CSE3081 Design and Analysis of Algorithms

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Chapter 14. Dynamic Programming



Introduction

- Dynamic programming solves problems by combining the solutions to subproblems.
 - similar to the divide-and-conquer method
 - "programming" refers to a tabular method, not to writing computer code
- Difference with divide-and-conquer
 - dynamic programming applies when the subproblems overlap
 - subproblems share subsubproblems
- When the subproblems overlap, a divide-and-conquer algorithm does more work than necessary
 - repeatedly solves common subsubproblems
- A dynamic programming algorithm solves each subsubproblem just once and then saves its answer in a table, avoiding recomputing.



Applications

- Dynamic programming typically applies to optimization problems.
 - Such problems can have many possible solutions.
 - Each solution has a value, and you want to find a solution with the optimal (minimum or maximum) value.
 - There may be multiple "optimal" solutions that achieve the optimal value.



Steps

- 1. Characterize the structure of an optimal solution.
- 2. Recursively define the value of an optimal solution.
- 3. Compute the value of an optimal solution, typically in a bottom-up fashion.
- 4. Construct an optimal solution from computed information
 - this step can be omitted if we are only interested in the value of the optimal solution, and not the solution itself.



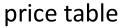
14.1 Rod Cutting



Introduction

- Serling Enterprises buys long steel rods and cuts them into shorter rods, which it then sells.
- Each cut is free.
- The company has a table giving, for i=1,2,..., the price p_i in dollars that they charge for a rod of length i inches.
- The length of each rod in inches is always an integer.
- The management of Serling Enterprises wants to know the best way to cut up the rods.

length i	1	2	3	4	5	6	7	8	9	10
price p_i	1	5	8	9	10	17	17	20	24	30

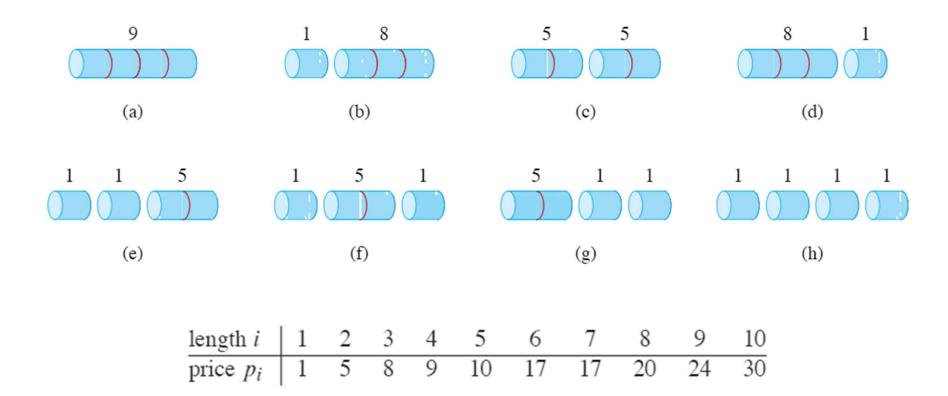






Example

• 8 possible ways of cutting up a rod of length 4.





The Rod-Cutting Problem

- Given a rod of length n inches and a table of prices p_i for $i=1,2,\ldots,n$, determine the maximum revenue r_n obtainable by cutting up the rod and selling the pieces.
- We can cut up a rod of length n in 2^{n-1} different ways.
 - We have an independent option of cutting or not cutting at distance i inches from the left end, for i=1,2,...,n-1.
- We will denote a decomposition into pieces using ordinary additive notation, such as 7 = 2 + 2 + 3.
- If an optimal solution cuts the rod into k pieces, for some $1 \le k \le n$, then optimal decomposition $n = i_1 + i_2 + \dots + i_k$ of the rod into pieces of lengths i_1, i_2, \dots, i_k provides the maximum corresponding revenue $r_n = p_{i_1} + p_{i_2} + \dots + p_{i_k}$.



The Rod-Cutting Problem: Example

Suppose this is the price table.

• Then,

- $r_1 = 1$ from solution 1 = 1 (no cuts),
- $r_2 = 5$ from solution 2 = 2 (no cuts),
- $r_3 = 8$ from solution 3 = 3 (no cuts),
- $r_4 = 10$ from solution 4 = 2 + 2,
- $r_5 = 13$ from solution 5 = 2 + 3,
- $r_6 = 17$ from solution 6 = 6 (no cuts),
- $-r_7 = 18$ from solution 7 = 1 + 6 or 7 = 2 + 2 + 3,
- $r_8 = 22$ from solution 8 = 2 + 6,
- $r_9 = 25$ from solution 9 = 3 + 6,
- $r_{10} = 30$ from solution 10 = 10 (no cuts)



The Rod-Cutting Problem: Insights

- Generally, we can express the values r_n for $n \ge 1$ in terms of optimal revenues from shorter rods.
- $r_n = \max\{p_n, r_1 + r_{n-1}, r_2 + r_{n-2}, \dots, r_{n-1} + r_1\}$
- The first argument p_n corresponds to making no cuts at all.
- In other cases, we make an initial cut of the rod into two pieces of size i and n-i for each i=1,2,3,...,n-1.
- Then, we can obtain optimal revenues r_i and r_{n-i} from those two pieces.



The Rod-Cutting Problem: Insights

- For each *i*, you make the first cut at place *i*.
- The overall optimal solution for this case incorporates optimal solutions to the two resulting subproblems.
 - maximizing revenue from each of those two pieces.

Optimal substructure

 optimal solutions to a problem incorporate optimal solutions to related subproblems, which you may solve independently.



The Rod-Cutting Problem: Insights

- We can view the decomposition as follows:
 - The left piece of length i
 - The right piece of length n-1
- The left piece can no longer be cut.
- We can only cut the right piece.
- This view does not limit the ways of cutting the rods.
- Then, we can make a simpler version of the equation for calculating r_n .
- $r_n = \max\{p_i + r_{n-i}: 1 \le i \le n\}$
 - When i=n, it means that we are not cutting the rod at all. We assume $r_0=0$.
- Now, an optimal solution embodies the solution to only one related subproblem rather than two.



The Rod-Cutting Problem: Divide-and-Conquer?

CUT-ROD: top-down and recursive

```
CUT-ROD(p, n)

1 if n == 0

2 return 0

3 q = -\infty

4 for i = 1 to n

5 q = \max\{q, p[i] + \text{CUT-ROD}(p, n - i)\}

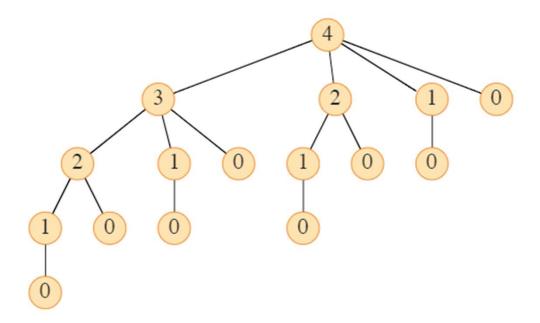
6 return q
```

Is this efficient?



The Rod-Cutting Problem: Divide-and-Conquer?

- CUT-ROD is inefficient because it is recursively called over and over again with the same parameter values. It solves the same problem repeatedly.
- Recursion tree





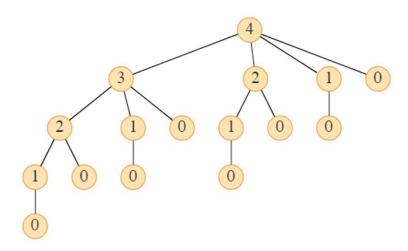
The Rod-Cutting Problem: Running Time of CUT-ROD

- T(n) denotes the total number of calls made to CUT-ROD(p, n) for a particular value of n.
- It is equal to the number of nodes in a subtree whose root is labeled n in the recursion tree.
- T(0) = 1, $T(n) = 1 + \sum_{j=0}^{n-1} T(j)$
- The initial 1 is for the call at the root.
- The term T(j) counts the number of calls (including recursive calls) due to call CUT-ROD(p, n-i), where j=n-i.
- Solving this recurrence relation, $T(n) = 2^n = \Theta(2^n)$.



The Rod-Cutting Problem: Running Time of CUT-ROD

- Conceptual explanation of why $T(n) = \Theta(2^n)$
 - A rod of length n has n-1 potential locations to cut.
 - Each possible way to cut up the rod makes a cut at some subset of these n-1 locations (including the empty set which corresponds to no cut.)
 - There are 2^{n-1} subsets.
 - Each leaf in the recursion tree corresponds to one possible way to cut up the rods. There are 2^{n-1} leaves.
 - Each node in the tree corresponds to a recursion call to CUT-ROD.
 - Thus, at least 2^{n-1} recursion calls should be made.





- Instead of solving the same problems repeatedly, arrange for each subproblem to be solved only once.
- The first time we solve a subproblem, we save its solution in a table.
- If we need to refer to this subproblem's solution again later, just look it up. Don't recompute it.
- We need memory to store solutions, so dynamic programming can be viewed as a time-memory trade-off.



- Top-down approach
 - We write the procedure recursively in a natural manner, but modify to save the result of each subproblem.
 - When solving a subproblem, the procedure first checks whether the solution has been previously saved.
 - If the solution is saved, the procedure simply returns the value.
 - If the solution is not saved, the procedure computes the value and saves it.



Top-down approach

```
MEMOIZED-CUT-ROD(p, n)
1 let r[0:n] be a new array // will remember solution values in r
2 for i = 0 to n
3 	 r[i] = -\infty
4 return MEMOIZED-CUT-ROD-AUX(p, n, r)
MEMOIZED-CUT-ROD-AUX(p, n, r)
1 if r[n] \ge 0 // already have a solution for length n?
 return r[n]
3 if n == 0
q = 0
5 else q = -\infty
      for i = 1 to n // i is the position of the first cut
          q = \max\{q, p[i] + \text{MEMOIZED-CUT-ROD-AUX}(p, n - i, r)\}
           // remember the solution value for length n
 r[n] = q
  return q
```



- Bottom-up approach
 - We solve the subproblems in size order, smallest first, storing the solution to each subproblem when it is first solved.
 - In this way, when solving a particular problem, there are already saved solutions for all of the smaller subproblems its solution depends upon.
 - You need to solve each subproblem only once, and when you first see it, you
 have already solved all of its prerequisite subproblems.
- In most cases, top-down and bottom-up approach of dynamic programming have the same asymptotic running time.
- However, the bottom-up approach often has much better constant factors, since it has lower overhead for procedure calls.



Bottom-up approach

```
BOTTOM-UP-CUT-ROD(p, n)

1 let r[0:n] be a new array // will remember solution values in r

2 r[0] = 0

3 for j = 1 to n // for increasing rod length j

4 q = -\infty

5 for i = 1 to j // i is the position of the first cut

6 q = \max\{q, p[i] + r[j-i]\}

7 r[j] = q // remember the solution value for length j

8 return r[n]
```



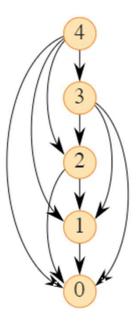
Running Time of Dynamic Programming Approaches

- Running time of BOTTOM-UP-CUT-ROD
 - We have a nested for loop.
 - Line 3-7 forms the outer for loop, where j is incremented from 1 to n.
 - Line 5-7 forms the inner for loop, which iterates j times.
 - Thus, the running time of BOTTOM-UP-CUT-ROD is $\Theta(n^2)$.
- Running time of MEMOIZED-CUT-ROD
 - Note that a recursive call to solve a previously solved problem returns immediately.
 - MEMOIZED-CUT-ROD-AUX solves each problem for sizes 0, 1, ..., n just once.
 - To solve a problem of size n, the for loop of lines 6-7 iterates n times.
 - The total number of iterations for this for loop, over all recursive calls of MEMOIZED-CUT-ROD, forms an arithmetic series $(1+2+\cdots+n)$, giving a total of $\Theta(n^2)$ iterations.



Subproblem Graph

- The subproblem graph shows how subproblems depend on one another.
- The subproblem graph for the rod-cutting problem.

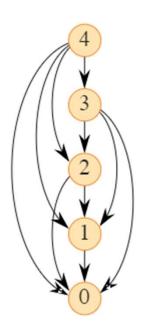


• For example, to solve a rod-cutting problem for length 4, we need the solutions to rod-cutting problem for lengths 0, 1, 2, 3.



Subproblem Graph

- Typically, the time to compute the solution to a subproblem is proportional to the degree (number of outgoing edges) of the corresponding vertex in the subproblem graph.
- The number of subproblems is equal to the number of vertices in the subproblem graph.
- In this common case, the running time of dynamic programming is linear in the number of vertices and edges.





Reconstructing a Solution

- The procedures MEMOIZED-CUT-ROD and BOTTOM-UP-CUT-ROD return the value of an optimal solution to the rod-cutting problem, but they do not return the solution itself: a list of piece sizes.
- In order to obtain the solution, the dynamic programming approach should record not only the optimal value but also the choice for each subproblem.



Reconstructing a Solution

- A procedure that records the optimal solution as well as the optimal value.
 - When p[i] + r[j i] is the maximum, we record s[j] = i.

i	0	1	2	3	4	5	6	7	8	9	10
r[i]	0	1	5	8	10	13	17	18	22	25	30
s[i]		1	2	3	2	2	6	1	2	3	10



Reconstructing a Solution

Now that we have the solution, we can print the cut locations.

```
PRINT-CUT-ROD-SOLUTION(p, n)

1 (r, s) = \text{EXTENDED-BOTTOM-UP-CUT-ROD}(p, n)

2 while n > 0

3 print s[n] // cut location for length n

4 n = n - s[n] // length of the remainder of the rod
```

i	0	1	2	3	4	5	6	7	8	9	10
r[i]	0	1	5	8	10	13	17	18	22	25	30
s[i]		1	2	3	2	2	6	1	2	3	10

- When n = 10, the procedure just prints 10.
- When n = 7, the procedure prints 1 6.



14.2 Matrix-Chain Multiplication



Matrix-Chain Multiplication Problem: Introduction

• Given a sequence (chain) $< A_1, A_2, ..., A_n >$ of n matrices to be multiplied, where the matrices aren't necessarily square, the goal is to compute the product

- $A_1A_2A_3...A_n$,
- while minimizing the number of scalar multiplications.



Matrix-Chain Multiplication: Example

• If the chain of matrices is $\langle A_1, A_2, A_3, A_4 \rangle$, then you can parenthesize the product $A_1A_2A_3A_4$ in five distinct ways:

```
(A_1(A_2(A_3A_4))),

(A_1((A_2A_3)A_4)),

((A_1A_2)(A_3A_4)),

((A_1(A_2A_3))A_4),

(((A_1A_2)A_3)A_4).
```

• How we parenthesize a chain of matrices can have a significant impact on the cost of evaluating the product.



Cost of Multiplying Two Rectangular Matrices

• The procedure RECTANGULAR-MATRIX-MULTIPLY computes $C = C + A \cdot B$ for three matrices $A = (a_{ij})$, $B = (b_{ij})$, and $C = (C_{ij})$, where A is $p \times q$, B is $q \times r$, and C is $p \times r$.

```
RECTANGULAR-MATRIX-MULTIPLY (A, B, C, p, q, r)

1 for i = 1 to p

2 for j = 1 to r

3 for k = 1 to q

4 c_{ij} = c_{ij} + a_{ik} \cdot b_{kj}
```

• The running time of RECTANGULAR-MATRIX-MULTIPLY is dominated by the number of scalar multiplications in line 4, which is pqr.



Example: Order of Multiplication Affects Cost

- We have a chain of three matrices $\langle A_1, A_2, A_3 \rangle$.
- Suppose their dimensions are 10×100 , 100×5 , and 5×50 , respectively.
- Case 1: $((A_1A_2)A_3)$
 - $10 \cdot 100 \cdot 5 = 5000$ scalar multiplications to compute $A_1 A_2$, plus
 - $10 \cdot 5 \cdot 50 = 2500$ scalar multiplications to multiply this matrix by A_3 .
 - Total: 7500 scalar multiplications
- Case 2: $(A_1(A_2A_3))$
 - $100 \cdot 5 \cdot 50 = 25000$ scalar multiplications to compute A_2A_3 , plus
 - $10 \cdot 100 \cdot 50 = 50000$ scalar multiplications to multiply A_1 by this matrix
 - Total: 75000 scalar multiplications



Problem Definition

- Matrix-chain multiplication problem
- Given a chain $\langle A_1, A_2, ..., A_n \rangle$ of n matrices, where for i=1,2,...,n, matrix A_i has dimension $p_{i-1} \times p_i$, fully parenthesize the product $A_1A_2 ... A_n$ in a way that minimizes the number of scalar multiplications. The input is the sequence of dimensions $\langle p_0, p_1, ..., p_n \rangle$.
- This problem does not entail actually multiplying matrices. The goal is only to determine an order for multiplying matrices that has the lowest cost.
- Typically, the time invested in determining this optimal order is more than paid for by the time saved later on when actually performing the matrix multiplications.
 - 7,500 vs. 75,000 scalar multiplications



Counting the Number of Parenthesizations

- Can't we just exhaustively check all possible parenthesizations?
- To find out, let us denote the number of alternative parenthesizations of sequence of n matrices by P(n).
- When n=1, there is just one matrix, so there is only one way to fully parenthesize the matrix product.
- When $n \ge 2$, a fully parenthesized matrix product is the product of two fully parenthesized matrix subproducts, and the split between the two subproducts may occur between kth and (k+1)st matrices for any $k=1,2,\ldots,n-1$.



Counting the Number of Parenthesizations

We obtain the recurrence

$$P(n) = \begin{cases} 1 & \text{if } n = 1, \\ \sum_{k=1}^{n-1} P(k)P(n-k) & \text{if } n \ge 2. \end{cases}$$

- When we solve the recurrence, $P(n) = \Omega(2^n)$.
- Thus, exhaustively searching all possible parenthesizations is not efficient.



Applying Dynamic Programming

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



Step 1: Structure of an Optimal Parenthesization

- Let $A_{i:j}$, where $i \leq j$, denote the matrix that results from evaluating the product $A_i A_{i+1} \dots A_j$.
- If i < j, to parenthesize the product $A_i A_{i+1} \dots A_j$, the product must split between A_k and A_{k+1} for some integer k in the range $i \le k < j$.
- That is, we first compute the matrices $A_{i:k}$ and $A_{k+1:j}$, and then multiply them together to produce the final product $A_{i:j}$.
- The cost
 - The cost of computing the matrix $A_{i:k}$, plus
 - the cost of computing the matrix $A_{k+1:j}$, plus
 - the cost of multiplying them together.



Step 1: Structure of an Optimal Parenthesization

- When computing $A_{i:j}$, suppose the optimal split is to split the product between A_k and A_{k+1} . In this solution, the parenthesization of each subchain must be optimal parenthesization.
- The subchain $A_{i:k}$ should have the optimal parenthesization.
- The subchain $A_{k+1:j}$ should have the optimal parenthesization.
- To build an optimal solution, we split the problem into two subproblems (optimally parenthesizing $A_iA_{i+1}\dots A_k$ and $A_{k+1}A_{k+2}\dots A_j$), find optimal solutions to the two subproblem instances, and then combine these optimal subproblem solutions.
- To ensure that we've examined the optimal split, we must consider all possible splits.



Step 2: A Recursive Solution

- Given the input dimensions $\langle p_0, p_1, p_2, \dots, p_n \rangle$, an index pair i, j specifies a subproblem.
- Let m[i,j] be the minimum number of scalar multiplications needed to compute $A_{1:n}$ is thus m[1,n].
- We can define m[i, j] recursively as follows:
- $m[i,j] = m[i,k] + m[k+1,j] + p_{i-1}p_kp_j$
- Since we do not know the optimal k, we need to try all possible values of k, k = i, i + 1, ..., j 1.



Step 2: A Recursive Solution

• Recursive definition for the minimum cost of parenthesizing the product $A_iA_{i+1}\dots A_j$ becomes

$$m[i,j] = \begin{cases} 0 & \text{if } i = j, \\ \min\{m[i,k] + m[k+1,j] + p_{i-1}p_kp_j : i \le k < j\} & \text{if } i < j. \end{cases}$$

- To construct the optimal solution, we define s[i,j] to be a value k at which you split the product $A_iA_{i+1} \dots A_j$ in an optimal parenthesization.
- That is, s[i,j] equals a value k such that $m[i,j] = m[i,k] + m[k+1,j] + p_{i-1}p_kp_j$.



- There is one subproblem for each choice of i and j satisfying $1 \le i \le j \le n$.
- Thus, the number of subproblems is $\Theta(n^2)$.
- A recursive algorithm may encounter each subproblem many times in different branches of its recursion tree.
- This property of overlapping subproblems is the second hallmark of when dynamic programming applies.
 - The first hallmark is the optimal substructure.
- We will use a tabular, bottom-up approach to solve the problem.

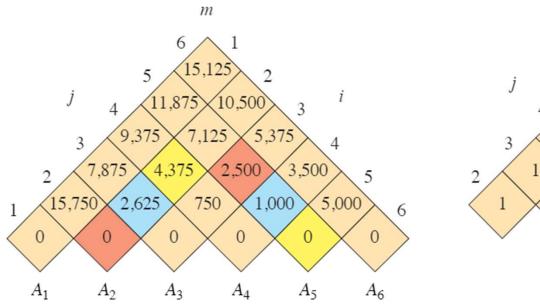


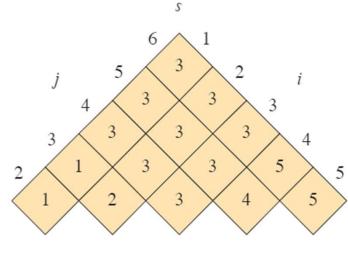
Procedure MATRIX-CHAIN-ORDER

```
MATRIX-CHAIN-ORDER (p, n)
  let m[1:n, 1:n] and s[1:n-1, 2:n] be new tables
                                 // chain length 1
2 for i = 1 to n
3 	 m[i,i] = 0
4 for l=2 to n
                    // l is the chain length
for i = 1 to n - l + 1 // chain begins at A_i
j = i + l - 1 // chain ends at A_i
          m[i,j] = \infty
          for k = i to j - 1 // try A_{i:k}A_{k+1:j}
8
              q = m[i,k] + m[k+1,j] + p_{i-1}p_kp_i
9
              if q < m[i, j]
10
                  m[i, j] = q // remember this cost
11
                  s[i, j] = k // remember this index
12
   return m and s
```



- Tables for *m* and *s*
 - The optimal number of scalar multiplications: 15,125





matrix	A_1	A_2	A_3	A_4	A_5	A_6
dimension	30×35	35×15	15 × 5	5×10	10×20	20×25



- Procedure MATRIX-CHAIN-ORDER has three nested for loops.
- Each loop index (l, i, and k) takes on at most n-1 values.
- Running time of the procedure: $\Theta(n^3)$
- The algorithm requires $\Theta(n^2)$ of memory space to store m and s tables.



Step 4: Constructing an Optimal Solution

Procedure for printing the optimal solution.

```
PRINT-OPTIMAL-PARENS (s, i, j)

1 if i == j

2 print "A"<sub>i</sub>

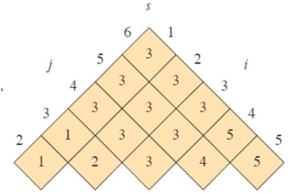
3 else print "("

4 PRINT-OPTIMAL-PARENS (s, i, s[i, j])

5 PRINT-OPTIMAL-PARENS (s, s[i, j] + 1, j)

6 print ")"
```

- Example
 - If we have the s table as in the right:
 - the optimal solution is $((A_1(A_2A_3))((A_4A_5)A_6))$.





14.4 Longest Common Subsequence



Introduction

- Biological applications often need to compare the DNA of two (or more) different organisms.
- A strand of DNA consists of a string of molecules called bases, where the possible bases are adenine, cytosine, guanine, and thymine.
- Representing each of these bases by its initial letter, we can express a strand of DNA as a string over the 4-element set {A, C, G, T}.
- For example, the DNA of one organism may be
 - $S_1 = ACCGGTCGAGTGCGCGGAAGCCGGCCGAA$
- and the DNA of another organism may be
 - $S_2 = GTCGTTCGGAATGCCGTTGCTCTGTAAA$



Introduction: DNA Strands

X = TCCCCGCTCTGCTCTGTCCGGTCACAGGACTTTTTGCCCTCTGTTCCCGGGTCCCTCAGGCGGCCACCCA GTGGGCACACTCCCAGGCGCCCCGGCCCCGCGCTCCCTCTGCCTTTCATTCCCAGCTGTCAAC ATCCTGGAAGCTTTGAAGCTCAGGAAAGAAGAAGAAATCCACTGAGAACAGTCTGTAAAGGTCCGTAGTGC CCCTCTATAAAAGCTCTGTGCATCCTGCCACTGAGGACTCCGAAGAGGTAGCAGTCTTCTGAAAGACTTC AACTGTGAGGACATGTCGTTCAGATTTGGCCAACATCTCATCAAGCCCTCTGTAGTGTTTCTCAAAACAG AACTGTCCTTCGCTCTTGTGAATAGGAAACCTGTGGTACCAGGACATGTCCTTGTGTGCCCGCTGCGGCC AGTGGAGCGCTTCCATGACCTGCGTCCTGATGAAGTGGCCGATTTGTTTCAGACGACCCAGAGAGTCGGG ACAGTGGTGGAAAAACATTTCCATGGGACCTCTCTCACCTTTTCCATGCAGGATGGCCCCGAAGCCGGAC AGACTGTGAAGCACGTTCACGTCCATGTTCTTCCCAGGAAGGCTGGAGACTTTCACAGGAATGACAGCAT CTATGAGGAGCTCCAGAAACATGACAAGGAGGACTTTCCTGCCTCTTGGAGATCAGAGGAGGAAATGGCA GCAGAAGCCGCAGCTCTGCGGGTCTACTTTCAGTGACACAGATGTTTTTCAGATCCTGAATTCCAGCAAA ATGCAGTTTCTTCATCTCACCATCCTGTATTCTTCAACCAGTGATCCCCCACCTCGGTCACTCCAACTCC CTTAAAATACCTAGACCTAAACGGCTCAGACAGGCAGATTTGAGGTTTCCCCCTGTCTCCTTATTCGGCA

Y = ATGTTAACCAAGGAATGGATCTGTGTCGTTCCACGTTCGAAGGCCTTTTCTGATGAAATGAAGATAGGTT TCAACTCCACAGGTTATTGTGGTATGATCTTAACCAAAAATGATGAAGTTTTCTCCAAGATTACTGAAAA ACCTGAATTGATTAACGATATCTTATTGGAATGTGGTTTCCCAAACACTTCTGGTCAAAAACCAAACGAA TGATAGAAGTAATTTCTATATGTATATGTCTATTCAATTTTTATTTCTAATGACTTTGAAATTTTATATT TACTTATTATTAATATTGTTGTATTACTTCCTTGAAAAAATATGTCTAAAGAGTCCTAATTTGGATTTTC TTTTCCTCCTAACTTCACTGTCCTGCGCCTGCTTTCCTAACGCACCATCGCTAATACACCAGCTTTCATT GCTTGTTGCTGCGCTATTGCTCGCATGGAACGTTTTCAGTGCGTCATCATCCTGGGATAAACTAAAGACT AAGTCACCAGTTTCATTTGAGGCTTTTCTCTGCGTGTTGACAAAAGAGGACAGATCAACCATATCACCTG GTTCACCATGAGAATGTCCTTACTTAGTTGCGAATTTGGTTCTGATCTGCCTTCAGACTTGGAATCATTC ATTACTGTCATTACTTTTAGCCTATTCATTTCTTCCTTGCCATCAGGTACAGGGATTTGACCACAGAGT GTTGAAGGGGTGCATCGTCCGCTCTCGTAATAACCCTCCGATACTATTTTCATTGTTGGCACGTTGCACT GAAAAGGGCACTTGGCACTGTGCACTTTTAATGTTTTCATTTTTCATCATCATCATCATCATATGGCATTTCAT AATGTTGACTCTTGTAGTTGAAGATTGAGTTTATCGTGTTACTGTTTCGTCTACCTTTCATATTATCAAT CAGGTTGCGGTGTTGCATGTGGGAGATAGGACTGCCCGATCTTCTTCTTCTCCACTACACATCC TGTCTTTTATCGCTATCCAGCACACGATTGAGCTGTGAATTGCCGCACTTTTTAGGATAACCATCCTTGG



Introduction

- We compare two strands of DNA to determine how "similar" the two strands are, as some measure of how closely related the two organisms are.
- We can define similarity in many different ways.
- For example, we can say that the two DNA strands are similar if one is a substring of the other.
- Alternatively, we could say that two strands are similar if the number of changes needed to turn one into the other is small. (Edit Distance Problem)



Introduction

- Yet another way to measure similarity of strands S_1 and S_2 is by finding a third strand S_3 in which the bases in S_3 appear in each of S_1 and S_2 .
- These bases must appear in the same order, but not necessarily consecutively.
- The longer the strand S_3 we can find, the more similar S_1 and S_2 are.
- In our example, the longest strand S_3 is:
- $S_3 = GTCGTCGGAAGCCGGCCGAA$
- We call this notion of similarity the longest common subsequence (LCS).



Subsequence

- A subsequence of given sequence is just the given sequence with 0 or more elements left out.
- Formally, given a sequence $X = \langle x_1, x_2, ..., x_m \rangle$, another sequence $Z = \langle z_1, z_2, ..., z_k \rangle$ is a subsequence of X if there exists a strictly increasing sequence $\langle i_1, i_2, ..., i_k \rangle$ of indices of X such that for all j = 1, 2, ..., k, we have $x_{i_j} = z_j$.
- For example, $Z = \langle B, C, D, B \rangle$ is a subsequence of $X = \langle A, B, C, B, D, A, B \rangle$ with corresponding index sequence $\langle 2, 3, 5, 7 \rangle$.



Common Subsequence

- Given two sequences X and Y, we say that a sequence Z is a common subsequence of X and Y if Z is a subsequence of both X and Y.
- For example, if $X = \langle A, B, C, B, D, A, B \rangle$ and $Y = \langle B, D, C, A, B, A \rangle$, the sequence $\langle B, C, A \rangle$ is a common subsequence of both X and Y.
- There is a longer common subsequence: $\langle B, C, B, A \rangle$. This is the longest common subsequence of X and Y. Its length is 4.
 - X and Y have no common subsequence of length 5 or higher.



Longest Common Subsequence Problem

- The input is two sequences $X = \langle x_1, x_2, ..., x_m \rangle$ and $Y = \langle y_1, y_2, ..., y_n \rangle$, and the goal is to find a maximum-length common subsequence of X and Y.
- We solve this problem using dynamic programming.



LCS: brute-force approach?

- We can solve the LCS problem with a brute-force approach: enumerate all subsequences of X and check each subsequence to see whether it is a also a subsequence of Y, keeping track of the longest subsequence we find.
- Since each subsequence of X corresponds to a subset of the indices $\{1, 2, ..., m\}$ of X.
- Because X has 2^m subsequences, this approach requires exponential time \rightarrow inefficient.



Step 1: Characterizing a Longest Common Subsequence

- The LCS problem has an optimal-substructure property.
- We define the *i*th prefix of X, for i = 0, 1, ..., m, as $X_i = \langle x_1, x_2, ..., x_i \rangle$.
- For example, if $X = \langle A, B, C, B, D, A, B \rangle$, then $X_4 = \langle A, B, C, B \rangle$.
 - X_0 is the empty sequence.



Step 1: Characterizing a Longest Common Subsequence

- Theorem 14.1 (Optimal substructure of an LCS)
- Let $X = \langle x_1, x_2, ..., x_m \rangle$ and $Y = \langle y_1, y_2, ..., y_n \rangle$ be sequences, and let $Z = \langle z_1, z_2, ..., z_k \rangle$ be any LCS of X and Y.
- 1. 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} .
- 2. If $x_m \neq y_n$ and $z_k \neq x_m$, then Z is an LCS of X_{m-1} and Y.
- 3. If $x_m \neq y_n$ and $z_k \neq y_n$, then Z is an LCS of X and Y_{n-1} .

Proof of Theorem 14.1

Case 1

- If $z_k \neq x_m$, then we could append $x_m = y_n$ to Z to obtain a common subsequence of X and Y of length k+1, contradicting the supposition that Z is a longest common subsequence of X and Y. Thus, we must have $z_k = x_m = y_n$.
- Suppose for the purpose of contradiction that there exists a common subsequence W of X_{m-1} and Y_{n-1} with length greater than k-1. Then, appending $x_m=y_n$ to W produces a common subsequence of X and Y whose length is greater than k, which is a contradiction.

Case 2

- 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 common subsequence of X_m and Y, contradicting the assumption that Z is an LCS of X and Y.
- Case 3: the proof is symmetric to case 2.



Step 2: A Recursive Solution

- If $x_m = y_n$, we first find an LCS of X_{m-1} and Y_{n-1} . Then, we append $x_m = y_n$ to this LCS to obtain an LCS of X and Y.
- If $x_m \neq y_n$, we need to solve two subproblems:
 - finding an LCS of X_{m-1} and Y
 - finding an LCS of X and Y_{n-1}
- whichever of these two LCSs is longer is an LCS of X and Y.
- Let us define c[i,j] to be the length of an LCS of the sequences X_i and Y_j .
- Then, the recursive formula is:

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



Step 3: Computing the Length of an LCS

- The LCS problem has $\Theta(mn)$ distinct subproblems.
- Dynamic programming can compute the solutions bottom-up.

```
LCS-LENGTH(X, Y, m, n)
 1 let b[1:m, 1:n] and c[0:m, 0:n] be new tables
 2 for i = 1 to m
        c[i,0] = 0
 4 for j = 0 to n
 5 	 c[0, j] = 0
 6 for i = 1 to m // compute table entries in row-major order
        for j = 1 to n
            if x_i == y_i
                c[i, j] = c[i - 1, j - 1] + 1
                b[i, j] = "\\\"
10
         elseif c[i - 1, j] \ge c[i, j - 1]
11
             c[i,j] = c[i-1,j]
12
                b[i, j] = "\uparrow"
13
            else c[i, j] = c[i, j - 1]
14
                b[i, j] = "\leftarrow"
15
   return c and b
```



Step 3: Computing the Length of an LCS

• The c table and the b table

	j	0	1	2	3	4	5	6
i		y_j	B	D	C	A	B	A
0	x_i	0	0	0	0	0	0	0
1	A	0	1	↑	↑	\1	← 1	\1
2	B	0	1	←1	← 1	↑ 1	\	←2
3	C	0	1 1	1 1	× 2	←2	1 2	1 2
4	B	0	\1	1	1 2	↑ 2	3	-3
5	D	0	1 1	\		1 2	↑ 3	1 3
6	A	0	1	1 2	1 2	<u></u>	1 3	4
7	В	0	\1	1 1 2	2 ↑ 2 ↑ 2	1 3	4	1 4



Step 3: Computing the Length of an LCS

• The running time of procedure LCS-LENGTH is $\Theta(mn)$, since each table entry takes $\Theta(1)$ time to compute.



Step 4: Constructing and LCS

Using the b table, we can construct the longest common subsequence.

```
PRINT-LCS(b, X, i, j)

1 if i == 0 or j == 0

2 return // the LCS has length 0

3 if b[i, j] == \text{``\cdot'}

4 PRINT-LCS(b, X, i - 1, j - 1)

5 print x_i // same as y_j

6 elseif b[i, j] == \text{``\cdot'}

7 PRINT-LCS(b, X, i - 1, j)

8 else PRINT-LCS(b, X, i, j - 1)
```



Improving the Code

- In the LCS algorithm, we can eliminate the b table altogether.
- Each c[i,j] entry depends on only three other table entries: c[i-1,j-1], c[i-1,j], and c[i,j-1].
- Given the value of c[i,j], we can determine in O(1) time which of these three values was used to compute c[i,j], without inspecting table b.
- If we need only the length of an LCS, we can reduce the c table as well.
- We need to use only two rows of table c at a time: the row being computed and the previous row.



Extra: Other Problems



Problem: World Series Odds

- Dodgers and Yankees are playing the World Series in which either team needs to win n games first.
- Suppose that each team has a 50% chance of winning any particular game.
- Let p(i, j) be the probability that if Dodgers needs i games to win, and Yankees needs j games, Dodgers will eventually win the series.
- Ex: if n = 4, p(2, 3) = 11/16
- Compute p(i, j) for an arbitrary n. $(0 \le i, j \le n)$



World Series Odds – Divide-and-Conquer

Recursive formulation

$$P(i,j) = \begin{cases} 1, & \text{if } i = 0 \text{ and } j > 0 \\ 0, & \text{if } i > 0 \text{ and } j = 0 \end{cases}$$
$$\frac{P(i-1,j) + P(i,j-1)}{2}, & \text{if } i > 0 \text{ and } j > 0 \end{cases}$$

- Worst-case time complexity
 - T(n): maximum time taken by a call to p(i, j)
 - T(n) is exponential!
- What is the problem of this approach?

World Series Odds – Dynamic Programming

• Instead of computing the same problem repeatedly, fill in a table as suggested below.

4	1	15/16	13/16	21/32	1/2	
3	1	7/8	11/16	1/2	11/32	
2	1	3/4	1/2	5/16	3/16	
1	1	1/2	1/4	1/8	1/16	
0		0	0	0	0	
\int_{i}	0	1	2	3	4	

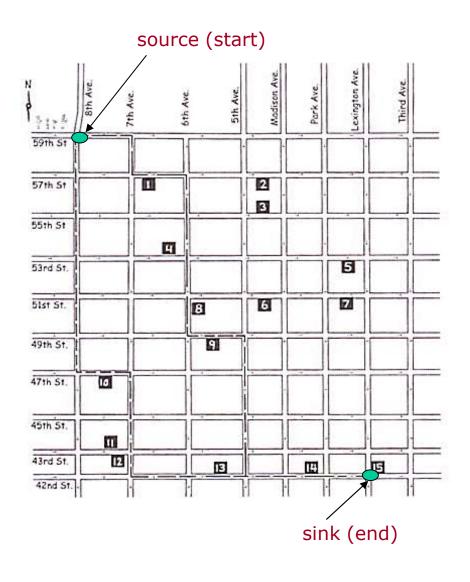
- Time Complexity: O(n²)
 - Much better than divide-and-conquer!



The Manhattan Tourist Problem

Problem

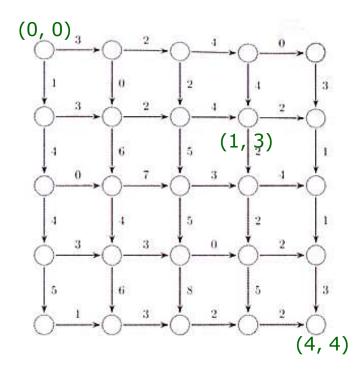
- Given two street corners in the borough of Manhattan in New York City, find the path between them with the maximum number of attractions, that is, a path of maximum overall weight.
- Assume that a tourist may move either EAST or SOUTH only.



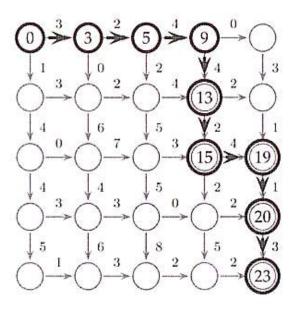


The Manhattan Tourist Problem

- Formal description
 - Given a weighted grid G of size (n, m) with two distinguished vertices, a source (0, 0) and a sink (n, m), find a <u>longest</u> path between them in its weighted graph



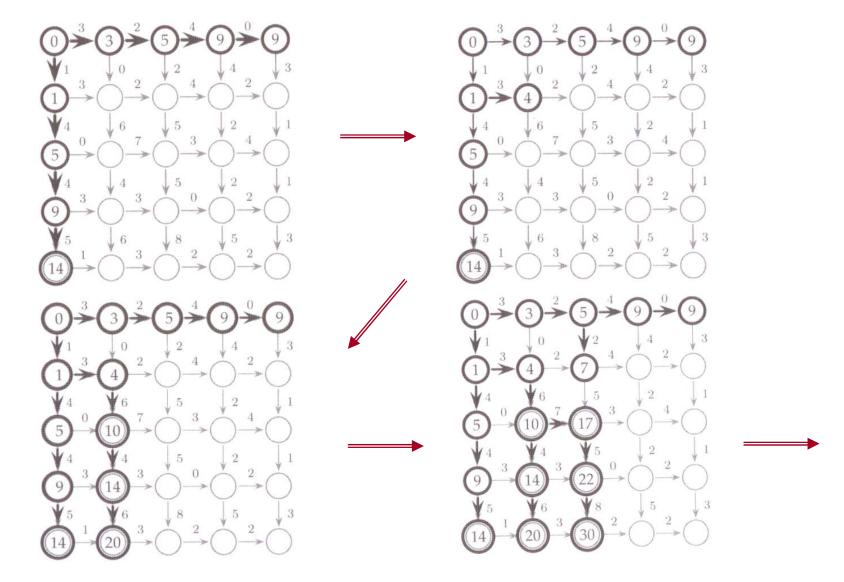
An example grid of size (4, 4)



A possible selection determined by a greedy approach

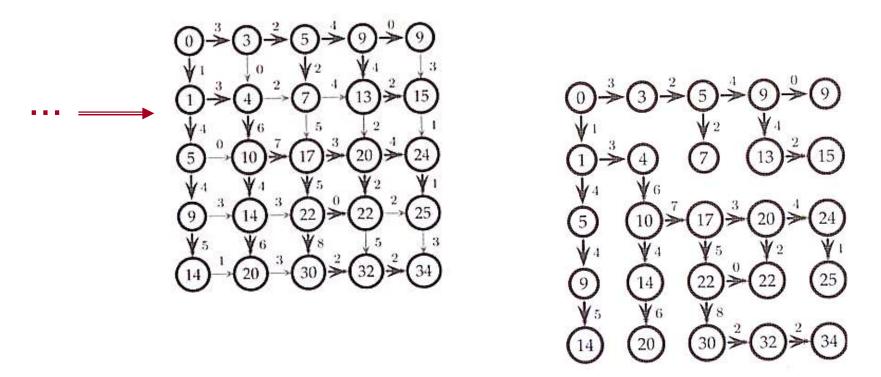


Dynamic Programming: A pictorial description





Dynamic Programming: A pictorial description (cont.)



Found longest path from source to every vertex in the grid



Dynamic Programming: Algorithm

Recursive relation

$$s_{i,j} \equiv$$
 the length of the longest path from $(0, 0)$ to (i, j)

$$s_{i,j} = \max \begin{cases} s_{i-1,j} + \text{weight between } (i-1,j) \text{ and } (i,j) \\ s_{i,j-1} + \text{weight between } (i,j-1) \text{ and } (i,j) \end{cases}$$

- Algorithm
- Time Complexity?

MANHATTANTOURIST(
$$\dot{\mathbf{w}}, \dot{\mathbf{w}}, n, m$$
)

$$1 \quad s_{0,0} \leftarrow 0$$

2 for
$$i \leftarrow 1$$
 to n

3
$$s_{i,0} \leftarrow s_{i-1,0} + \overset{\downarrow}{w}_{i,0}$$

4 for
$$j \leftarrow 1$$
 to m

5
$$s_{0,j} \leftarrow s_{0,j-1} + \overrightarrow{w}_{0,j}$$

6 for
$$i \leftarrow 1$$
 to n

7 for
$$j \leftarrow 1$$
 to m

8
$$s_{i,j} \leftarrow \max \begin{cases} s_{i-1,j} + \overset{\downarrow}{w}_{i,j} \\ s_{i,j-1} + \overset{\downarrow}{w}_{i,j} \end{cases}$$

9 return $s_{n,m}$



Edit Distance

• **Problem:** Given two strings $A = a_1 a_2 \cdots a_n$ and $B = b_1 b_2 \cdots b_m (n, m \ge 0)$, consider the problem of transforming A into B by either inserting a character into A, deleting a character from A, or replacing a character in A by another. Suppose that the costs per insertion, deletion, and replacement are c_{ins} , c_{del} , and c_{repl} , respectively. Then, find the minimum cost of transforming A into B.

$$A = S - N O W Y$$

 $B = S U N N - Y$

Let $\overline{C[i,j]}$ be the minimum cost to transform $A_i = a_1 a_2 \cdots a_i$ into $B_i =$ $b_1 b_2 \cdots b_j$ for $i \in [0, n]$ and $j \in [0, m]$.

$$C[0,0] = 0$$

 $C[i,0] = i * c_{del}, i \in [1,n]$
 $C[0,j] = j * c_{ins}, j \in [1,m]$

O(mn)

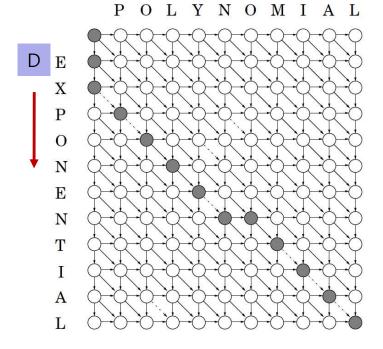
$$C[i,j] = \left\{ \begin{array}{l} C[i-1,j-1] & \text{if } a_i = b_j \\ C[i,j-1] + c_{ins} \\ C[i-1,j] + c_{del} \\ C[i-1,j-1] + c_{repl} \end{array} \right\} & \text{if } a_i \neq b_j \\ \end{array}, i \in [1,n], j \in [1,m]$$

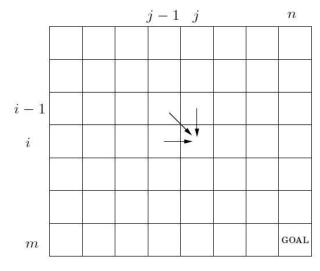


Edit Distance

• Example

$$A = E X P O N E N - T I A L B = - - P O L Y N O M I A L$$





		P	O	L	Y	N	O	M	I	A	L
	0	1	2	3	4	5	6	7	8	9	10
\mathbf{E}	1	1	2	3	4	5	6	7	8	9	10
X	2	2	2	3	4	5	6	7	8	9	10
P	3	2	3	3	4	5	6	7	8	9	10
O	4	3	2	3	4	5	5	6	7	8	9
N	5	4	3	3	4	4	5	6	7	8	9
\mathbf{E}	6	5	4	4	4	5	5	6	7	8	9
N	7	6	5	5	5	4	5	6	7	8	9
\mathbf{T}	8	7	6	6	6	5	5	6	7	8	9
I	9	8	7	7	7	6	6	6	6	7	8
A	10	9	8	8	8	7	7	7	7	6	7
L	11	10	9	8	9	8	8	8	8	7	6



End of Class

Questions?

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