

COMS4231: Analysis of Algorithms

Fall 19

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Last time

- Lower bound for sorting based on comparisons: $\Omega(n \log n)$
- Sorting beyond comparisons, in $O(n)$ time
 - Count Sort, Bucket Sort
 - Radix Sort

Today

- Dynamic Programming (new technique)
 - Fibonacci
 - 0-1 Knapsack
 - Longest Common Subsequence

Dynamic Programming

Reduce problem to smaller problems

like D&C but possibly *overlapping subproblems*

Memoization:

do not solve the same problem instance repeatedly,
solve it once and record the result to reuse it (if needed)

Bottom-up (iterative) version: Problems are solved from
smaller to larger and solutions tabulated

Top-down (recursive) version: Before initiating **recursive**
call, **check if solution was already computed** previously.

Example: Fibonacci numbers

- $F_0 = 0, F_1 = 1$
- $F_n = F_{n-1} + F_{n-2}$ for $n \geq 2$
- Sequence: 0, 1, 1, 2, 3, 5, 8, 13, 21,...

Recursive algorithm **FIB**(n):

if $n = 0$ **then return** 0

else if $n = 1$ **then return** 1

else return FIB($n - 1$) + FIB($n - 2$)

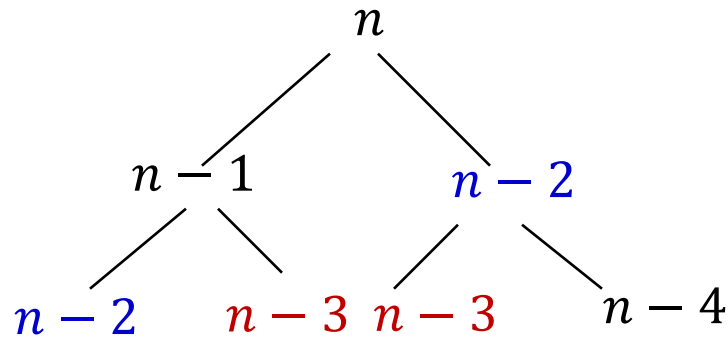
Complexity: $T(n) = T(n - 1) + T(n - 2) + O(1)$

Grows at least as fast as the Fibonacci numbers

$$F_n \approx \varphi^n / \sqrt{5}, \text{ where } \varphi = (1 + \sqrt{5})/2 = \text{golden ratio}$$

Fibonacci numbers ctd.

- **FIB** is called only on n distinct arguments, but it is called repeatedly with the **same arguments**



Tabulate the result, so no need to evaluate it again on the same argument \Rightarrow complexity $O(n)$

New algorithm **FIB**(n):

$F[0] = 0; F[1] = 1;$

for $i = 2$ **to** n **do**

$F[i] = F[i - 1] + F[i - 2]$

return $F[n]$

(Actually, in this case we don't need an array: only need to remember the last two values)

often for Optimization Problems

Optimization Problem:

- For given instance of the problem, there is a set of “feasible” solutions, involving a number of choices (or decisions)
- Every solution has a cost or a value: metric for evaluating solutions
- Goal: find an optimal solution:
 - min-cost
 - or max-value

Dynamic Programming for Optimization Problems

Main principles:

1) Optimal Substructure:

Problem can be reduced to a set of smaller subproblems;
Optimal solution for whole involves optimal solutions for subproblems

2) Memoization:

Subproblems are solved from smaller to larger and solutions tabulated

Problem: 0-1 Knapsack

- n items, with given integer weights w_i , values $v_i, i = 1, \dots, n$
- Knapsack with weight capacity W
- **Goal:** Choose a subset of items that fits in the knapsack and has **maximum value**
i.e, choose a subset $S \subseteq \{1, \dots, n\}$ that
maximizes $\sum_{i \in S} v_i$
subject to $\sum_{i \in S} w_i \leq W$

Example

Item	Weight	Value	Knapsack capacity: 13
1	4	25	
2	6	30	
3	2	10	
4	5	27	
5	7	35	

Some **feasible solutions**, and their value

$\{1,2,3\}$: 65, $\{1,3,4\}$: 62, $\{1,3,5\}$: 70, $\{2,3,4\}$: 67,
 $\{2,5\}$: 65, $\{4,5\}$: 62,

Optimal solution: $\{1,3,5\}$, value 70

Reduction to smaller subproblems: "last step/item analysis"

Should we take the n^{th} item?

- If we take it:

we have capacity $W - w_n$ left and can pick any subset from the first $n - 1$ items \rightarrow ($n - 1$ size subproblem)

- If we don't take it: we have capacity W for the first $n - 1$ items \rightarrow ($n - 1$ size subproblem)

Overall optimum = the best of these 2 options!

Let $M(b, i)$ = maximum value we can get with knapsack of capacity b , using a subset of the first i items only

$$M(b, i) = \max\{ M(b - w_i, i - 1) + v_i, M(b, i - 1) \} \text{ if } b \geq w_i$$

else $M(b, i) = M(b, i - 1)$

Base case: $M(b, 0) = 0$ for all b

Recursive Algorithm (not D.P.)

$M(b, i) = \max\{ M(b - w_i, i - 1) + v_i, M(b, i - 1) \}$ if $b \geq w_i$
else $M(b, i) = M(b, i - 1)$

Base case: $M(b, 0) = 0$ for all b

Rec-KNAP(w, v, b, i)

[max value that can be obtained for capacity b from first i items only]

if $i = 0$ or $b = 0$ then return 0

else if $b < w_i$ then return Rec-KNAP($w, v, b, i - 1$)

else return $\max\{ \text{Rec-KNAP}(w, v, b - w_i, i - 1) + v_i, \text{Rec-KNAP}(w, v, b, i - 1) \}$

Main call: Rec-KNAP(w, v, W, n)

Time Complexity of Recursive algorithm

- A call for i items may generate two recursive calls with $i - 1$ items.
- $T(i) = 2T(i - 1) + O(1)$
- $\Rightarrow T(n) = \Theta(2^n)$
- But: many of these recursive calls solve the same problem $M(b, i)$:
 - at most nW different arguments.
 - better if $W \ll 2^n$

DP Algorithm (iterative)

DP-KNAP(w, v, W)

for $b = 0$ **to** W **do** $M(b, 0) = 0$

for $i = 1$ **to** n **do**

for $b = 0$ **to** W **do**

if $b \geq w_i$ **and** $M(b - w_i, i - 1) + v_i > M(b, i - 1)$

then $M(b, i) = M(b - w_i, i - 1) + v_i$

else $M(b, i) = M(b, i - 1)$

Return $M(W, n)$

Running time: $O(nW)$

Ok if W is “small”

(strictly, not a *polynomial-time* algorithm if weights given in binary)

DP Algorithm: recursive version

Initialize $M(b,i) = \text{"?"}$ at the beginning for all b,i

Main call: $\text{Rec-KNAP}(w, v, W, n)$

$\text{Rec-KNAP}(w, v, b, i)$

if $i = 0$ **or** $b = 0$ **then return** 0

if $M(b, i) \neq \text{"?"}$ **then return** $M(b, i)$

else if $b < w_i$ **then** $r = \text{Rec-KNAP}(w, v, b, i - 1)$

else $r = \max\{ \text{Rec-KNAP}(w, v, b - w_i, i - 1) + v_i ,$
 $\text{Rec-KNAP}(w, v, b, i - 1) \}$

$M(b, i) = r$

Return r

Has same (worst-case) complexity as the iterative DP algorithm

Recovering an optimal solution

Record which case generates $M(b, i)$ for every b, i

for $b = 0$ **to** W **do** $M(b, 0) = 0$

for $i = 1$ **to** n **do**

for $b = 0$ **to** W **do**

if $b \geq w_i$ **and** $M(b - w_i, i - 1) + v_i > M(b, i - 1)$

then $\{M(b, i) = M(b - w_i, i - 1) + v_i ; s(b, i) = 1\}$

else $\{M(b, i) = M(b, i - 1) ; s(b, i) = 0\}$

Return $M(W, n)$ **and** s

Optimal Solution:

$b = W; S = \emptyset$

for $i = n$ **down to** 1 **do**

if $s(b, i) = 1$ **then** $\{S = S \cup \{i\}; b = b - w_i\}$

return S

$s(b, i) = 1$ iff (an)
optimal solution for
 $M(b, i)$ uses item i

Longest Common Subsequence (LCS)

- Given two sequences $x[1 \dots m]$, $y[1 \dots n]$, find a longest common subsequence (gaps allowed)

$x = A \ T \ C \ T \ T \ A \ G$
 $y = T \ G \ C \ A \ T \ A$

The diagram illustrates the alignment of two sequences, x and y. Sequence x is "A T C T T A G" and sequence y is "T G C A T A". Lines connect the following characters: x[1] (A) to y[1] (T), x[3] (C) to y[3] (C), x[4] (T) to y[5] (T), and x[6] (A) to y[6] (A). This alignment shows a common subsequence of length 4: "A C T A".

- Applications: computational biology, *diff*
- Naive way: Take every subsequence of x check against $y \rightarrow \text{Time: } 2^m n$