

1 Greedy Algorithms

Suppose we want to solve a problem, and we're able to come up with some recursive formulation of the problem that would give us a nice dynamic programming algorithm. But then, upon further inspection, we notice that any optimal solution only depends on looking up the optimal solution to one other subproblem. A greedy algorithm is an algorithm which exploits such a structure, ignoring other possible choices. Greedy algorithms can be seen as a refinement of dynamic programming; in order to prove that a greedy algorithm is correct, we must prove that to compute an entry in our table, it is sufficient to consider at most one other table entry; that is, at each point in the algorithm, we can make a "greedy", locally-optimal choice, and guarantee that a globally-optimal solution still exists. Instead of considering multiple choices to solve a subproblem, greedy algorithms only consider a single subproblem, so they run extremely quickly – generally, linear or close-to-linear in the problem size.

Unfortunately, greedy algorithms do not always give the optimal solution, but they frequently give good (approximate) solutions. To give a correct greedy algorithm one must first identify optimal substructure (as in dynamic programming), and then argue that at each step, you only need to consider one subproblem. That is, even though there may be many possible subproblems to recurse on, given our selection of subproblem, there is always an optimal solution that contains the optimal solution to the selected subproblem.

1.1 Activity Selection Problem

One problem, which has a very nice (correct) greedy algorithm, is the Activity Selection Problem. In this problem, we have a number of activities. Your goal is to choose a subset of the activities to participate in. Each activity has a start time and end time, and you can't participate in multiple activities at once. Thus, given n activities a_1, a_2, \dots, a_n where a_i has start time s_i and finish time f_i , we want to find a maximum set of non-conflicting activities.

The activity selection problem has many applications, most notably in scheduling jobs to run on a single machine.

1.1.1 Optimal Substructure

Let's start by considering a subset of the activities. In particular, we'll be interested in considering the set of activities $S_{i,j}$ that start after activity a_i finishes and end before activity a_j starts. That is, $S_{i,j} = \{a_k \mid f_i \leq s_k, f_k \leq s_j\}$. We can participate in these activities between a_i and a_j . Let $A_{i,j}$ be a maximum subset of non-conflicting activities from the subset $S_{i,j}$. Our first intuition would be to approach this by using dynamic programming. Suppose some $a_k \in A_{i,j}$, then we can break down the optimal subsolution $A_{i,j}$ as follows

$$|A_{i,j}| = 1 + |A_{i,k}| + |A_{k,j}|$$

where $A_{i,k}$ is the best set for $S_{i,k}$ (before a_k), and $A_{k,j}$ is the best set for after a_k . Another way of interpreting this expression is to say "once we place a_k in our optimal set, we can only consider optimal solutions to subproblems that do not conflict with a_k ."

Thus, we can immediately come up with a recurrence that allows us to come up with a dynamic programming algorithm to solve the problem.

$$|A_{i,j}| = \max_{a_k \in S_{i,j}} 1 + |A_{i,k}| + |A_{k,j}|$$

This problem requires us to fill in a table of size n^2 , so the dynamic programming algorithm will run in $\Omega(n^2)$ time. The actual runtime is $O(n^3)$ since filling in a single entry might take $O(n)$ time.

But we can do better! We will show that we only need to consider the a_k with the smallest finishing time, which immediately allows us to search for the optimal activity selection in linear time.

Claim 1. *For each $S_{i,j}$, there is an optimal solution $A_{i,j}$ containing $a_k \in S_{i,j}$ of minimum finishing time f_k .*

Note that if the claim is true, when f_k is minimum, then $A_{i,k}$ is empty, as no activities can finish before a_k ; thus, our optimal solution only depends on one other subproblem $A_{k,j}$ (giving us a linear time algorithm).

Here, we prove the claim.

Proof. Let a_k be the activity of minimum finishing time in $S_{i,j}$. Let $A_{i,j}$ be some maximum set of non-conflicting activities. Consider $A'_{i,j} = A_{i,j} \setminus \{a_l\} \cup \{a_k\}$ where a_l is the activity of minimum finishing time in $A_{i,j}$. It's clear that $|A'_{i,j}| = |A_{i,j}|$. We need to show that $A'_{i,j}$ does not have conflicting activities. We know $a_l \in A_{i,j} \subset S_{i,j}$. This implies $f_l \geq f_k$, since a_k has the minimum finishing time in $S_{i,j}$.

All $a_t \in A_{i,j} \setminus \{a_l\}$ don't conflict with a_l , which means that $s_t \geq f_l$, which means that $s_t \geq f_k$, so this means that no activity in $A_{i,j} \setminus \{a_l\}$ can conflict with a_k . Thus, $A'_{i,j}$ is an optimal solution. \square

Due to the above claim, the expression for $A_{i,j}$ from before simplifies to the following expression in terms of $a_k \subseteq S_{i,j}$, the activity with minimum finishing time f_k .

$$\begin{aligned} |A_{i,j}| &= 1 + |A_{k,j}| \\ A_{i,j} &= A_{k,j} \cup \{a_k\} \end{aligned}$$

Algorithm Greedy-AS assumes that the activities are presorted in nondecreasing order of their finishing time, so that if $i < j$, $f_i \leq f_j$.

Algorithm 1: Greedy-AS(a)

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A ← {a1} // activity of min fi
k ← 1
for m = 2 → n do
    if sm ≥ fk then
        // am starts after last activity in A
        A ← A ∪ {am}
        k ← m
return A

```

By the above claim, this algorithm will produce a legal, optimal solution via a greedy selection of activities. There may be multiple optimal solutions, but there always exists a solution that includes a_k with the minimum finishing time. The algorithm does a single pass over the activities, and thus only requires $O(n)$ time – a dramatic improvement from the trivial dynamic programming solution. If the algorithm also needed to sort the activities by f_i , then its runtime would be $O(n \log n)$ which is still better than the original dynamic programming solution.

1.2 Scheduling

Consider another problem that can be solved greedily. We are given n jobs which all need a common resource. Let w_j be the weight (or importance) and l_j be the length (time required) of job j . Our output is an ordering of jobs. We define the completion time c_j of job j to be the sum of the lengths of jobs in the ordering up to and including l_j . Our goal is to output an ordering of jobs that minimizes the weighted sum of completion times $\sum_j w_j c_j$.

1.2.1 Intuition

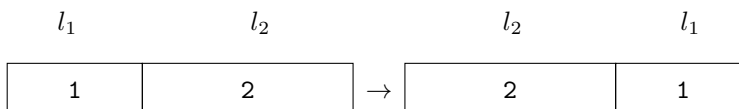
Our intuition tells us that if all jobs have the same length, then we prefer larger weighted jobs to appear earlier in the order. If jobs all have equal weights, then we prefer shorter length jobs in the order.



In the first case, assuming they all have equal weights of 1, $\sum_{i=1}^3 w_i c_i = 1 + 3 + 6 = 10$. In the second case, $\sum_{i=1}^3 w_i c_i = 3 + 5 + 6 = 14$.

1.2.2 Optimal Substructure

What do we do in the cases where $l_i < l_j$ and $w_i < w_j$? Consider the optimal ordering of jobs. Suppose we have a job i that is followed by job j in the optimal order. Consider swapping jobs i and j . The example below swaps jobs 1 and 2.



Note that swapping jobs i and j does not alter the completion times for every other job and only changes the completion times for i and j . c_i increases by l_j and c_j decreases by l_i . This means that our objective function $\sum_i w_i c_i$ changes by $w_i l_j - w_j l_i$. Since we assumed our order was optimal originally, our objective function cannot decrease after swapping the jobs. This means,

$$w_i l_j - w_j l_i \geq 0$$

which implies,

$$\frac{l_j}{w_j} \geq \frac{l_i}{w_i}$$

Therefore, we want to process jobs in increasing order of $\frac{l_i}{w_i}$, the ratio of the length to the weight of each job. The algorithm also does a single pass over jobs, and thus only requires $O(n)$ time, assuming the jobs were ordered by $\frac{l_i}{w_i}$. Like previously, if the algorithm also needed to sort the jobs based on the ratio of length to weight, then its runtime would be $O(n \log n)$.

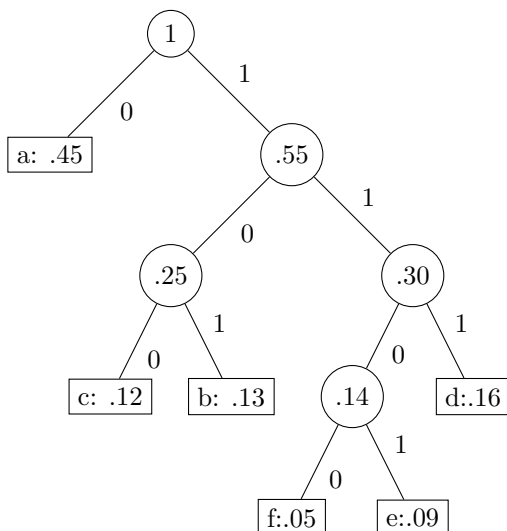
1.3 Optimal Codes

Our third example comes from the field of information theory. In ASCII, there is a fixed 8 bit code for each character. Suppose we want to incorporate information about frequencies of characters to obtain shorter encodings. What if we want to represent characters by codes of different lengths depending on each character's frequencies? We explore a greedy solution to find the optimal encoding of characters.

To create optimal codes, we want a way to encode and decode our sequence. To encode the sequence, we would just have to concatenate the code of each character together. How about for decoding? Consider the following codes of characters: $a \rightarrow 0, b \rightarrow 1, c \rightarrow 01$. However, when decoding, when we encounter 01, this could be decoded as "ab" or "c". Therefore, our codes need to be **prefix free**: no codeword is a prefix of another.

1.3.1 Tree Representation

We may think of representing our codes in a tree structure, where the codewords represent the leaves of our tree. An example is shown below:



Above, in addition to the characters $\{a, b, c, d, e, f\}$, we've included *frequency information*. That is, $f(a) = 0.45$ means that the probability of a random character in this language being equal to a is $.45$.

The code for each character can be found by concatenating the bits of the path from the root to the leaves. By convention, every left branch is given the bit 0 and every right branch is given the bit 1.

As long as the characters are on the leaves of this tree, the corresponding code will be prefix-free. This is because one string is a prefix of another if and only if the node corresponding to the first is an ancestor of the node corresponding to the second. No leaf is an ancestor of any other leaf, so the code is prefix-free.

1.3.2 How good is a code?

Suppose we have a set of characters C with frequencies $f(c)$ so that $\sum_{c \in C} f(c) = 1$. That is, $f(c)$ can be thought of as the probability of using a letter c in this language. The *cost*, in terms of bits, of a character $c \in C$ when using the coding scheme represented by a tree T is just the depth in the tree T : $\text{cost}(c) = d_T(c)$. For example, in the tree above, e has depth 4 in the tree, and requires 4 bits to represent. The average cost of the tree is

$$B(T) = \mathbb{E}_{c \in C} d_T(c) = \sum_{c \in C} f(c) d_T(c).$$

We say that a tree T is *optimal* if this expected cost $B(T)$ is as small as possible.

1.3.3 Huffman Codes

In 1951, David A. Huffman, in his MIT information theory class, was given the choice of a term paper or final exam. Huffman chose to do the term paper rather than take the final exam. He found greedy algorithm to find the most efficient binary code, which we know today as **Huffman codes**.

The basic idea is this: build subtrees for subsets of characters and merge them from the bottom up, combining the two trees with the characters of minimum total frequency.

Input: Set of characters $C = \{c_1, c_2, \dots, c_n\}$ of size n , and $F = \{f(c_1), f(c_2), \dots, f(c_n)\}$, a set of frequencies.

- Create nodes N_k for each character c_k , with key $f(c_k)$.
- Let **current** denote the set $\{N_1, \dots, N_n\}$ of nodes.
- while **current** has length more than one:
 - Find the two nodes N_i and N_j in **current** with the minimum frequencies and create a new intermediate node I with N_i and N_j as its children, so that $I.key = N_i.key + N_j.key$.
 - Add I to **current** and remove N_i, N_j .
- Return the only entry of **current**, which is the root of the tree.

Figure 1: A high-level version of the HUFFMAN CODING algorithm.

The tree shown above results from running this algorithm on the letters with those frequencies; see the slides for an illustration of this process.

1.3.4 Proof of Correctness

This algorithm works, but at first it's not at all obvious why. For a rigorous proof, refer to Lemmas 16.2 and 16.3 in CLRS. However, we'll sketch the idea below. Formally, the proof goes by induction. Recall that after iteration t in Algorithm 1, we have a list **current**, which contains the roots of subtrees that we still need to merge up. We will maintain the following inductive hypothesis:

- **Inductive Hypothesis:** Suppose we have completed t iterations of the loop in Algorithm 1. Then there exists a way to merge the subtrees in **current** that is optimal.
- For the **base case**, we observe that when $t = 0$, **current** is just the set of all characters, and definitionally there exists an optimal tree made out of these nodes.
- For the **inductive step**, we need to show that if the inductive hypothesis holds at step $t - 1$, then it holds at step t . We'll sketch this later.
- Finally, to **conclude** the argument, we see that at the end of the algorithm, there is only one element in **current**, and in this case the inductive hypothesis reads that there is a way to merge this single subtree to obtain an optimal subtree. That's just a convoluted way of saying that the single tree we return is optimal, and so we are done.

All that remains to show is the inductive step. We first observe the following claim:

Claim 2. *We are given a set of characters C and a set of its associated frequencies F where $f(c)$ is the frequency of character c . Let x and y be the characters with the two smallest frequencies. There exists an optimal coding tree for C such that x, y are sibling leaves.*

Proof. Let T be the optimal coding tree for C . The optimal coding tree must be a full binary tree, that is, every non-leaf node must have two children. Let a, b be characters that are sibling leaves of maximum depth. We define the number of bits to encode c as $d_T(c)$ and the number of bits needed for the coding tree as $B(T) = \sum_c f(c)d_T(c)$.

We can replace a, b by x, y without increasing the total number of bits needed for the coding tree.¹ If we swap x and a , the change in cost becomes

$$f(x)d_T(a) + f(a)d_T(x) - f(x)d_T(x) - f(a)d_T(a) = (f(x) - f(a))(d_T(a) - d_T(x)) \leq 0$$

¹For simplicity, we ignore the case where a, b, x, y are not distinct. For more details, see Lemma 16.2 in CLRS.

Therefore, there swapping a, b with x, y will not increase our objective function $B(T)$. Hence, there exists an optimal coding tree where x, y are siblings in the tree. \square

Claim 2 shows that there exists an optimal coding tree where x and y are sibling leaves, that is, there is an optimal code that makes the same greedy choice as the algorithm. However, this is only immediately helpful for the first iteration of the inductive step, when all of the elements of **current** are indeed leaves. In order to make this idea work for all t , we need one more claim.

Claim 3. *Let C be a set of characters, and let T be a coding tree for C . Imagine creating C' from C by collapsing all the characters in a subtree rooted at a node N with key $k = N.\text{key}$ into a single character c' with frequency k . Then the corresponding tree T' is optimal for C' .*

Conversely, suppose that a tree T' that is an optimal coding tree for an alphabet C' . Let $c' \in C'$ be a character with frequency $f(c')$. Introduce new characters c''_1, \dots, c''_r with total frequency $\sum_{i=1}^r f(c''_i) = f(c')$. Let T'' be an optimal coding tree on c''_1, \dots, c''_r . Then the tree T on the alphabet $C = (C' \setminus \{c'\}) \cup \{c''_1, \dots, c''_r\}$ that has the leaf c' replaced with the subtree T'' is optimal.

Proof. Let T and T' be the two trees described in the lemma, and consider the difference of their costs.

$$\begin{aligned}
B(T) - B(T') &= \sum_{c \in C} f(c) \cdot d_T(c) - \sum_{c \in C'} f(c) \cdot d_{T'}(c) \\
&= \left(\sum_{i=1}^r f(c''_i) d_T(c''_i) \right) - f(c') d_{T'}(c') \\
&= \left(\sum_{i=1}^r f(c''_i) (d_{T''}(c''_i) + d_{T'}(c')) \right) - f(c') d_{T'}(c') \\
&= \sum_{i=1}^r f(c''_i) d_{T''}(c''_i) + d_{T'}(c') \sum_{i=1}^r f(c''_i) - f(c') d_{T'}(c') \\
&= \sum_{i=1}^t f(c''_i) d_{T''}(c''_i)
\end{aligned}$$

where the last line used the fact that $\sum_{i=1}^r f(c''_i) = f(c')$, and so the last two terms cancelled. This means that the difference in the cost between these two trees *only* depends on T'' , it doesn't depend at all about the structure of T . Thus, T is optimal if and only if T' is optimal. \square

The two Claims together prove the inductive step, because the second claim implies that the logic of the first claim holds, even for newly created intermediate nodes I .

Note: The proof in CLRS has the same basic steps (Lemmas 16.2 and 16.3 instead of the claims above), although phrased slightly differently. The sketch above is pretty sketchy, so if the above is hard to follow, please check out CLRS for a more detailed version.