

# EE360C: ALGORITHMS

## DYNAMIC PROGRAMMING

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# A REVIEW OF ALGORITHMIC TECHNIQUES

## Greedy

Build up a solution incrementally, myopically optimizing some local criterion

## Divide and Conquer

Break up a problem into two (or more) subproblems, solve each subproblem independently, and combine the solutions to the subproblems to form a solution to the original problem

## Dynamic Programming

Break up a problem into a series of overlapping subproblems and build up solutions to larger and larger subproblems.

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Bellman pioneered the systematic study of dynamic programming in the 1950s.

## Etymology

- Dynamic programming = planning over time
- Secretary of Defense at the time was hostile to mathematical research
- Bellman sought an impressive name to avoid confrontation
  - “something not even a Congressman could object to”

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**Dynamic Programming** is an algorithm design technique that, like divide and conquer, relies on solutions to sub problems to solve a problem

- in dynamic programming, however, the subproblems are not independent
- a dynamic programming algorithm solves each of these subproblems just once, saving time in comparison to a traditional divide and conquer approach

Dynamic programming is commonly used for **optimization problems** in which many possible solutions exist; the algorithm helps us find one of the possibilities

- every solution to the problem has a value
- the algorithm helps us find a solution with the optimal value

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Dynamic programming algorithms typically involve the following steps:

- characterize the structure of an optimal solution
- recursively define the value of an optimal solution
- compute the value of an optimal solution in a bottom-up fashion
- construct an optimal solution from the computed information

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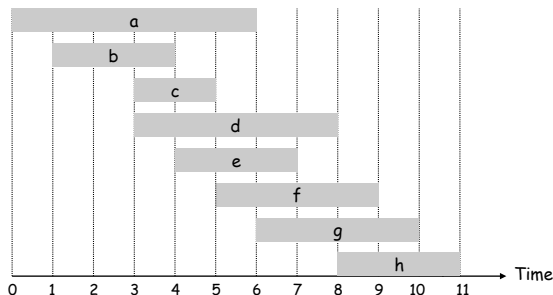
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# WEIGHTED INTERVAL SCHEDULING

The weighted interval scheduling problem:

- Job  $j$  starts at  $s_j$ , finishes at  $f_j$ , and has weight or value  $v_j$
- Two jobs are *compatible* if they do not overlap
- Goal: find maximum weight subset of mutually compatible jobs



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# UNWEIGHTED INTERVAL SCHEDULING

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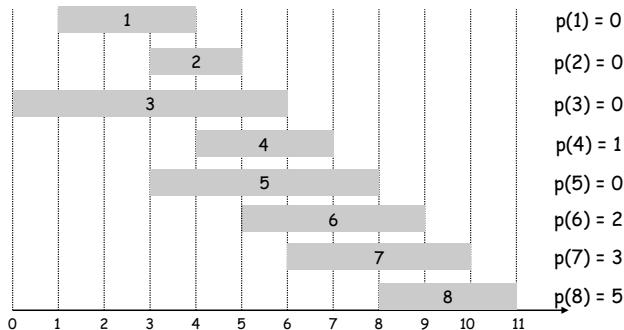
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Recall that greedy algorithm that works perfectly (optimally) if all the weights are 1. The greedy algorithm can fail spectacularly if arbitrary weights are allowed.

# BACK TO WEIGHTED INTERVAL SCHEDULING

Label (and sort) jobs by finishing time ( $f_1 \leq f_2 \leq \dots \leq f_n$ ).  
Define  $p(j)$  to be the largest index  $i < j$  such that job  $i$  is compatible with job  $j$ .



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# WEIGHTED INTERVAL SCHEDULING: BINARY CHOICE

$\text{OPT}(j)$  is the value of the optimal solution to the problem consisting of job request  $1, 2, \dots, j$ .

- **Case 1:** OPT selects job  $j$ .
  - then OPT can't include any of the jobs in the range  $p(j) + 1, p(j) + 2, \dots, j - 1$
  - OPT must include the optimal solution to the problem consisting of remaining compatible jobs  $1, 2, \dots, p(j)$
- **Case 2:** OPT does not select job  $j$ .
  - OPT must include the optimal solution to the problem consisting of remaining compatible jobs  $1, 2, \dots, j - 1$

$$\text{OPT}(j) = \begin{cases} 0 & \text{if } j = 0 \\ \max\{v_j + \text{OPT}(p(j)), \text{OPT}(j - 1)\} & \text{otherwise} \end{cases}$$

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# WEIGHTED INTERVAL SCHEDULING: BRUTE FORCE

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**Input:**  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$

**Sort** jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .

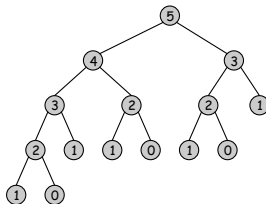
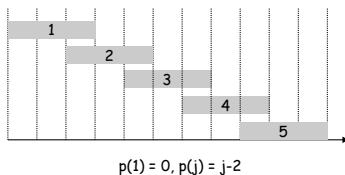
**Compute**  $p(1), p(2), \dots, p(n)$

```
Compute-Opt(j) {  
    if (j = 0)  
        return 0  
    else  
        return max( $v_j + \text{Compute-Opt}(p(j))$ ,  $\text{Compute-Opt}(j-1)$ )  
}
```

# WEIGHTED INTERVAL SCHEDULING: BRUTE FORCE IS BAD

The recursive algorithm fails spectacularly because of redundant sub-problems (it's an exponential algorithm)

- the number of recursive calls for a family of “layered” instances grows like the Fibonacci sequence

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# WEIGHTED INTERVAL SCHEDULING: MEMOIZATION

**Memoization:** store the results of each sub-problem in a cache; look them up as needed.

**Input:**  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$

**Sort** jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .

**Compute**  $p(1), p(2), \dots, p(n)$

**for**  $j = 1$  to  $n$

$M[j] = \text{empty}$   $\leftarrow$  global array

$M[j] = 0$

**M-Compute-Opt**( $j$ ) {

**if** ( $M[j]$  is empty)

$M[j] = \max(w_j + \text{M-Compute-Opt}(p(j)), \text{M-Compute-Opt}(j-1))$

**return**  $M[j]$

}

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# WEIGHTED INTERVAL SCHEDULING: MEMOIZATION IS GOOD

What is the running time of the memoized algorithm?  $O(n \lg n)$

- Sort by finish time:  $O(n \lg n)$
- Computing  $p(\cdot)$ :  $O(n)$  after sorting by start time
- $\text{M-Compute-Opt}(j)$ : each invocation takes  $O(1)$  time and either:
  - returns an existing value  $M[j]$
  - fills in one new entry  $M[j]$  and makes two recursive calls
- Define a progress measure  $\Phi$  as the number of nonempty entries in  $M[\cdot]$ .
  - initially  $\Phi = 0$ ; throughout  $\Phi \leq n$
  - a call to  $\text{M-Compute-Opt}(j)$  increases  $\Phi$  by at most 1;  $\Phi$  cannot be increased more than  $n$  times; there are at most  $2n$  recursive calls
- so the overall running time of  $\text{M-Compute-Opt}(n)$  is  $O(n)$

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# WEIGHTED INTERVAL SCHEDULING: FINDING A SOLUTION

This algorithm has only computed the optimal value (i.e., the weight of the optimal schedule). What if we want the solution itself?

```
Run M-Compute-Opt(n)
Run Find-Solution(n)

Find-Solution(j) {
    if (j = 0)
        output nothing
    else if ( $v_j + M[p(j)] > M[j-1]$ )
        print j
        Find-Solution(p(j))
    else
        Find-Solution(j-1)
}
```

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# BOTTOM-UP WEIGHTED INTERVAL SCHEDULING

To do this via dynamic programming, basically, we unwind the recursion...

**Input:**  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$

**Sort** jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .

**Compute**  $p(1), p(2), \dots, p(n)$

```
Iterative-Compute-Opt {  
    M[0] = 0  
    for j = 1 to n  
        M[j] = max( $v_j + M[p(j)]$ , M[j-1])  
}
```

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# WHEN DYNAMIC PROGRAMMING APPLIES

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For an optimization problem to be a good candidate for dynamic programming, it should exhibit:

- optimal substructure
- overlapping problems



A problem exhibits the optimal substructure property if the optimal solution contains within it optimal solutions to subproblems

- be careful, though, optimal substructure may also make a problem a good candidate for a greedy solution

We use the optimal substructure by then solving the subproblems bottom-up, combining the subproblem solutions as we move up.

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# OVERLAPPING SUBPROBLEMS

If the recursive solution is going to solve the same subproblem multiple times, then we're wasting computation

- this is in contrast to problems that generate unique subproblems at every split, for which divide and conquer is a good fit
- one alternative for overlapping subproblems is *memoization*
  - it's like dynamic programming, but it fills the table in using a top-down approach that's more like traditional recursion
  - you create a table to use in a recursive algorithm, and the recursive calls fill in the table as they return
  - results can then be just computed once and looked up thereafter
- more traditional dynamic programming is "bottom up"

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# SEGMENTED LEAST SQUARES

## Least Squares

- Foundational problem in statistic and numerical analysis
- Given  $n$  points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ , find a line  $y = ax + b$  that minimizes the sum of the squared error:

$$SSE = \sum_{i=1}^n (y_i - ax_i - b)^2$$

## Solution

The solution from calculus tells us that the minimum error is achieved when:

$$a = \frac{n \sum_i x_i y_i - (\sum_i x_i)(\sum_i y_i)}{n \sum_i x_i^2 - (\sum_i x_i)^2}$$

and

$$b = \frac{\sum_i y_i - a \sum_i x_i}{n}$$

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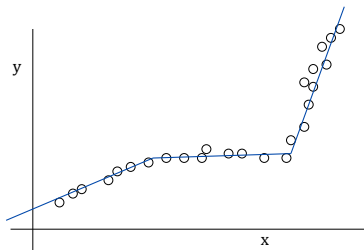
# SEGMENTED LEAST SQUARES

## Segmented Least Squares

- Points lie roughly on a sequence of several line segments
- Given  $n$  points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$  with  $x_1 < x_2 < \dots < x_n$ , find a sequence of lines that minimizes  $f(x)$

## Question

What is a reasonable choice for  $f(x)$  to balance accuracy (goodness of fit) and parsimony (number of lines)

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# SEGMENTED LEAST SQUARES

## The Tradeoff

We have to tradeoff the number of lines for the summed error

Find a sequence of lines that minimizes

- The sum of the sums of the squared errors  $E$  in each segment
- The number of lines  $L$

This results in a tradeoff function  $E + cL$  for some constant  $c$

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# DYNAMIC PROGRAMMING: MULTIWAY CHOICE

## Notation

- $OPT(j)$  = minimum cost for points  $p_1, p_2, \dots, p_j$
- $e(i, j)$  = minimum sum of squares for points  $p_i, p_{i+1}, \dots, p_j$

## To Compute $OPT(j)$

- Last segment uses points  $p_i, p_{i+1}, \dots, p_j$  for some  $i$
- Cost =  $e(i, j) + c + OPT(i - 1)$

$$OPT(j) = \begin{cases} 0 & \text{if } j = 0 \\ \min_{1 \leq i \leq j} (e(i, j) + c + OPT(i - 1)) & \text{otherwise} \end{cases}$$

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# SEGMENTED LEAST SQUARES: ALGORITHM

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```
INPUT:  $n, p_1, \dots, p_n, c$ 

Segmented-Least-Squares() {
     $M[0] = 0$ 
    for  $j = 1$  to  $n$ 
        for  $i = 1$  to  $j$ 
            compute the least square error  $e_{ij}$  for
            the segment  $p_i, \dots, p_j$ 

        for  $j = 1$  to  $n$ 
             $M[j] = \min_{1 \leq i \leq j} (e_{ij} + c + M[i-1])$ 

    return  $M[n]$ 
}
```

## Running Time

- Bottleneck: computing  $e(i, j)$  for  $O(n^2)$  pairs,  $O(n)$  per pair using the previous formula

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# THE KNAPSACK PROBLEM

## Knapsack Problem

- Given  $n$  objects and a “knapsack”
- Item  $i$  weighs  $w_i > 0$  kilograms and has value  $v_i > 0$
- Knapsack has a capacity of  $W$  kilograms
- Goal: fill the knapsack so as to maximize the total value

## Greedy?

Remember the greedy algorithm we explored was not optimal...

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# KNAPSACK: A FALSE START

Definition:  $OPT(i)$  is the max profit for the subset of items  $1, \dots, i$

- Case 1:  $OPT$  does not select item  $i$ 
  - $OPT$  selects the best of  $1, 2, \dots, i - 1$
- Case 2:  $OPT$  selects item  $i$ 
  - accepting item  $i$  does not immediately imply that we will have to reject other items
  - without knowing what other items were selected before  $i$ , we don't even know if we have enough room for  $i$

## Conclusion

We need more subproblems!

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# DYNAMIC PROGRAMMING: ADDING A VARIABLE

Definition:  $OPT(i, w)$  is the max profit of items  $1, \dots, i$  with weight limit  $w$

- Case 1:  $OPT$  does not select item  $i$ 
  - $OPT$  selects best of  $\{1, 2, \dots, i - 1\}$  using weight limit  $w$
- Case 2:  $OPT$  selects item  $i$ 
  - new weight limit  $w - w_i$
  - $OPT$  selects best of  $\{1, 2, \dots, i - 1\}$  using this new weight limit

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i - 1, w) & \text{if } w_i > w \\ \max(OPT(i - 1, w), v_i + OPT(i - 1, w - w_i)) & \text{otherwise} \end{cases}$$

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# KNAPSACK: BOTTOM UP

```
Input:  $n, w_1, \dots, w_N, v_1, \dots, v_N$ 

for  $w = 0$  to  $W$ 
     $M[0, w] = 0$ 

for  $i = 1$  to  $n$ 
    for  $w = 1$  to  $W$ 
        if ( $w_i > w$ )
             $M[i, w] = M[i-1, w]$ 
        else
             $M[i, w] = \max \{M[i-1, w], v_i + M[i-1, w-w_i]\}$ 

return  $M[n, W]$ 
```

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## Running Time

- $\theta(nW)$
- But I swear I've heard that the knapsack algorithm is NP-Complete!
- What does that mean?
- How is that possible? Didn't I just write down a polynomial time algorithm?
- Nope. It's "pseudo-polynomial"

## An Approximation Algorithm

There is a polynomial time algorithm that produces a feasible solution whose value is within 0.01% of the optimal.

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# THE COIN CHANGING PROBLEM

## The Problem

You are given  $k$  denominations of coins,  $d_1, d_2, \dots, d_k$  (all integers). Assume  $d_1 = 1$  so it is always possible to make change for any amount of money. We want to find an algorithm that makes change for an amount of money  $n$  using as few coins as possible.

## First Question

The coin changing problem exhibits optimal substructure. Consider any optimal solution to make change for  $n$  cents using our  $k$  denominations of coins. Consider breaking that solution into two different pieces along any coin boundary. Suppose that one half of the solution amounts to  $b$  cents and the other half to  $n - b$  cents. Then the solution to the each half must be an optimal way to make change for  $b$  cents (or  $n - b$  cents) using the same  $k$  denominations of coins. **Prove this.**

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## Second Question

Let  $C[p]$  be the minimum number of coins of the  $k$  denominations that sum to  $p$  cents. **Recursively define the value of the optimal solution.** To get you started, there must exist some “first coin”  $d_i$ , where  $d_i \leq p$ .

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# THE COIN CHANGING PROBLEM

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## Third Question

Provide an algorithm (e.g., pseudocode) to compute the value of the optimal solution bottom up. Provide another algorithm to construct the optimal solution from the computed information.

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## The Algorithm to Reconstruct the Solution

MAKECHANGE( $S, d, n$ )

```
1  while  $n > 0$ 
2      Print  $S[n]$ 
3       $n \leftarrow n - d[S[n]]$ 
```

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## Fourth Question

What is the running time of your algorithm?

## Running Time

CHANGE is  $\Theta(nk)$  (which is pseudo-polynomial in  $k$ , the input size and dependent on  $n$ , just like knapsack was dependent on  $W$ ). MAKECHANGE is  $O(n)$  since  $n$  is reduced by at least 1 in every iteration of the **while** loop. The total space requirement is  $\Theta(n)$

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# RNA SECONDARY STRUCTURE

## Secondary Structure

A set of pairs  $S = \{(b_i, b_j)\}$  that satisfy:

- **Watson and Crick:**  $S$  is a matching and each pair in  $S$  is a Watson-Crick component (matching A to U or C to G)
- **No Sharp Turns:** The ends of each pair are separated by at least 4 intervening bases. That is, if  $(b_i, b_j) \in S$ , then  $i < j - 4$ .
- **Non-Crossing:** If  $(b_i, b_j)$  and  $(b_k, b_l)$  are two pairs in  $S$ , then we cannot have  $i < k < j < l$

## Free Energy

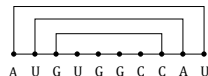
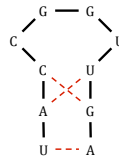
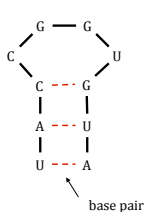
The usual hypothesis is that an RNA molecule will form the secondary structure that has the optimum total *free energy*, which we approximate by the number of base pairs.

## Goal

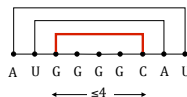
Given an RNA molecule  $B = b_1 b_2 \dots b_n$ , find a secondary structure  $S$  that maximizes the number of base pairs.

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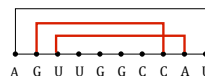
Examples.



ok



sharp turn



crossing

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## Notation

$OPT(i, j)$  is the maximum number of base pairs in a secondary structure of the substring  $b_i b_{i+1} \dots b_j$

- Case 1: if  $i \geq j - 4$ 
  - $OPT(i, j) = 0$  by the no sharp turns condition
- Case 2: Base  $b_j$  is not involved in a pair.
  - $OPT(i, j) = OPT(i, j - 1)$
- Case 3: Base  $b_j$  pairs with  $b_t$  for some  $i \leq t < j - 4$ 
  - The non crossing constraint decouples the resulting subproblems
  - $OPT(i, j) = 1 + \max_t (OPT(i, t - 1) + OPT(t + 1, j - 1))$  such that  $i \leq t < j - 4$  and  $(b_j, b_t)$  is a Watson and Crick pair

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# BOTTOM UP DYNAMIC PROGRAMMING OVER INTERVALS

## Question

What order to solve the subproblems?

## Answer

Do the shortest intervals first.

```
RNA( $b_1, \dots, b_n$ ) {  
  for  $k = 5, 6, \dots, n-1$   
    for  $i = 1, 2, \dots, n-k$   
       $j = i + k$   
      Compute  $M[i, j]$   
  
  return  $M[1, n]$   using recurrence  
}
```

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# DYNAMIC PROGRAMMING SUMMARY

## Recipe

- Characterize the structure of the problem
- Recursively define the value of an optimal solution
- Compute the value of the optimal solution
- Construct the optimal solution from computed information

## Dynamic Programming Techniques

- Binary choice: weighted interval scheduling.
- Multi-way choice segmented least squares.
- Adding a new variable: knapsack
- Dynamic programming over intervals: RNA secondary structure

## Top-down vs. Bottom-up?

Different people have different intuitions

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# LONGEST COMMON SUBSEQUENCE

This is a common problem in biological applications: find the longest common sequence of ACTG in two different pieces of DNA

- useful in finding similar proteins, etc.

Given a sequence  $X = \langle x_1, x_2, \dots, x_m \rangle$ , a sequence  $Z = \langle z_1, z_2, \dots, z_k \rangle$  is a **subsequence** of  $X$  if there exists a strictly increasing sequence  $\langle i_1, i_2, \dots, i_k \rangle$  of indices of  $X$  such that for all  $j = 1, 2, \dots, k$ ,  $x_{i_j} = z_j$ .

(For example  $Z = \langle B, C, D, B \rangle$  is a subsequence of  $X = \langle A, B, C, B, D, A, B \rangle$ .)

Our goal, is, given two sequences  $X$  and  $Y$ , find the longest common subsequence of the two sequences.

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# CHARACTERIZING THE LONGEST COMMON SUBSEQUENCE

## Theorem: Optimal Substructure of an LCS

Let  $X = \langle x_1, x_2, \dots, x_m \rangle$  and  $Y = \langle y_1, y_2, \dots, y_n \rangle$  be sequences and let  $Z = \langle z_1, z_2, \dots, z_k \rangle$  be any LCS of  $X$  and  $Y$ .

- ❶ If  $x_m = y_n$ , then  $z_k = x_m = y_n$  and  $Z_{k-1}$  is an LCS of  $X_{m-1}$  and  $Y_{n-1}$ .
- ❷ If  $x_m \neq y_n$ , then  $z_k \neq x_m$  implies that  $Z$  is an LCS of  $X_{m-1}$  and  $Y$ .
- ❸ if  $x_m \neq y_n$ , then  $z_k \neq y_n$  implies that  $Z$  is an LCS of  $X$  and  $Y_{n-1}$ .

## Proof (Part I)

If  $z_k \neq x_m$ , then we could append  $x_m = y_n$  to  $Z$  to get a common subsequence with length  $k + 1$ , which would contradict the premise. So  $z_k = x_m = y_n$ . And therefore  $Z_{k-1}$ , the prefix of  $Z$  with  $z_k$  removed is a longest common subsequence of  $X_{m-1}$  and  $Y_{n-1}$ .

## Proof (Parts II and III)

If  $z_k \neq x_m$ , then  $Z$  is a common subsequence of  $X_{m-1}$  and  $Y$ . If there were a common subsequence  $W$  of  $X_{m-1}$  and  $Y$  with length greater than  $k$ , then  $W$  would also be a subsequence of  $X$  and  $Y$ , which contradicts the premise that  $Z$  was an LCS.

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# A RECURSIVE SOLUTION TO LCS

If  $x_m = y_n$ , there is one subproblem to solve: find the LCS of  $X_{m-1}$  and of  $Y_{n-1}$ . If  $x_m \neq y_n$ , then there are two subproblems to solve: find the LCS of  $X_{m-1}$  and  $Y$  and the LCS of  $X$  and  $Y_{n-1}$ .

$$c[i, j] = \begin{cases} 0 & \text{if } i = 0 \text{ or } j = 0 \\ c[i - 1, j - 1] + 1 & \text{if } i, j > 0 \text{ and } x_i = y_j \\ \max(c[i, j - 1], c[i - 1, j]) & \text{if } i, j > 0 \text{ and } x_i \neq y_j \end{cases}$$

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# COMPUTING THE LENGTH OF AN LCS

We use dynamic programming to compute solutions to the  $\Theta(mn)$  subproblems in a bottom-up fashion

- we create a table  $c[0 \dots m, 0 \dots n]$  and compute its entries in row major order (filling in the first row, left to right, then the second row, etc.)
- we keep a second table  $b[0 \dots m, 0 \dots n]$  whose entries point to the table entry that corresponds to the best subproblem for the problem  $b[i, j]$  (to help in reconstructing the optimal solution)

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# COMPUTING THE LENGTH OF AN LCS (CONT.)

LCS-LENGTH( $X, Y$ )

```

1   $m \leftarrow \text{length}[X]$ 
2   $n \leftarrow \text{length}[Y]$ 
3  for  $i \leftarrow 1$  to  $m$ 
4      do  $c[i, 0] \leftarrow 0$ 
5  for  $j \leftarrow 1$  to  $n$ 
6      do  $c[0, j] \leftarrow 0$ 
7  for  $i \leftarrow 1$  to  $m$ 
8      do for  $j \leftarrow 1$  to  $n$ 
9          do if  $x_i = y_j$ 
10             then  $c[i, j] \leftarrow c[i - 1, j - 1] + 1$ 
11                  $b[i, j] \leftarrow \nwarrow$ 
12             else if  $c[i - 1, j] \geq c[i, j - 1]$ 
13                 then  $c[i, j] \leftarrow c[i - 1, j]$ 
14                      $b[i, j] \leftarrow \uparrow$ 
15                 else  $c[i, j] \leftarrow c[i, j - 1]$ 
16                      $b[i, j] \leftarrow \leftarrow$ 
17  return  $c$  and  $b$ 

```

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# AN EXAMPLE

		$j$	0	1	2	3	4	5	6
		$y_j$		<b>B</b>	<i>D</i>	<b>C</b>	<i>A</i>	<b>B</b>	<b>A</b>
0	$x_i$		0	0	0	0	0	0	0
1	<i>A</i>		0	↑ 0	↑ 0	↑ 0	↖ 1	← 1	↖ 1
2	<b>B</b>		0	↖ 1	← 1	← 1	↑ 1	↖ 2	← 2
3	<b>C</b>		0	↑ 1	↑ 1	↖ 2	← 2	↑ 2	↑ 2
4	<b>B</b>		0	↖ 1	↑ 1	↑ 2	↑ 2	↖ 3	← 3
5	<i>D</i>		0	↑ 1	↖ 2	↑ 2	↑ 2	↑ 3	↑ 3
6	<b>A</b>		0	↑ 1	↑ 2	↑ 2	↖ 3	↑ 3	↖ 4
7	<i>B</i>		0	↖ 1	↑ 2	↑ 2	↑ 3	↖ 4	↑ 4

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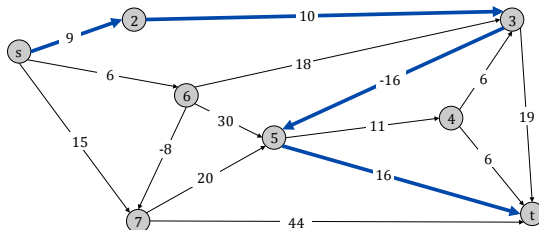
# SHORTEST PATHS

## Shortest Path Problem

Given a direct graph  $G = (V, E)$  with the edge weights  $c_{vw}$  (which can include negative edge weights), find the shortest path from node  $s$  to node  $t$ .

## Example

Nodes represent agents in a financial setting and  $c_{vw}$  is the cost of a transaction in which we buy from agent  $v$  and sell immediately to  $w$



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## Observation

If some path from  $s$  to  $t$  contains a negative cost cycle, there does not exist a shortest  $s$ - $t$  path; otherwise there exists a path that is a simple path.

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# SHORTEST PATHS AND DYNAMIC PROGRAMMING

Definition:  $OPT(i, v)$  is the length of the shortest  $v-t$  path  $P$  using at most  $i$  edges.

- Case 1:  $P$  uses at most  $i - 1$  edges
  - $OPT(i, v) = OPT(i - 1, v)$
- Case 2:  $P$  uses exactly  $i$  edges
  - if  $(v, w)$  is the first edge, the  $OPT$  uses  $(v, w)$ , and then selects the best  $w-t$  path using at most  $i - 1$  edges

$$OPT(i, v) = \begin{cases} 0 & \text{if } i = 0 \\ \min(OPT(i - 1, v), \min_{(v, w) \in E} (OPT(i - 1, w) + c_{vw})) & \text{otherwise} \end{cases}$$

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See the book for the Bellman-Ford algorithm implementation and a discussion of the running time.

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# DISTANCE VECTOR ROUTING

## Communication Network

- nodes = routers
- edges = direct communication link
- cost of edge = delay on link (which is naturally non-negative, but we use Bellman-Ford algorithm anyway!)

## Dijkstra's Algorithm

Requires global information of network

## Bellman-Ford

Uses only local knowledge of neighboring nodes

## Synchronization

We don't expect the routers to run in lock-step; Bellman-Ford can tolerate asynchronous routing updates and can be shown to still converge on the correct shortest path.

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# DISTANCE VECTOR PROTOCOL

## Distance Vector Protocol

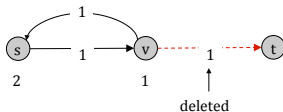
- Each router maintains a vector of shortest path lengths to every other node (distances) and the first hop on each path (directions)
- Algorithm: each router performs  $n$  separate computations, one for each potential destination node
- “Routing by rumor”

## Examples

RIP, Xerox XNS RIP, Novell's IPX RIP, Cisco's IGRP, DEC's DNA Phase IV, AppleTalk's RTMP

## Caveat

Edge costs may change during the algorithm; inconsistent router state can lead to the **counting to infinity** problem

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## Link State Routing

- Each router stores the entire path (not just the distance and next hop)
- Based on Dijkstra's algorithm
- Avoids the counting to infinity problem
- Requires significantly more storage

## Examples

Border Gateway Protocol (BGP), Open Shortest Path First (OSPF)

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