_i)$  is wrong. That means we used the wrong edge to go from  $v_{i-1}$  to  $v_i$ . This means that

$$d(v_i) = d(v_{i-1}) + w_{v_{i-1}v_i} > \text{shortest path to } v_i.$$

Let  $a$  be a vertex with guaranteed correct distance, and  $b$  one otherwise. Assume that this path with  $a, b$  is the correct shortest path from  $s$  to  $v_i$ . However,  $v_i$  was popped from the queue before  $b$ . This means that the shortest path to  $b$  is  $\geq$  the shortest path to  $v_i$ . With some algebraic manipulation as well as asserting that there are no negative edge weights (check the lecture notes), then we have that the shortest path to  $v_i$  is strictly greater than the shortest path to  $v_i$ , a contradiction. Therefore, there cannot exist such a  $v_i$ . ■

distance + weight, we will get the **minimum spanning tree**. This algorithm is known as Prim's algorithm.

**Theorem 7.** Prim's algorithm finds an MST.

**Proof.** Let  $e_1, e_2, \dots, e_n - 1$  be the edges used by Prim, by the order in which Prim uses these edges.  $e_1, \dots, e_{i-1}$  are combined in some MST, but  $e_i$  is the first edge such that  $e_1, \dots, e_i$  is not combined in any MST.

Then, there is some MST  $U$  that uses  $e_1, \dots, e_{i-1}$ . This  $U$  must have an edge connecting the inside and outside that doesn't use  $e_i$ . Since the graph must be connected, there must now be a cycle.

However, because Prim decided to use this edge  $ab$  as well as  $e_i$ , which means that  $w_{ab} > w_{e_i}$ . Then,  $U + e_i - ab$  is a spanning tree of smaller weight, which means that  $U$  was not an MST, which is a contradiction. ■

## Lecture 8: Prim's and Kruskal

Note that if we run DFS, BFS, and Dijkstra, we get trees based on which edges we used to traverse the graph. Dijkstra's is called the shortest path tree. These trees are also known as **spanning trees**.

**Definition 9.** A **spanning tree** is a tree that reaches all nodes of the graph using only edges of that graph.

### 3.5 Prim's

Note that if we replace the priority in Dijkstra with just the weight of the edges instead of the current