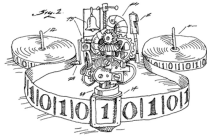


EECS 376: Foundations of Computer Science

Lecture 05 - Dynamic Programming 2



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Dynamic Programming Review

- **Step 1:** Write a **recursive formulation** of the solution. **Bound the number of distinct subproblems** that ever appear in your formulation
- **Step 2:** Create a table representing distinct subproblems. Fill in the table from the **bottom-up**.
- **Runtime:** (#subproblem \times time per subproblem)

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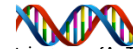
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Longest Common Subsequence

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Motivation: DNA Comparison



- Your DNA is a (*long*) string over $\{A, T, C, G\}$.
- “Humans and chimps are 98.9% similar.”
 - X : ACCGGTCGAGTGCGCGGAAGCCGGCCGAA
 - Y : GTCGTTCCGAATGCCGTTGCTCTGTAA
- The length of the *longest common subsequence* between two genomes is a measure of *similarity*.

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Longest Common Subsequence

- Given strings $X[1..m]$ and $Y[1..n]$
- **Goal:** find the *length* of a **longest common subsequence** of X and Y
 - A **subsequence** of X is a string obtainable from X by deleting chars (may not be consecutive)
 - A **common subsequence** of X and Y is a subsequence of both X and Y
- **Example:**
 - “CT” is a common subsequence of “CGATG” and “CATGT”.
 - **Q:** What’s the longest?
- **Q:** What’s a brute force solution?
 - Each character of X and Y is either deleted or not:
 - Runtime: $O(2^m 2^n)$

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Recurrence for LCS : First, define the function

- **Def:** $LCS(i, j)$ = length of a LCS of $X[1..i]$ and $Y[1..j]$.
 - $i = 0$ means X is the empty string
 - $j = 0$ means Y is the empty string
- **Goal:** $LCS(m, n)$ (compute the length of LCS, will compute LCS itself soon)
- **Example:** Suppose $X = \text{“ATGCC”}$ and $Y = \text{“TAGC”}$.
 - **Q:** What’s $LCS(1, 0)$? 0
 - **Q:** What’s $LCS(5, 3)$? 2 TGC or AG
 - **Q:** What’s $LCS(4, 4)$? 3 TGC or AGC
 - **Q:** What’s $LCS(5, 4)$? 3 TGC or AGC

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Recurrence for LCS

- **Def:** $LCS(i, j)$ = length of a LCS of $X[1..i]$ and $Y[1..j]$.
 - $i = 0$ means X is the empty string
 - $j = 0$ means Y is the empty string
- **Goal:** return $LCS(m, n)$
- What is a recursion for $LCS(i, j)$?

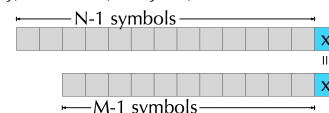
$$LCS(i, j) = \begin{cases} ? & \text{if } i = 0 \text{ or } j = 0 \\ \text{Pause and Think...} & \text{if } X[i] = Y[j] \\ & \text{if } X[i] \neq Y[j] \end{cases}$$

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Recurrence for LCS

- Def:** $LCS(i, j)$ = length of a LCS of $X[1..i]$ and $Y[1..j]$.
 (所有即 sol 中的 char 都 - 排成可以放一起)
 那有某个 optimal sol 中包含
 因为把它 delete 掉 (i, j--)
 length ++
- **Case 1:** $X[i] = Y[j]$ (ends with the same character)
 - **Example:** $X[1..i] = \text{“CTGCA”}$ and $Y[1..j] = \text{“TCGA”}$
 - **Claim.** There exists an optimal solution OPT that matches $X[i]$ and $Y[j]$. 然后看下一个
 - **Proof by contradiction.**
 - Suppose for contradiction: all optimal sol OPT do not match $X[i] = Y[j]$ (Say, $X[i] = \text{“A”}$).
 - OPT + “A” is an LCS too. But OPT + “A” is longer than OPT.
 - So, OPT is not optimal, contradiction.
 - $LCS(i, j) = 1 + LCS(i - 1, j - 1)$



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- $LCS(i, j) = \max\{LCS(i - 1, j), LCS(i, j - 1)\}$

$$LCS(i, j) = \begin{cases} 0 & i = 0 \text{ or } j = 0 \\ 1 + LCS(i-1, j-1) & X[i] = Y[j] \\ ? & X[i] \neq Y[j] \end{cases}$$

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考虑了 X 中的 p 和 27
每个 y 中的 p 分别 match 时的情况

Example

Task: compute LCS of
• "APOCRYPHAL"
• "POLYPEPTIDE"

		A	P	O	C	R	Y	P	H	A	L
A	0	0	0	0	0	0	0	0	0	0	0
P	0	0	1	1	1	1	1	1	1	1	1
O	0	0	1	2							
L	0										
Y	0										
P	0										
E	0										
P	0										
T	0										
I	0										
D	0										
E	0										

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Example

Task: compute LCS of
• "APOCRYPHAL"
• "POLYPEPTIDE"

		A	P	O	C	R	Y	P	H	A	L
A	0	0	0	0	0	0	0	0	0	0	0
P	0	0	1	1	1	1	1	1	1	1	1
O	0	0	1	2	2	2	2	2	2	2	2
L	0	0	1	2	2	2	2	2	2	2	3
Y	0	0	1	2	2	2	3	3	3	3	3
P	0	0	1	2	2	2	3	4	4	4	4
E	0	0	1	2	2	2	3	4	4	4	4
P	0	0	1	2	2	2	3	4	4	4	4
T	0	0	1	2	2	2	3	4	4	4	4
I	0	0	1	2	2	2	3	4	4	4	4
D	0	0	1	2	2	2	3	4	4	4	4
E	0	0	1	2	2	2	3	4	4	4	4

Q: What is the length of LCS?
A: 4

Q: What is the LCS itself?
A: POYP

Exercise: Think how to edit
the program so that it can
return the LCS itself.

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Time and Space Complexity

- How efficient is this algorithm? Computing a single table entry requires a constant number of operations. Since there are $(N+1) \cdot (M+1)$ entries, constructing the table takes $O(NM)$ time and requires $O(NM)$ space.
- Backtracking also does a constant number of operations per entry on the path, and the path length is at most $N + M + 1$, so backtracking takes $O(N + M)$ time.
- Thus, the total complexity of this algorithm is $O(NM)$ in both time and space.

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0-1 KNAPSACK

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0-1 Knapsack Problem

Input:

- n items t_1, \dots, t_n : each with value v_i and weight w_i .
- W : the total capacity of the 0-1 knapsack.
- All v_i, w_i, W are integers.

Output: A set of items $S \subseteq \{1, \dots, n\}$ such that

- (not too heavy) $\sum_{i \in S} w_i \leq W$
- (max value) $\sum_{i \in S} v_i$ as large as possible



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$Knapsack(\{t_1, \dots, t_n\}, W)$

Recursive Strategy:
Take t_n ?

Reject: $val_{rej} = Knapsack(\{t_1, \dots, t_{n-1}\}, W)$

Accept: $val_{acc} = v_n + Knapsack(\{t_1, \dots, t_{n-1}\}, W - w_n)$

- Do not know whether optimal solution actually has t_n .
— Again, try both and take the better one.

The recurrence:

$$Knapsack(\{t_1, \dots, t_n\}, W) = \max \begin{cases} Knapsack(\{t_1, \dots, t_{n-1}\}, W) \\ v_n + Knapsack(\{t_1, \dots, t_{n-1}\}, W - w_n) \end{cases}$$

The base cases:

- $Knapsack(\emptyset, W') = 0$ for all W'
- $Knapsack(\{t_1, \dots, t_i\}, 0) = 0$ for all i

- What is the size of the table?
- What is time spent per cell?

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$$t_1: V_1=3, W_1=1; t_2: V_2=5, W_2=4; t_3: V_3=6, W_3=4$$

$$W=7$$

	$i=0$	$i=1$	$i=2$	$i=3$
$W=0$	0	0	0	0
$W=1$	0	$m(0,3)=3$	$m(1,0)=3$	$m(5,0)=5$
$W=2$	0	3	3	5
$W=3$	0	3	3	5
$W=4$	0	3	$m(3,4)=4$	5
$W=5$	0	3	$\max(3,5)=5$	$m(5,3+6)=9$
$W=6$	0	3	5	9
$W=7$	0	3	5	9

Knapsack Solution

- $Knapsack(\{t_1, \dots, t_n\}, W)$

$O(nW)$ ops

table = 2D-array, indexed from 0 to n and 0 to W
table[i, W'] stores subproblem $Knapsack(\{t_1, \dots, t_i\}, W')$

- For $j = 0, \dots, n$:
— table[j, 0] = 0 Base Case 1: no weight to carry \rightarrow no value
- For $k = 0, \dots, W$:
— table[0, k] = 0 Base Case 2: no items to choose from \rightarrow no value

For $j = 1, \dots, n$:

$O(nW)$ loops

For $k = 1, \dots, W$:

reject = table[j - 1, k]

accept = $(v_j + \text{table}[j - 1, k - w_j])$ if $k \geq w_j$ else $-\infty$

table[j, k] = max{reject, accept}

Return table[n, W]

$O(1)$ operations/loop

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Is this polynomial in input size?

- What is the “input size” in term of n and W ?
- Is there $\text{poly}(n, \log W)$ -time algorithm?
 - No, unless $P = NP$.
 - You will learn about “ P vs. NP ” after the midterm.

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Wrap Up

- We have seen more examples of dynamic programming
- **Next:**
 - Using graphs (instead of 1D/2D tables) in dynamic programming algo.
 - For: shortest path problems.

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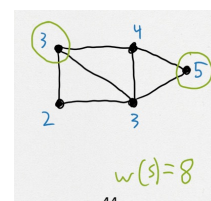
BONUS: MAXIMUM WEIGHT INDEPENDENT SET OF TREES

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Max Weight Independent Set

- **Input:** Graph G with a weight w_v on each **node** v
- A set of nodes S is **independent** if no edges between nodes in S .
- **Output:** **Independent set** S maximizing $\sum_{s \in S} w_s$
- NP-Hard ☹

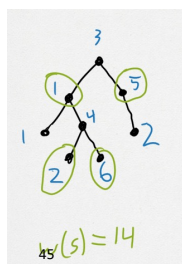


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Max Weight Independent Set **On Trees**

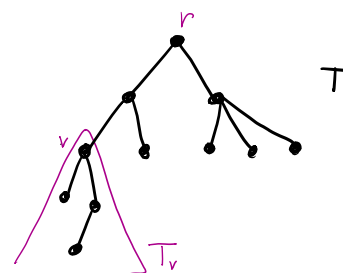
- **Input:** **Tree** T with a weight w_v on each **node** v
- A set of nodes S is **independent** if no edges between nodes in S .
- **Output:** **Independent set** S maximizing $\sum_{s \in S} w_s$
- Admits linear-time algo!



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Coming up with the recurrence

- $\text{MWIS}(v)$: max weight independent set of subtree T_v rooted at v
- **Goal:** $\text{MWIS}(r)$ where r is the root of T



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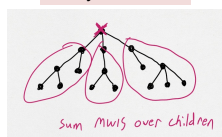
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Coming up with the recurrence

- $\text{MWIS}(v)$: max independent set of subtree T_v rooted at v
- **Goal:** $\text{MWIS}(r)$ where r is the root of T

Recursive Strategy:
Consider the root node r .
Include r in S ?

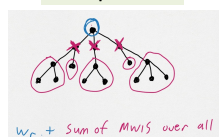
Reject root r



$$\text{MWIS}(r) = \sum_{v \text{ child of } r} \text{MWIS}(v)$$

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Accept root r



$$\text{MWIS}(r) = w_r + \sum_{v \text{ grandchild of } r} \text{MWIS}(v)$$

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Max Weight Independent Set **On Trees**

- $\text{MWIS}(T)$: assume T nonempty
Computes max possible total weight of MWIS
(Could easily tweak algorithm to also output that independent set)
- r = root node of T
table = array, indexed by tree nodes memo[v] stores MWIS of subtree rooted at v
- For each node v in T , in which order???
-
- Return table[r]

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Max Weight Independent Set **On Trees**

- MWIS(T) assumes T is nonempty
Computes max possible total weight of MWIS
(Could easily tweak algorithm to also output that independent set)

r = root node of T
table = array, indexed by tree nodes memo[v] stores MWIS of subtree rooted at v

For each node v in T, in ascending tree order process all descendants before v itself

reject $\leftarrow \sum_{c \text{ child of } v} \text{table}[c]$ don't take v_i
accept $\leftarrow w_v + \sum_{g \text{ grandchild of } v} \text{table}[g]$ take v_i , reject its children
table[v] $\leftarrow \max\{\text{reject}, \text{accept}\}$

Return table[r]

n loops
Ops/loop **depends on # of children/grandchildren**
Tree of branching factor b has $O(nb^2)$ ops total

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Max Weight Independent Set **On Trees**

- Today:**
– $O(nb^2)$ time
- Challenge:**
– $O(n)$ time
– **Hint: Same algorithm. Smarter analysis.**
Show that $\sum_{v \in T} |\text{grandchildren}(v)| = O(n)$

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Upshot: Dynamic Programming on Trees

Many hard problems
usually become easy on trees
via dynamic programming

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Longest Palindromic Substring (Another classic coding problem)

- Given a string $X[1..n]$
- Goal:** Find the length of the longest palindrome in X .
• A **palindrome** is a string that's equal to its reverse
- Example:** "aba", "aca", and "ada" are the longest palindromes in $X = \text{"abacada"}$ (Note: "aacaa" doesn't count!)
- Q:** What's a brute force algorithm?
• Try every substring $X[i..j]$

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Recurrence for PAL

- Given a string $X[1..n]$
- Let $PAL(i, j)$ be a Boolean (T/F) value for whether $X[i..j]$ is a palindrome

Sometimes the problem
we solve might seem quite
"different"

$$PAL(i, j) = \begin{cases} X[i] = X[j] & j \leq i + 1 \\ X[i] = X[j] \text{ and } PAL(i + 1, j - 1) & j > i + 1 \end{cases}$$

Q: Given this recurrence, how do we
find the length of a longest palindrome?

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