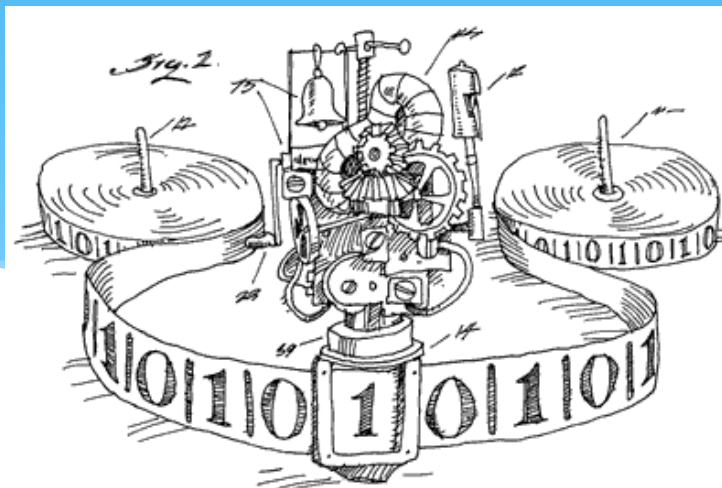


# EECS 376: Foundations of Computer Science

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18 January 2023

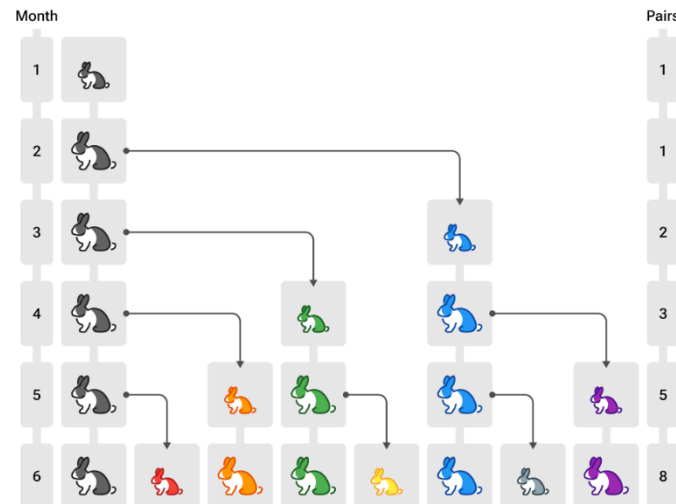


“If you can solve it, it is an exercise; otherwise, it is a research problem”

- Richard E. Bellman,

The Primary Expositor of Dynamic Programming

# Algorithmic Strategy: Dynamic Programming



“An interesting question is, ‘Where did the name, dynamic programming, come from?’ The 1950s were not good years for mathematical research...

I had to do something to shield [SecDef] Wilson and the Air Force from the fact that I was really doing mathematics...

I was interested in planning, in decision making, in thinking. But planning is not a good word... I decided to use the word, ‘programming.’

...It’s impossible to use the word ‘dynamic’ in a pejorative sense...

Thus, I thought dynamic programming was a good name.

It was something not even a Congressman could object to.”

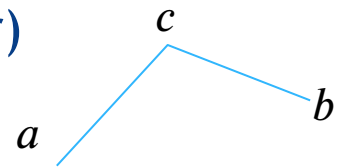


# Dynamic Programming

**High-level Idea:** Break a big problem into smaller (easier) subproblems—like D&C—but also exploiting:

1. Principle of “optimal substructure”:  
any “piece” of an optimal structure is itself optimal.

**Example:** A subpath of any shortest path is itself a shortest path between its endpoints.



2. Overlapping sub-problems: “many” subproblems re-occur “many” times.

**Example:** When computing the Fibonacci sequence using the rule  $F_n = F_{n-1} + F_{n-2}$ , “many” recursive calls will be repeated.

# Warm-Up: Fibonacci

- \* Recurrence for Fibonacci:

$$F(n) = \begin{cases} 1 & \text{if } n = 0 \text{ or } n = 1 \\ F(n-1) + F(n-2) & \text{if } n \geq 2 \end{cases}$$

- \* Given a recurrence, three ways to compute its values:
- \* **Top-down Recursive (Naïve):** Starting at desired input, **recurse down** to base case(s)
- \* **Top-down w/ Memoization:** Same as naïve, but **save results** as they're computed, **reusing** already-computed results
- \* **Bottom-up Table:** Start from base case(s), **build up** to desired result
- \* All these are '**mechanical translations**' of the recurrence

# Fib: Naïve Implementation

- \* Recurrence for Fibonacci:

$$F(n) = \begin{cases} 1 & \text{if } n = 0 \text{ or } n = 1 \\ F(n-1) + F(n-2) & \text{if } n \geq 2 \end{cases}$$

- \* **Top-down Recursive (Naïve):**

- \* **Algorithm:**  $Fib(n)$

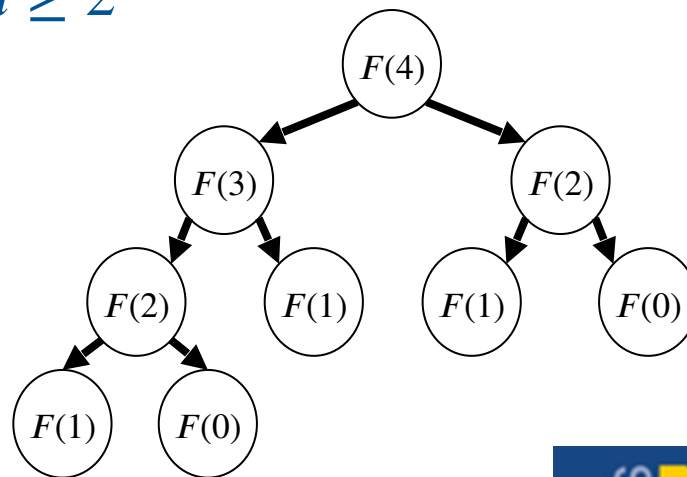
- \* If  $n = 0$  OR  $n = 1$

- \* Return 1

- \* Return  $Fib(n-1) + Fib(n-2)$

- \* Pro: direct translation of recurrence

- \* Con: *exponential* runtime:  $T(n) = T(n-1) + T(n-2) + O(1)$



# Fib: Top-Down w/Memoization

- \* Recurrence for Fibonacci:

$$F(n) = \begin{cases} 1 & \text{if } n = 0 \text{ or } n = 1 \\ F(n-1) + F(n-2) & \text{if } n \geq 2 \end{cases}$$

- \* **Top-down Memoization:**

- \*  $memo[0\dots n] := \text{empty table}$

- \* **Algorithm:**  $Fib(n)$

- \* If  $n = 0$  OR  $n = 1$

- \* **Return** 1

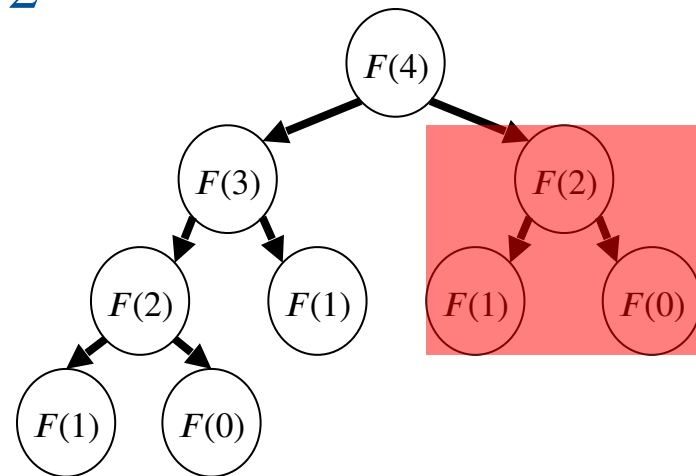
- \* If  $memo[n]$  undefined:

- \*  $memo[n] := Fib(n-1) + Fib(n-2)$

- \* **Return**  $memo[n]$

- \* **Pros:** much faster (but how much?)

- \* **Con:** global memo, clumsy impl., hard to analyze runtime



# Fib: Bottom Up

- \* Recurrence for Fibonacci:

$$F(n) = \begin{cases} 1 & \text{if } n = 0 \text{ or } n = 1 \\ F(n-1) + F(n-2) & \text{if } n \geq 2 \end{cases}$$

- \* **Bottom-up Table:**

- \* **Algorithm:**  $Fib(n)$

- \* allocate  $table[0..n]$

- \*  $table(0) := 1$

- \*  $table(1) := 1$

- \* For  $i = 2$  to  $n$ :

- \*  $table(i) := table(i-1) + table(i-2)$

- \* Return  $table(n)$

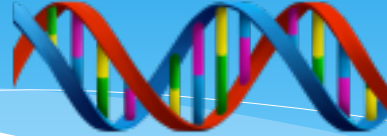
- \* **Pro:** much faster, no globals, easy to analyze runtime

- \* **Cons(?):** must compute *entire* table of smaller results (but usually end up doing this anyway, in every strategy)

1	1	2	3	5	8	13	21	34	55
0	1	2	3	4	5	6	7	8	9



# DNA Comparison



- \* Your DNA is a (*long*) string over {A, T, C, G}.
  - \* Small chance of random insertions, deletions, edits
- \* “Humans and chimps are 98.9% similar.”
  - \* *X*: ACCGGTCGAGTGCGCGGAAGCCGGCCGAA
  - \* *Y*: GTCGTTCGGAATGCCGTTGCTCTGTAA
- \* The length of the longest common subsequence between two genomes is a measure of similarity.
- \* How efficiently can we compute an LCS of *X*, *Y*?
  - \* |human genome|  $\approx$  3bil, |chimp genome|  $\approx$  2.8bil

# Longest Common Subsequence

- \* **Definition:** A *subsequence* of a string  $s$  is a (not necessarily contiguous) subset of the characters of  $s$ , in their original order.
  - \* **Example:** for  $s$  = “Fibonacci sequence”
    - \* “Fun”
    - \* “seen”
    - \* “cse”
    - \* ...
- \* **Goal:** Given strings  $X[1..n]$  and  $Y[1..m]$ , find a *longest common subsequence* of  $X$  and  $Y$ .
  - \* Largest string obtainable from  $X$  and  $Y$  by deleting chars
- \* **Example:** “Gole” is an LCS of “Google” and “Go Blue”.
- \* **Q:** What’s a brute force solution?
  - \* Each character of  $X$  and  $Y$  is either deleted or not.

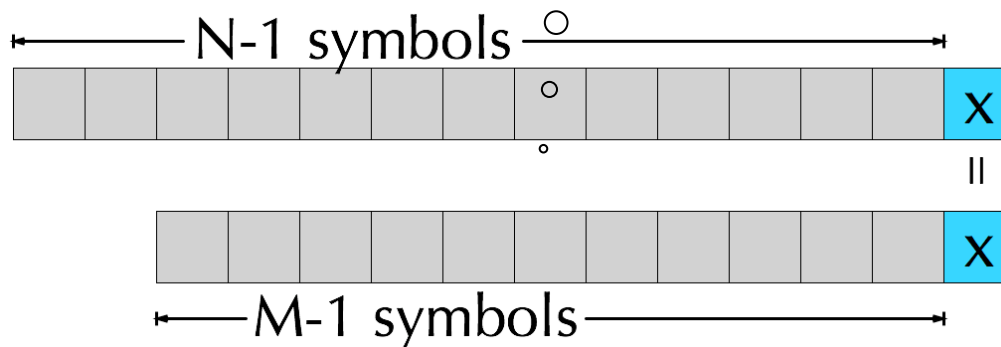
# Dynamic Programming for LCS (and every other DP problem)

- \* Let  $X[1 \dots n]$  and  $Y[1 \dots m]$  be two given strings.
- \* **Key Idea #1:** to start, focus on the *length* of an LCS
  - \* (After finding length, finding an actual LCS will be easy!)
- \* **Key Idea #2:** discover a *recurrence* for LCS length, relative to suitable *substrings* (subproblems)
  - \* Which substrings to consider? How to relate them?
  - \* An “art”!
- \* Define  $LCS(i, j) :=$  length of LCS of  $X[1 \dots i]$  and  $Y[1 \dots j]$ .
  - \* Subproblems are (pairs of) *prefixes* of  $X$  and  $Y$ .

# LCS Recurrence (Part 1)

- \*  $LCS(i, j) :=$  length of LCS of  $X[1..i]$  and  $Y[1..j]$ .
- \* If the last characters are equal, i.e.,  $X[i] = Y[j]$ :
  - \*  $LCS(i, j) = 1 + LCS(i - 1, j - 1)$ .

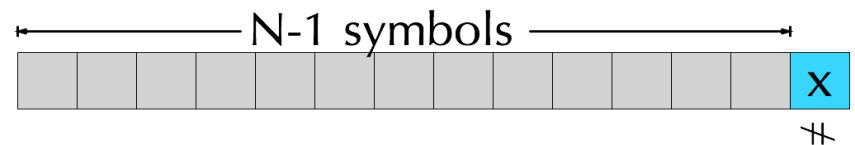
Principle of Optimality



# LCS Recurrence (Part 2)

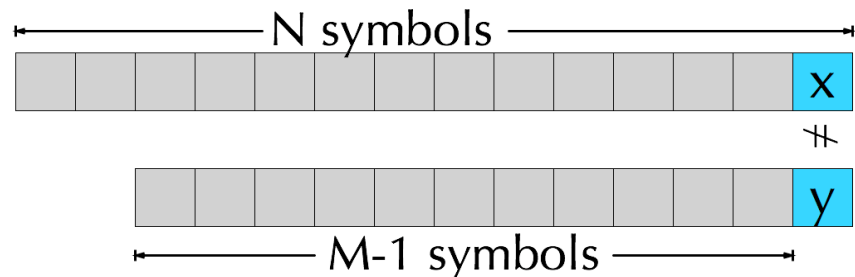
- \*  $LCS(i, j) :=$  length of LCS of  $X[1 \dots i]$  and  $Y[1 \dots j]$ .
- \* If the last characters are **not** equal, i.e.,  $X[i] \neq Y[j]$ :
- \*  $LCS(i, j) =$  Maximum of the only two options:

$LCS(i-1, j)$



and

$LCS(i, j-1)$



# Full Recurrence for LCS

\*  $LCS(i, j)$  = LCS length of  $X[1 \dots i]$  and  $Y[1 \dots j]$ .

\* Then:

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

\* **Naïve Implementation:** Exponential runtime!

\* **Bottom up:** There are  $O(nm)$  values of interest:  $LCS(i, j)$  for  $0 \leq i \leq n$  and  $0 \leq j \leq m$  (overlapping sub-problems)

<https://www.cs.usfca.edu/~galles/visualization/DPLCS.html>

# LCS Dynamic Programming in Action

- \* See hand-written notes (or visualization webpage):
  1. Bottom-up table filling
  2. Recovering an LCS itself from the lengths

# Longest **Increasing** Subsequence

- \* Given: an array of numbers  $A[1 \dots n]$
- \* **Goal:** Find (the length of) a *longest increasing subsequence* of  $A$ .
  - \* Longest increasing array obtainable by deleting entries of  $A$
- \* **Example:**  $[5, 6, 7, 8]$  is an increasing subsequence of  $[5, 6, 0, 7, 1, 2, 8, 4, 0, 5, 3]$ .
  - \* **Q:** What's a longest one?
- \* **Q:** What's a brute force algorithm?
  - \* Each entry is either deleted or not:  $\geq 2^n$  time!



# Recurrence for LIS?

- \* Given an array of integers  $A[1..n]$
- \* Let  $LIS(i)$  be the length of a longest increasing subsequence of  $A[1..i]$ .
- \* **Q:** What's  $LIS(4)$  if  $A = [1, 2, 0, 4, 3]$ ? If  $A = [1, 2, 3, 0, 5]$ ?
- \* **Q:** Can we determine whether  $A[i]$  extends an LIS of  $A[1..j]$  by only looking at  $A[i]$  and  $A[j]$ ?
  - \* **No.** We need more information to determine whether  $LIS(i) \geq 1 + LIS(j)$ .

# Recurrence for $LIS_{at}$ ?

- \* Given an array of integers  $A[1..n]$
- \* Let  $LIS_{at}(i)$  be the length of a longest increasing subsequence of  $A[1..i]$  **that ends with  $A[i]$** .
- \* **Q:** What's  $LIS_{at}(4)$  if  $A = [1, 2, 0, 4, 3]$ ?  
If  $A = [1, 2, 3, 0, 5]$ ?
- \* **Q:** Can we determine if  $A[i]$  extends an LIS of  $A[1..j]$  **that ends with  $A[j]$**  by only looking at  $A[i]$  and  $A[j]$ ?
  - \* **Yes.** If  $A[i] > A[j]$ , then  $LIS_{at}(i) \geq 1 + LIS_{at}(j)$ .

# Recurrence for $LIS_{at}$

We don't know where the second-to-last element of the LIS ending at  $A[i]$  is, so we consider every element that it *could* be!

- \* Given an array of integers  $A[1..n]$
- \* Let  $LIS_{at}(i)$  be the length of a longest increasing subsequence of  $A[1..i]$  that ends with  $A[i]$

$$LIS_{at}(i) = \begin{cases} 1 & \text{if } i = 1 \text{ or } A[j] \geq A[i] \text{ for all } j < i \\ 1 + \max \{ LIS_{at}(j) \mid A[j] < A[i] \text{ and } j < i \} & \text{otherwise} \end{cases}$$

**Q:** Given this recurrence, how to compute the length of an LIS of  $A$ ? An LIS itself?

<http://rosulek.github.io/vamonos/demos/lis.html>

# LIS Dynamic Programming in Action

- \* See hand-written notes (or visualization webpage):
  - \*  $A = [5, 6, 0, 7, 1, 2, 8, 4, 0, 5, 3]$
  - \* Bottom-up table filling, with “back pointers”
  - \* Recovering an LIS itself from the table