

Lecture 21: Dynamic Programming III

Lecture Overview

- Subproblems for strings
- Parenthesization
- Edit distance (& longest common subseq.)
- Knapsack
- Pseudopolynomial Time

Review:

- * 5 easy steps to dynamic programming

- | | |
|---|---------------------------------|
| (a) define subproblems | count # subproblems |
| (b) guess (part of solution) | count # choices |
| (c) relate subproblem solutions | compute time/subproblem |
| (d) recurse + memoize | time = time/subproblem · # sub- |
| problems | |
| OR build DP table bottom-up | |
| check subproblems acyclic/topological order | |
| (e) solve original problem: = a subproblem | |
| OR by combining subproblem solutions | ⇒ extra time |

- * problems from L20 (text justification, Blackjack) are on sequences (words, cards)

- * useful problems for strings/sequences x :

suffixes $x[i:]$	} $\Theta(x)$	← cheaper ⇒ use if possible
prefixes $x[:i]$		
substrings $x[i:j]$	} $\Theta(x^2)$	

Parenthesization:

Optimal evaluation of associative expression $A[0] \cdot A[1] \cdots A[n-1]$ — e.g., multiplying rectangular matrices

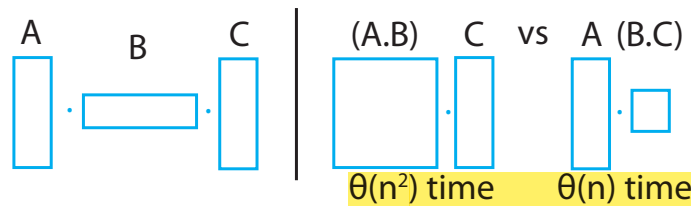


Figure 1:

2. guessing = outermost multiplication $\underbrace{(\dots)}_{\uparrow_{k-1}} \underbrace{(\dots)}_{\uparrow_k}$

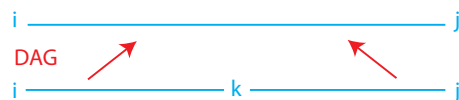
\Rightarrow # choices = $O(n)$

1. subproblems = ~~prefixes & suffixes?~~ **NO**
 = **cost of substring** $A[i:j]$

\Rightarrow # subproblems = $\Theta(n^2)$

3. recurrence:

- $DP[i, j] = \min(DP[i, k] + DP[k, j] + \text{cost of multiplying } (A[i] \cdots A[k-1]) \text{ by } (A[k] \cdots A[j-1]))$ for k in range($i+1, j$)



- $DP[i, i+1] = 0$
 \Rightarrow cost per subproblem = $O(j-i) = O(n)$

4. topological order: increasing substring size. Total time = $O(n^3)$

5. original problem = $DP[0, n]$
 (& use parent pointers to recover parens.)

NOTE: Above DP is not shortest paths in the subproblem DAG! Two dependencies \Rightarrow **not path!**

Edit Distance

Used for DNA comparison, diff, CVS/SVN/..., spellchecking (typos), plagiarism detection, etc.

Given two strings x & y , what is the cheapest possible sequence of character edits (insert c , delete c , replace $c \rightarrow c'$) to transform x into y ?

- cost of edit depends only on characters c, c'
- for example in DNA, $C \rightarrow G$ common mutation \implies low cost
- cost of sequence = sum of costs of edits
- If insert & delete cost 1, replace costs 0, minimum edit distance equivalent to finding longest common subsequence. Note that a subsequence is sequential but not necessarily contiguous.
- for example **H I E R O G L Y P H O L O G Y** vs. **M I C H A E L A N G E L O**
 \implies **HELLO**

Subproblems for multiple strings/sequences

- combine suffix/prefix/substring subproblems
- multiply state spaces
- still polynomial for $O(1)$ strings

Edit Distance DP

(1) subproblems: $c(i, j) = \text{edit-distance}(x[i:], y[j:])$ for $0 \leq i < |x|, 0 \leq j < |y|$
 $\implies \Theta(|x| \cdot |y|)$ subproblems

(2) guess whether, to turn x into y , (3 choices):

- $x[i]$ deleted
- $y[j]$ inserted
- $x[i]$ replaced by $y[j]$

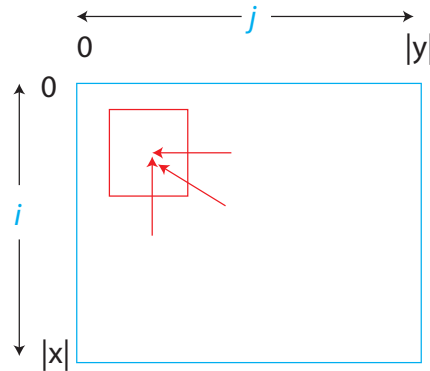
(3) recurrence: $c(i, j) = \text{maximum of}$:

- $\text{cost}(\text{delete } x[i]) + c(i + 1, j)$ if $i < |x|$,
- $\text{cost}(\text{insert } y[j]) + c(i, j + 1)$ if $j < |y|$,
- $\text{cost}(\text{replace } x[i] \rightarrow y[j]) + c(i + 1, j + 1)$ if $i < |x| \& j < |y|$

base case: $c(|x|, |y|) = 0$

$\implies \Theta(1)$ time per subproblem

(4) topological order: DAG in 2D table:



- bottom-up **OR** right to left
- only need to keep last 2 rows/columns
 \implies linear space
- total time = $\Theta(|x| \cdot |y|)$

(5) original problem: $c(0, 0)$

Knapsack:

Knapsack of size S you want to pack

- item i has integer size s_i & real value v_i
- goal: choose subset of items of maximum total value subject to total size $\leq S$

First Attempt:

1. ~~subproblem = value for suffix i~~ : **WRONG**
2. guessing = whether to include item $i \implies \# \text{ choices} = 2$
3. recurrence:
 - $DP[i] = \max(DP[i+1], v_i + DP[i+1] \text{ if } s_i \leq S?)$
 - **not enough information to know whether item i fits — how much space is left?**
GUESS!

Correct:

1. subproblem = value for suffix i :
 given knapsack of size X
 $\implies \# \text{ subproblems} = O(nS)$ **!**

3. recurrence:

- $DP[i, X] = \max(DP[i + 1, X], v_i + DP[i + 1, X - s_i] \text{ if } s_i \leq X)$
- $DP[n, X] = 0$
 \implies time per subproblem $= O(1)$

4. topological order: for i in $n, \dots, 0$: for X in $0, \dots, S$

total time $= O(nS)$

5. original problem $= DP[0, S]$

(& use parent pointers to recover subset)

AMAZING: effectively trying all possible subsets! ...but is this actually fast?

Polynomial time

Polynomial time = polynomial in input size

- here $\Theta(n)$ if number S fits in a word
- $O(n \lg S)$ in general
- S is exponential in $\lg S$ (not polynomial)

Pseudopolynomial Time

Pseudopolynomial time = polynomial in the problem size AND the numbers (here: S, s_i 's, v_i 's) in input. $\Theta(nS)$ is pseudopolynomial.

Remember:
 polynomial — GOOD
 exponential — BAD
 pseudopoly — SO SO

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