

COMS4231: Analysis of Algorithms

Fall 19

Alex Andoni

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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( $n - 2$ )
```

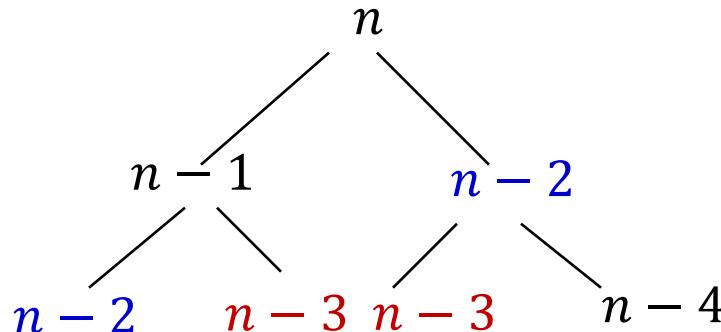
Complexity: $T(n) = T(n - 1) + T(n - 2) + O(1)$

Grows at least as fast as the Fibonacci numbers

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

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

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

Recursive Algorithm (not D.P.)

```
M(b, i) = max{ M(b - wi, i - 1) + vi, M(b, i - 1) } if b ≥ wi
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{ Rec-KNAP($w, v, b - w_i, i - 1$) + v_i ,
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 $M(b,i) = "?"$ at the beginning for all b,i

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

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

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

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

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

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

$$\begin{array}{ccccccc} x & = & A & T & C & T & T & A & G \\ & & \diagup & & | & \diagdown & & | \\ y & = & T & G & C & A & T & A \end{array}$$

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