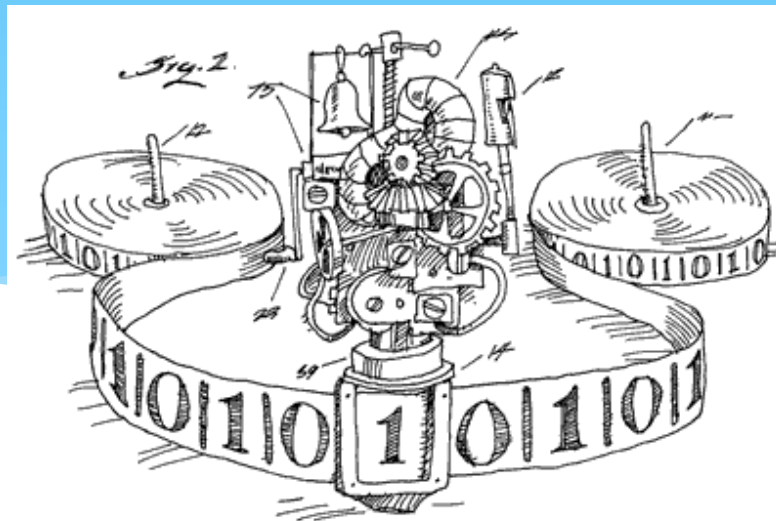


EECS 376: Foundations of Computer Science

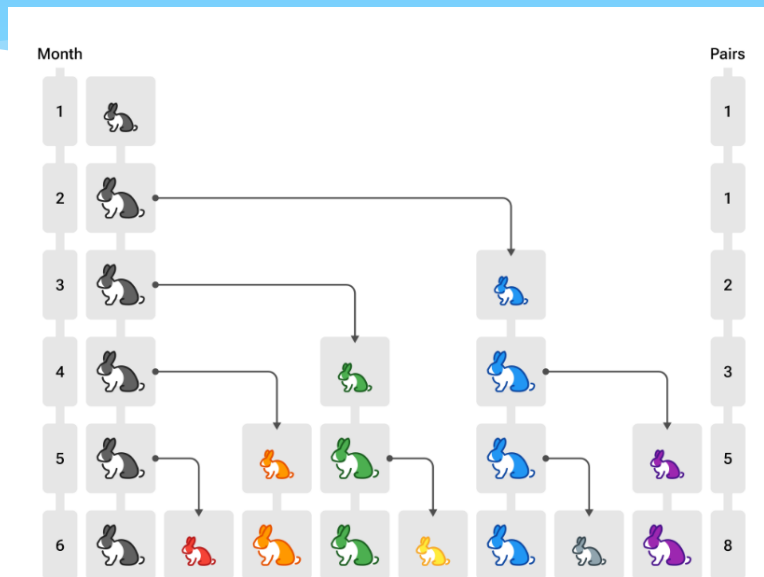
Seth Pettie

Lecture 4



“If you can solve it, it is an exercise; otherwise, it is a research problem”
-- Richard E. Bellman

Algorithmic Strategy: Dynamic Programming



Recap

- * **Previously:** divide and conquer
 - * A recurrence – break into smaller sub-problems and combine
 - * Design goal is to minimize the number of recursive calls k and time to combine $O(n^d)$
 - * Examples: Closest pair, Karatsuba

Dynamic Programming

- * **Today:** dynamic programming
 - * A recurrence – break into smaller sub-problems and combine
 - * ~~Design goal is to minimize the number of recursive calls k and time to combine $O(n^d)$~~
 - * Don't worry about minimizing number of recursive calls!
 - * **Idea: Maximize number of *repeated* recursive calls**

Dynamic Programming?

- * Dynamic programming is *not*:
 - * Dynamic, or 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.**” – Richard Bellman

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)
- * **Dynamic programming**
 - * **Top-down with 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 '**translate**' the recurrence into an algorithm

Fib: Naïve Implementation

- * The x th Fibonacci number, for x a non-negative integer:

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

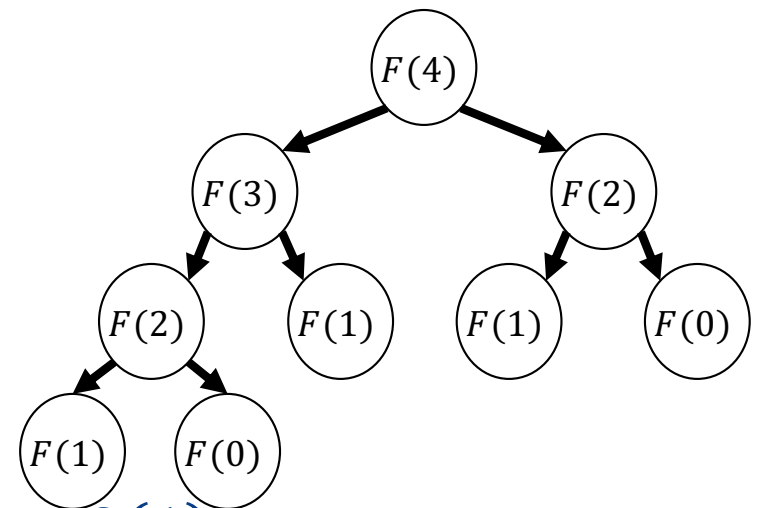
- * **Top-down recursive (naïve):**

```
F( $n$ ): //  $n \geq 0$  an integer
if  $n = 0$  or  $n = 1$  then return 1
return F( $n - 1$ ) + F( $n - 2$ )
```

- * **Pro:** direct translation of recurrence
- * **Con:** exponential runtime:

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

$$= O(F(n)) = O(\varphi^n) = O(1.62^n)$$



Fib: Memoization

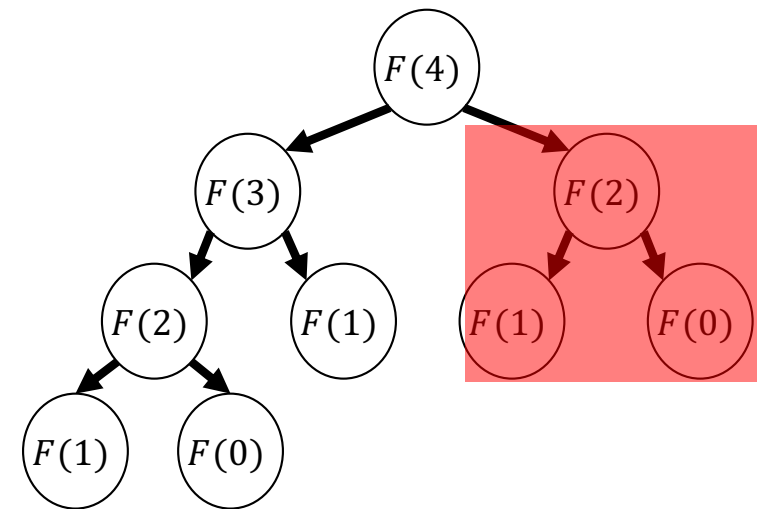
- * The x th Fibonacci number, for n a non-negative integer:

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

- * **Top-down memoization:**

```

allocate  $F[1..n]$  // entries initially NULL
 $F[1] \leftarrow 1, F[2] \leftarrow 1$ 
M-F( $n$ ): // memoized implementation of  $F_n$ 
if  $F[n] = \text{NULL}$  then
     $F[n] \leftarrow \mathbf{MF}(n-1) + \mathbf{MF}(n-2)$ 
return  $F[n]$ 
  
```



- * **Pros:** much faster (but how much?)
- * **Con:** requires accessing global memory, hard to analyze runtime

Fib: Bottom up (dynamic programming)

- * 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:**

```

DP-F(x): // table implementation of  $F_n$ 
allocate  $F[1..n]$ 
 $F[0] \leftarrow 1, F[1] \leftarrow 1$ 
for  $i = 2..n$ 
     $F[i] \leftarrow F[i-1] + F[i-2]$ 
return  $F[n]$ 

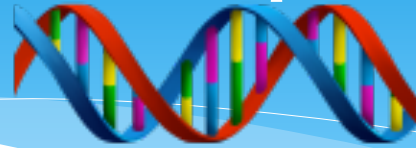
```

- * **Q:** What is the runtime of this algorithm?
- * $T(n) = O(n)$

x	F[x]
0	1
1	1
2	2
3	3
4	5
5	8
6	13
7	21

- * **Pro:** much faster, no globals, easier to analyze runtime
- * **Cons:** must compute *entire* table of smaller results (but usually end up doing this anyway, in every strategy)

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 : ACCGGT**CGAGTGCGCGGAAGCCGGCCGAA**
 - * Y : **GTCGTT**CGGAATGCCGTT**GCTCTGTAA**
- * 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 ?
 - * $|\text{human genome}| \approx 3\text{bil}$, $|\text{chimp genome}| \approx 2.8\text{bil}$

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
 - * 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: $O(2^{m+n})$

Better way?

- * Where to begin? If we can just find one pair of characters that *must* be matched in an LCS, then we can delete those characters and recurse
- * **Idea 1:** Given $X[1..m]$ and $Y[1..n]$, suppose last characters are the same, i.e. $X[m] = Y[n]$
- * **Q:** Should we always match $X[m]$ and $Y[n]$?
 - * **Example:** $X = \text{"ACTG"} , Y = \text{"ATAG"}$. **Q:** Is matching the blue G's a mistake? What about matching the red A's?
 - * Yes, if $X[m] = Y[n]$. Why? (Can matching $X[m]$ and $Y[n]$ ever be a mistake?)

Better way?

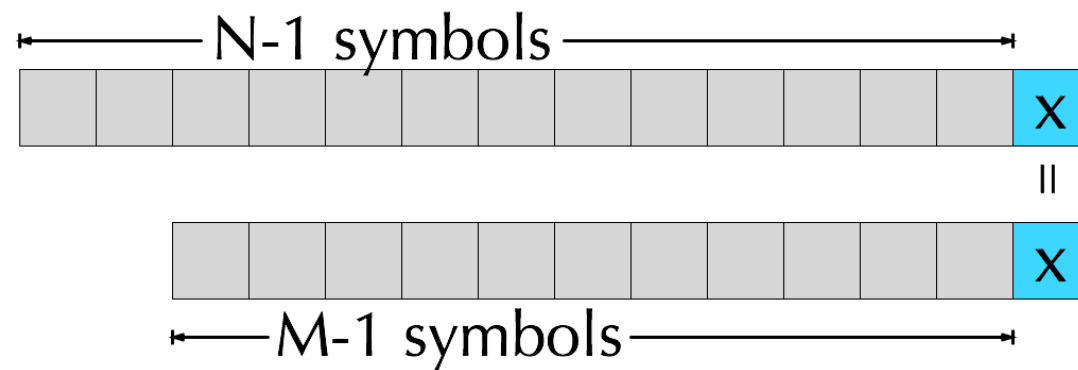
- * Given strings $X[1..m]$ and $Y[1..n]$
- * Let $LCS(i, j)$ denote the *length* of a longest common subsequence of $X[1..i]$ and $Y[1..j]$. ($i = 0$ or $j = 0$ denotes the empty string)
- * **Idea 2:** If $LCS(i, j - 1) = LCS(i, j) - 1$ or $LCS(i - 1, j) = LCS(i, j) - 1$, then the match $X[i]$ with $Y[j]$ is in a longest common subsequence of $X[1..i]$ and $Y[1..j]$.
 - * We don't have to find the longest common subsequence directly, all we need is to find the *length* of the longest common subsequence, $LCS(i, j)$!
- * **Example:** Suppose $X = \text{"ATGCC"}$ and $Y = \text{"TAGC"}$.
 - * Q: What's $LCS(1, 0)$?
 - * Q: What's $LCS(5, 3)$?
 - * Q: What's $LCS(4, 4)$?
 - * Q: What's $LCS(5, 4)$?

Recurrence for LCS

- * **Step 1:** *To find a dynamic programming algorithm, always start with a recurrence*
- * We want a recurrence for $LCS(i, j)$
- * **Q:** What should the base case be?
 - * **Base case:** if $i = 0$ or $j = 0$ (empty string); $LCS(i, j) = 0$
- * **Case 1:** $X[i] = Y[j]$ (ends with the same character)
 - * **Example:** $X[1..i] = \text{"CTGCA"}$ and $Y[1..j] = \text{"TCGA"}$
 - * $LCS(i, j) = 1 + LCS(i - 1, j - 1)$

Recurrence for *LCS*

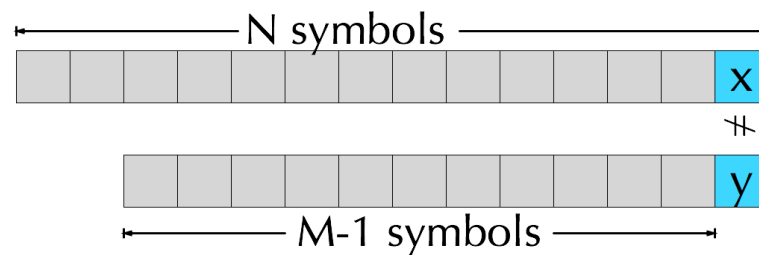
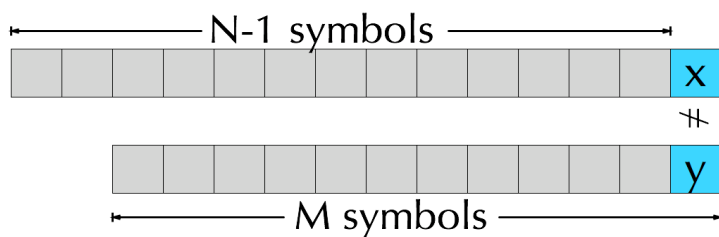
- * **Case 1:** $X[i] = Y[j]$ (ends with the same character)
 - * **Example:** $X[1..i] = \text{"CTGCA"}$ and $Y[1..j] = \text{"TCGA"}$
 - * $LCS(i, j) = 1 + LCS(i - 1, j - 1)$



Recurrence for LCS

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

- * **Case 2:** $X[i] \neq Y[j]$ (end with different characters)
 - * **Example:** $X[1..i] = \text{“GTCA”}$ and $Y[1..j] = \text{“GTC”}$
 - * At least one of the letters is not part of LCS
 - * **Q:** How do we know which one?
 - * Try both! $LCS(i, j) = \max\{LCS(i - 1, j), LCS(i, j - 1)\}$



Recurrence for *LCS*

- * Given strings $X[1..m]$ and $Y[1..n]$
- * Let $LCS(i, j)$ denote the length of a longest common subsequence of $X[1..i]$ and $Y[1..j]$

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

Q: Given this recurrence, how do we find the length of an LCS of X and Y ?

$LCS(m, n)$

Table Implementation of LCS

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

LCS($X[1..m], Y[1..n]$): // table implementation of *LCS*

allocate $L[0..m][0..n]$

$L[0][1..n] \leftarrow 0, L[1..m][0] \leftarrow 0$

for $i = 1..m$ and $j = 1..n$: // fill columns of row i , one by one

if $X[i] = Y[j]$ **then** $L[i][j] \leftarrow 1 + L[i - 1][j - 1]$

else $L[i][j] \leftarrow \max\{L[i - 1][j], L[i][j - 1]\}$

return $L[m][n]$

Runtime: $O(mn)$

Q: How could we recover actual LCS, not just its length?
Store “prev” pointer in $L[i][j]$ depending on case.

Filling the table

- * Try this visualization out!

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

Longest Increasing Subsequence

(A classic coding problem)

- * Given an array of integers $A[1..n]$
- * **Goal:** Find the length of a **longest increasing subsequence** of A
 - * largest inc. array obtainable by deleting parts of A
- * **Example:** $[5,6,7]$ is an increasing subsequence of $[5, 6, 0, 7, 1, 2, 0, 4, 0]$. **Q:** longest?
- * **Q:** What's a brute force solution?
 - * Each integer is either deleted or not, $O(2^n)$ 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 $A = [1,1,2,1,3]$? $A = [5,6,8,2,3]$?
- * Before: divided between last element(s) and rest of list(s). Can we do the same here?
- * **Q:** Can we determine if $A[i]$ extends LIS of $A[1..i-1]$ by only looking at $A[i]$ and $A[i-1]$?
 - * **Example:** $A[1..i-1] = [5,6,8,2]$, $A[i] = 3$
 - * **No.** We need more information before we can conclude that $LIS(i) = 1 + LIS(i-1)$

Recurrence for LIS_{at} ?

The subproblems we solve depend on how we solve the problem recursively

- * 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 at $A[i]$**
- * Q: What's $LIS_{at}(4)$ if $A = [1,1,2,1,3]$? $A = [5,6,8,2,3]$?
- * Q: Can we determine if $A[i]$ extends LIS of $A[1..j]$ **ending at $A[j]$** by only looking at $A[i]$ and $A[j]$?
 - * Example: $A[1..j] = [1,1,2]$, $A[i] = 3$
 - * **Yes.** If $A[i] > A[j]$, then
$$LIS_{at}(i) \geq 1 + LIS_{at}(j)$$

Recurrence for LIS_{at}

$$LIS_{at}(i) = \begin{cases} 0 & i = 0 \\ 1 + \max \left\{ LIS_{at}(j) \mid \begin{array}{l} (A[j] < A[i] \text{ and } j < i) \\ \text{or } j = 0 \end{array} \right\} & i \neq 0 \end{cases}$$

```
LIS( $A[1..n]$ ): // table implementation of  $LCS$   
allocate  $L[0..n]$   
 $L[0] \leftarrow 0$   
for  $i = 1..n$ : // fill table  
     $l \leftarrow 0$   
    for  $j = 1..i - 1$ :  
        if  $A[j] < A[i]$ :  $l \leftarrow \max\{l, L[j]\}$   
     $L[i] \leftarrow l + 1$   
return ?
```

- * The conversion from recurrence to table is *mechanical*
- * **Q:** Given this recurrence, how do we determine the length of a LIS?

Recurrence for LIS_{at}

$$LIS_{at}(i) = \begin{cases} 0 & i = 0 \\ 1 + \max \left\{ LIS_{at}(j) \mid \begin{array}{l} (A[j] < A[i] \text{ and } j < i) \\ \text{or } j = 0 \end{array} \right\} & i \neq 0 \end{cases}$$

LIS($A[1..n]$): // table implementation of *LCS*

allocate $L[0..n]$

$L[0] \leftarrow 0$

for $i = 1..n$: // fill table

$l \leftarrow 0$

for $j = 1..i - 1$:

if $A[j] < A[i]$: $l \leftarrow \max\{l, L[j]\}$

$L[i] \leftarrow l + 1$

return $\max_{1 \leq i \leq n} L[i]$

Runtime: $O(n^2)$

- * The conversion from recurrence to table is *mechanical*
- * **Q:** Given this recurrence, how do we determine the length of a LIS?
 - * $LIS(n) = \max_{1 \leq i \leq n} LIS_{at}(i)$
- * **Q:** Runtime?