### MA 252: Data Structures and Algorithms Lecture 32

http://www.iitg.ernet.in/psm/indexing\_ma252/y12/index.html

#### Partha Sarathi Mandal

Dept. of Mathematics, IIT Guwahati

### The All-Pairs Shortest Paths Problem

Given a weighted digraph G = (V, E) with weight function w: E → R, (R, is the set of real numbers) determine the length of the shortest path i.e., distance between all pairs of vertices in G. Here we assume that there are no cycles with zero or negative cost.

#### **SSSP**

- Dijkstra's SSSP algorithm requires *all* edge weights to be nonnegative
  - even more restrictive than outlawing negative weight cycles
  - Runtime O(E log V) if priority queue is implemented with a binary heap.
  - Runtime  $O(V \log V + E)$  priority queue is implemented with a Fibonacci heap.
- Bellman-Ford SSSP algorithm can handle negative edge weights
  - even "handles" negative weight cycles by reporting they exist
  - Runtime O(V E)

#### All pairs shortest paths

- Simple approach
  - Call Dijkstra's |V| times
  - $O(|V| |E| \log |V|) / O(|V|^2 \log |V| + |V| |E|))$
  - Call Bellman-Ford |V| times
  - $O(|V|^2 |E|)$
- A dynamic programming solution. Only assumes no negative weight cycles.
  - First version is  $\Theta(|V|^4)$
  - Repeated squaring reduces to  $\Theta(|V|^3 \log |V|)$
- Floyd-Warshall  $-\Theta(|V|^3)$
- Johnson's algorithm  $O(|V|^2 \log |V| + |V| |E|)$

### Dynamic programming

#### Computing Fibonacci Numbers

#### Fibonacci numbers:

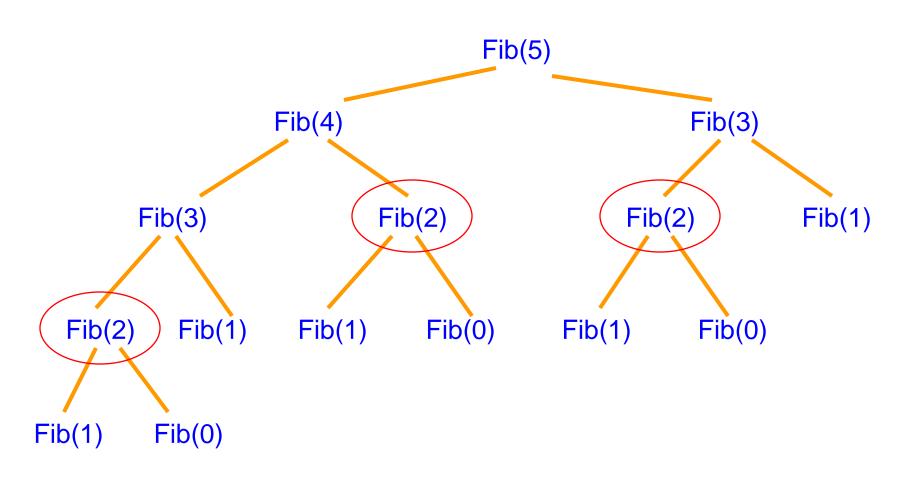
- $-F_0 = 0$   $-F_1 = 1$  $-F_n = F_{n-1} + F_{n-2}$  for n > 1
- Sequence is 0, 1, 1, 2, 3, 5, 8, 13, ...

#### Computing Fibonacci Numbers

• Obvious recursive algorithm:

- Fib(*n*):
  - if n = 0 or 1 then return n
  - else return (Fib(n-1) + Fib(n-2))

#### Recursion Tree for Fib(5)



#### How Many Recursive Calls?

- If all leaves had the same depth, then there would be about  $2^n$  recursive calls.
- But this is over-counting.
- However with more careful counting it can be shown that it is  $\Omega((1.6)^n)$
- Exponential!

#### Save Work

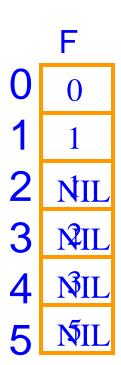
- Wasteful approach repeat work unnecessarily
  - Fib(2) is computed **three** times
- Instead, compute Fib(2) once, store result in a table, and access it when needed

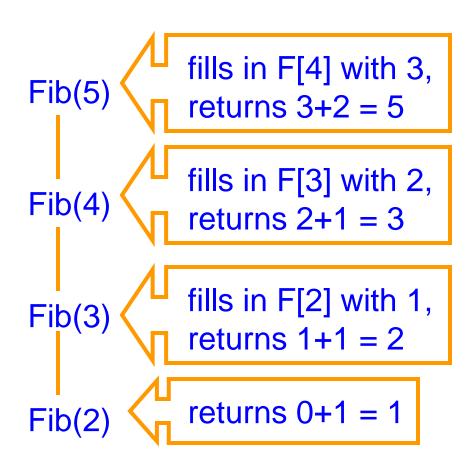
#### More Efficient Recursive Algo

- F[0] := 0; F[1] := 1; F[n] := Fib(n);
  Fib(n):

  if n = 0 or 1 then return F[n]
  if F[n-1] = NIL then F[n-1] := Fib(n-1)
  if F[n-2] = NIL then F[n-2] := Fib(n-2)
  return (F[n-1] + F[n-2])
- computes each F[i] only once

#### Example of Memoized Fib



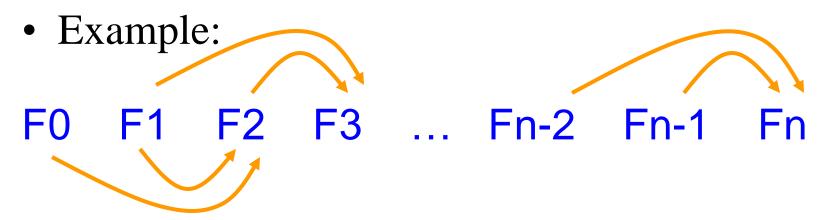


#### Get Rid of the Recursion

- Recursion adds overhead
  - extra time for function calls
  - extra space to store information on the runtime stack about each currently active function call
- Avoid the recursion overhead by filling in the table entries **bottom up**, *instead* of **top down**.

#### Subproblem Dependencies

 Figure out which subproblems rely on which other subproblems



### Order for Computing Subproblems

- Then figure out an order for computing the subproblems that respects the dependencies:
  - when you are solving a subproblem, you have already solved all the subproblems on which it depends
- Example: Just solve them in the order  $F_0$ ,  $F_1$ ,  $F_2$ ,  $F_3$ ,...



#### DP Solution for Fibonacci

- 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]
- Can perform application-specific optimizations
  - e.g., save space by only keeping last two numbers computed

## Dynamic Programming (DP) Paradigm

- DP is typically applied to Optimization problems.
- DP can be applied when a problem exhibits:
- Optimal substructure:
  - Is an optimal solution to the problem contains within it optimal solutions to subproblems.
- Overlapping subproblems:
  - If recursive algorithm revisits the same problem over and over again.

# Dynamic Programming (DP) Paradigm

- DP can be applied when the solution of a problem includes solutions to subproblems
- We need to find a recursive formula for the solution
- We can recursively solve subproblems, starting from the trivial case, and save their solutions in memory
- In the end we'll get the solution of the whole problem

#### Dynamic programming

- One of the most important algorithm tools!
- Very common interview question

- Method for solving problems where optimal solutions can be defined in terms of optimal solutions to sub-problems AND
- the sub-problems are overlapping

### Identifying a dynamic programming problem

• The solution can be defined with respect to solutions to subproblems

• The subproblems created are overlapping, that is we see the same subproblems repeated

### Two main ideas for dynamic programming

• Identify a solution to the problem with respect to **smaller** subproblems

$$-F(n) = F(n-1) + F(n-2)$$

• Bottom up: start with solutions to the smallest problems and build solutions to the larger problems

```
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] P. S. Mandal, IITG
```

use an array to store solutions to subproblems

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

ABA?

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = ABACDABAB

**ABA** 

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

ACA?

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

**ACA** 

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

DCA?

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B



• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

AADAA?

• For a sequence  $X = x_1, x_2, ..., x_n$ , a subsequence is a subset of the sequence defined by a set of increasing indices  $(i_1, i_2, ..., i_k)$  where  $1 \le i_1 < i_2 < ... < i_k \le n$ 

X = A B A C D A B A B

**AADAA** 

#### LCS problem

- Given two sequences X and Y, a common subsequence is a subsequence that occurs in both X and Y
- Given two sequences  $X = x_1, x_2, ..., x_m$  and  $Y = y_1, y_2, ..., y_n$ , What is the **longest** common subsequence?

$$X = A B C B D A B$$
  
 $Y = B D C A B A$ 

#### LCS problem

- Given two sequences X and Y, a common subsequence is a subsequence that occurs in both X and Y
- Given two sequences  $X = x_1, x_2, ..., x_m$  and  $Y = y_1, y_2, ..., y_n$ , What is the **longest** common subsequence?

$$X = A B C B D A B$$

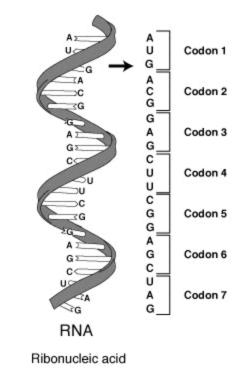
$$Y = B D C A B A$$

The sequences {B, D, A, B} of length 4 are the LCS of *X* and *Y*, since there is no common subsequence of length 5 or greater.

#### LCS problem

Application:

comparison of two DNA strings



**Brute force** algorithm would compare each subsequence of X with the symbols in Y

#### LCS Algorithm

- **Brute-force algorithm:** For every subsequence of *x*, check if it's a subsequence of *y* 
  - How many subsequences of x are there?
  - What will be the running time of the brute-force algorithm?

#### LCS Algorithm

- if |X| = m, |Y| = n, then there are  $2^m$  subsequences of x; we must compare each with Y (n comparisons)
- So the running time of the brute-force algorithm is O(n 2<sup>m</sup>)
- Notice that the LCS problem has *optimal substructure*: solutions of subproblems are parts of the final solution.
- Subproblems:
  - "find LCS of pairs of *prefixes* of X and Y"

#### LCS Algorithm

- First we'll find the length of LCS. Later we'll modify the algorithm to find LCS itself.
- Define  $X_i$ ,  $Y_j$  to be the prefixes of X and Y of length i and j respectively
- Define c[i,j] to be the length of LCS of  $X_i$  and  $Y_j$
- Then the length of LCS of X and Y will be c[m,n]
- c[m,n] is the final solution.

$$X = A B C B D A B$$

$$Y = B D C A B A$$

$$X = ABCBDA$$
?

$$Y = B D C A B$$
?

Is the last character part of the LCS?

$$X = ABCBDA$$
?

$$Y = B D C A B$$
?

Two cases: either the characters are the same or they're different

$$X = ABCBDA$$

$$LCS$$
The characters are part of the LCS 
$$Y = BDCABA$$

If they're the same

$$LCS(X,Y) = LCS(X_{m-1}, Y_{n-1}) + 1$$

P. S. Mandal, IITG

$$X = ABCBDAB$$

$$LCS$$

$$Y = BDCABA$$

If they're different

$$LCS(X,Y) = LCS(X_{m-1},Y)$$

P. S. Mandal, IITG

$$X = ABCBDAB$$

$$LCS$$

$$Y = BDCABA$$

If they're different

$$LCS(X,Y) = LCS(X,Y_{n-1})$$

P. S. Mandal, IITG

$$X = ABCBDAB$$

$$Y = BDCABA$$

$$X = ABCBDAB$$

$$Y = B D C A B A$$

If they're different
P. S. Mandal, IITG

$$X = A B C B D A B$$

$$Y = B D C A B A$$

$$LCS(X,Y) = \begin{cases} 1 + LCS(X_{m-1}, Y_{n-1}) & \text{if } x_m = y_n \\ \max(LCS(X_{m-1}, Y), LCS(X, Y_{n-1}) & \text{otherwise} \end{cases}$$