CS 374: Algorithms & Models of Computation, Spring 2017

More Dynamic Programming

Lecture 14 March 9, 2017

What is the running time of the following?

Consider computing f(x, y) by recursive function + memoization.

$$f(x,y) = \sum_{i=1}^{x+y-1} x * f(x+y-i,i-1),$$

$$f(0,y) = y \qquad f(x,0) = x.$$

The resulting algorithm when computing f(n, n) would take: (A) O(n)(B) $O(n \log n)$ (C) $O(n^2)$ (D) $O(n^3)$

(E) The function is ill defined - it can not be computed.

Recipe for Dynamic Programming

- Develop a recursive backtracking style algorithm A for given problem.
- Identify structure of subproblems generated by A on an instance
 I of size n
 - Estimate number of different subproblems generated as a function of *n*. Is it polynomial or exponential in *n*?
 - If the number of problems is "small" (polynomial) then they typically have some "clean" structure.
- Sewrite subproblems in a compact fashion.
- Rewrite recursive algorithm in terms of notation for subproblems.
- Onvert to iterative algorithm by bottom up evaluation in an appropriate order.
- Optimize further with data structures and/or additional ideas.

A variation

Input A string $w \in \Sigma^*$ and access to a language $L \subseteq \Sigma^*$ via function IsStringinL(string x) that decides whether x is in L, and non-negative integer k Goal Decide if $w \in L^k$ using IsStringinL(string x) as a black box sub-routine

Example

Suppose *L* is *English* and we have a procedure to check whether a string/word is in the *English* dictionary.

- Is the string "isthisanenglishsentence" in *English*⁵?
- Is the string "isthisanenglishsentence" in *English*⁴?
- Is "asinineat" in *English*²?
- Is "asinineat" in *English*⁴?
- Is "zibzzzad" in English1?

Recursive Solution

When is $w \in L^k$?

Recursive Solution

When is $w \in L^k$? k = 0: $w \in L^k$ iff $w = \epsilon$ k = 1: $w \in L^k$ iff $w \in L$ k > 1: $w \in L^k$ if w = uv with $u \in L$ and $v \in L^{k-1}$

Recursive Solution

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When is w \in L^k?
k = 0: w \in L^k iff w = \epsilon
k = 1: w \in L^k iff w \in L
k > 1: w \in L^k if w = uv with u \in L and v \in L^{k-1}
Assume w is stored in array A[1...n]
IsStringinLk(A[1..n], k):
     If (\mathbf{k} = \mathbf{0})
          If (n = 0) Output YES
          Else Ouput NO
     If (\mathbf{k} = \mathbf{1})
          Output IsStringinL(A[1..n])
     Else
          For (i = 1 \text{ to } n - 1) do
                (i = 1 \text{ to } n - 1) \text{ do}
If (IsStringinL(A[1..i]) \text{ and } IsStringinLk(A[1..i], k - 1))
                     Output YES
```

Output NO

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     Else
          For (i = 1 \text{ to } n - 1) do
               If (IsStringinL(A[1..i])) and IsStringinLk(A[i + 1..n], k - 1))
                     Output YES
     Output NO
```

 How many distinct sub-problems are generated by IsStringinLk(A[1..n], k)?

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- How many distinct sub-problems are generated by IsStringinLk(A[1...n], k)? O(nk)
- How much space?

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Output NO

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- How much space? O(nk) pause
- Running time?

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Output NO

- How many distinct sub-problems are generated by IsStringinLk(A[1..n], k)? O(nk)
- How much space? O(nk) pause
- Running time? O(n²k)

Another variant

Question: What if we want to check if $w \in L^i$ for some $0 \le i \le k$? That is, is $w \in \bigcup_{i=0}^k L^i$?



Definition

A string is a palindrome if $w = w^R$. Examples: *I*, *RACECAR*, *MALAYALAM*, *DOOFFOOD*

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Problem: Given a string w find the *longest subsequence* of w that is a palindrome.

Example

MAHDYNAMICPROGRAMZLETMESHOWYOUTHEM has MHYMRORMYHM as a palindromic subsequence

Assume w is stored in an array A[1..n]

LPS(A[1..n]): length of longest palindromic subsequence of A.

Recursive expression/code?

Part I

Edit Distance and Sequence Alignment

Spell Checking Problem

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Question: Given two strings $x_1x_2...x_n$ and $y_1y_2...y_m$ what is a *distance* between them?

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Edit Distance: minimum number of "edits" to transform x into y.

Edit Distance

Definition

Edit distance between two words X and Y is the number of letter insertions, letter deletions and letter substitutions required to obtain Y from X.

Example

The edit distance between FOOD and MONEY is at most 4:

 $\underline{F}OOD \rightarrow MO\underline{O}D \rightarrow MON\underline{O}D \rightarrow MON\underline{E}\underline{D} \rightarrow MONEY$

Edit Distance: Alternate View

Alignment

Place words one on top of the other, with gaps in the first word indicating insertions, and gaps in the second word indicating deletions.

F O O D M O N E Y

FOOD MONEY

Edit Distance: Alternate View

Alignment

Place words one on top of the other, with gaps in the first word indicating insertions, and gaps in the second word indicating deletions.

$\begin{array}{cccc} F & O & O & & D \\ M & O & N & E & Y \end{array}$

Formally, an alignment is a set M of pairs (i, j) such that each index appears at most once, and there is no "crossing": i < i' and i is matched to j implies i' is matched to j' > j. In the above example, this is $M = \{(1, 1), (2, 2), (3, 3), (4, 5)\}$.

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Place words one on top of the other, with gaps in the first word indicating insertions, and gaps in the second word indicating deletions.

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Edit Distance Problem

Problem

Given two words, find the edit distance between them, i.e., an alignment of smallest cost.

Applications

- Spell-checkers and Dictionaries
- Onix diff
- S DNA sequence alignment ... but, we need a new metric

Similarity Metric

Definition

For two strings X and Y, the cost of alignment M is

- **(**Gap penalty] For each gap in the alignment, we incur a cost δ .
- (a) [Mismatch cost] For each pair p and q that have been matched in M, we incur cost α_{pq} ; typically $\alpha_{pp} = 0$.

Similarity Metric

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Edit distance is special case when $\delta = \alpha_{pq} = 1$.

An Example

Example

Alternative:

Or a really stupid solution (delete string, insert other string):

 $Cost = 19\delta$.

What is the edit distance between...

What is the minimum edit distance for the following two strings, if insertion/deletion/change of a single character cost 1 unit?



(A) 1
(B) 2
(C) 3
(D) 4
(E) 5

What is the edit distance between...

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Sequence Alignment

- Input Given two words **X** and **Y**, and gap penalty δ and mismatch costs α_{pq}
 - Goal Find alignment of minimum cost

Let $X = \alpha x$ and $Y = \beta y$

 α, eta : strings.

x and y single characters.

Think about optimal edit distance between X and Y as alignment, and consider last column of alignment of the two strings:

α	x	or	$\boldsymbol{\alpha}$	x	or	αx	
β	y		βy			β	y

Observation

Prefixes must have optimal alignment!

Problem Structure

Observation

Let $X = x_1 x_2 \cdots x_m$ and $Y = y_1 y_2 \cdots y_n$. If (m, n) are not matched then either the mth position of X remains unmatched or the nth position of Y remains unmatched.

• Case x_m and y_n are matched.

- Pay mismatch cost \$\alpha_{x_m y_n}\$ plus cost of aligning strings \$x_1 \cdots x_{m-1}\$ and \$y_1 \cdots y_{n-1}\$
- **2** Case x_m is unmatched.

0 Pay gap penalty plus cost of aligning $x_1 \cdots x_{m-1}$ and $y_1 \cdots y_n$

- Solution Case *y_n* is unmatched.
 - Pay gap penalty plus cost of aligning $x_1 \cdots x_m$ and $y_1 \cdots y_{n-1}$

Subproblems and Recurrence

Optimal Costs

Let Opt(i, j) be optimal cost of aligning $x_1 \cdots x_i$ and $y_1 \cdots y_j$. Then

$$Opt(i,j) = \min \begin{cases} \alpha_{x_iy_j} + Opt(i-1,j-1), \\ \delta + Opt(i-1,j), \\ \delta + Opt(i,j-1) \end{cases}$$

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Base Cases: $Opt(i, 0) = \delta \cdot i$ and $Opt(0, j) = \delta \cdot j$

Recursive Algorithm

Assume X is stored in array A[1..m] and Y is stored in B[1..n]Array *COST* stores cost of matching two chars. Thus *COST*[*a*, *b*] give the cost of matching character *a* to character *b*.

```
 \begin{array}{l} \textit{EDIST}(\textit{A}[1..m],\textit{B}[1..n]) \\ \text{If} (m = 0) \text{ return } n\delta \\ \text{If} (n = 0) \text{ return } m\delta \\ m_1 = \delta + \textit{EDIST}(\textit{A}[1..(m - 1)],\textit{B}[1..n]) \\ m_2 = \delta + \textit{EDIST}(\textit{A}[1..m],\textit{B}[1..(n - 1)])) \\ m_3 = \textit{COST}[\textit{A}[m],\textit{B}[n]] + \textit{EDIST}(\textit{A}[1..(m - 1)],\textit{B}[1..(n - 1)]) \\ \text{ return } \min(m_1,m_2,m_3) \end{array}
```

Example

DEED and DREAD

DEED PREAD

<u>DEED</u> LEAD DREAD

Memoizing the Recursive Algorithm

```
int M[0..m][0..n]
Initialize all entries of M[i][j] to \infty
return EDIST(A[1..m], B[1..n])
```

```
EDIST(A[1..m], B[1..n])
    If (M[i][j] < \infty) return M[i][j] (* return stored value *)
    If (\mathbf{m} = \mathbf{0})
         M[i][i] = n\delta
    ElseIf (n = 0)
         M[i][j] = m\delta
    Else
         m_1 = \delta + EDIST(A[1..(m-1)], B[1..n])
         m_2 = \delta + EDIST(A[1..m], B[1..(n-1)]))
         m_3 = COST[A[m], B[n]] + EDIST(A[1..(m-1)], B[1..(n-1)])
         M[i][j] = \min(m_1, m_2, m_3)
    return M[i][j]
```

Removing Recursion to obtain Iterative Algorithm

```
\begin{split} \textit{EDIST}(A[1..m], B[1..n]) & \text{int} \quad M[0..m][0..n] \\ \textit{for } i = 1 \text{ to } m \text{ do } M[i, 0] = i\delta \\ \textit{for } j = 1 \text{ to } n \text{ do } M[0, j] = j\delta \end{split} \\ \textit{for } i = 1 \text{ to } m \text{ do } \\ \textit{for } j = 1 \text{ to } n \text{ do } \\ \textit{for } j = 1 \text{ to } n \text{ do } \\ M[i][j] = \min \begin{cases} \alpha_{x_i y_j} + M[i-1][j-1], \\ \delta + M[i-1][j], \\ \delta + M[i][j-1] \end{cases} \end{split}
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```

Analysis

Removing Recursion to obtain Iterative Algorithm

```
EDIST (A[1..m], B[1..n])
int M[0..m][0..n]
for i = 1 to m do M[i, 0] = i\delta
for j = 1 to n do M[0, j] = j\delta
for i = 1 to m do
for j = 1 to n do
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```

Analysis

```
Running time is O(mn).
```

Space used is O(mn).

Matrix and DAG of Computation

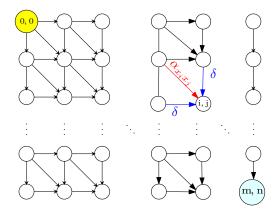


Figure: Iterative algorithm in previous slide computes values in row order.

Example

DEED and DREAD

Sequence Alignment in Practice

- Typically the DNA sequences that are aligned are about 10⁵ letters long!
- So about 10¹⁰ operations and 10¹⁰ bytes needed
- The killer is the 10GB storage
- On we reduce space requirements?

Optimizing Space

Recall

$$M(i,j) = \min \begin{cases} \alpha_{x_i y_j} + M(i-1,j-1), \\ \delta + M(i-1,j), \\ \delta + M(i,j-1) \end{cases}$$

- 2 Entries in *j*th column only depend on (j 1)st column and earlier entries in *j*th column
- Only store the current column and the previous column reusing space; N(i, 0) stores M(i, j 1) and N(i, 1) stores M(i, j)

Computing in column order to save space

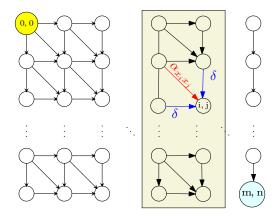


Figure: M(i, j) only depends on previous column values. Keep only two columns and compute in column order.

Space Efficient Algorithm

for all *i* do
$$N[i, 0] = i\delta$$

for $j = 1$ to *n* do
 $N[0, 1] = j\delta$ (* corresponds to $M(0, j)$ *)
for $i = 1$ to *m* do
 $N[i, 1] = \min \begin{cases} \alpha_{x_i y_j} + N[i - 1, 0] \\ \delta + N[i - 1, 1] \\ \delta + N[i, 0] \end{cases}$
for $i = 1$ to *m* do
Copy $N[i, 0] = N[i, 1]$

Analysis

Running time is O(mn) and space used is O(2m) = O(m)

Analyzing Space Efficiency

- From the m × n matrix M we can construct the actual alignment (exercise)
- Matrix N computes cost of optimal alignment but no way to construct the actual alignment
- Space efficient computation of alignment? More complicated algorithm see notes and Kleinberg-Tardos book.

Part II

Longest Common Subsequence Problem

LCS Problem

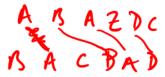
Definition

LCS between two strings X and Y is the length of longest common subsequence between X and Y.

Example

LCS between ABAZDC and BACBAD is

A B D ABAD



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Derive a dynamic programming algorithm for the problem.

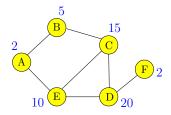
Part III

Maximum Weighted Independent Set in Trees

Maximum Weight Independent Set Problem

Input Graph G = (V, E) and weights $w(v) \ge 0$ for each $v \in V$

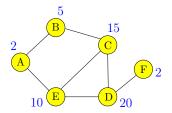
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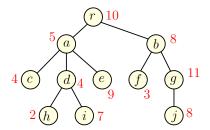


Maximum weight independent set in above graph: $\{B, D\}$

Maximum Weight Independent Set in a Tree

Input Tree T = (V, E) and weights $w(v) \ge 0$ for each $v \in V$

Goal Find maximum weight independent set in T



Maximum weight independent set in above tree: ??

For an arbitrary graph G:

- Number vertices as v_1, v_2, \ldots, v_n
- Solutions Find recursively optimum solutions without v_n (recurse on $G v_n$) and with v_n (recurse on $G v_n N(v_n)$ & include v_n).
- Saw that if graph *G* is arbitrary there was no good ordering that resulted in a small number of subproblems.

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What about a tree?

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What about a tree? Natural candidate for v_n is root r of T?

Natural candidate for v_n is root r of T? Let \mathcal{O} be an optimum solution to the whole problem.

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- Case $r \in \mathcal{O}$: None of the children of r can be in \mathcal{O} . $\mathcal{O} \{r\}$ contains an optimum solution for each subtree of Thanging at a grandchild of r.

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How many of them?

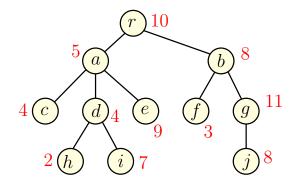
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Subproblems? Subtrees of T rooted at nodes in T.

How many of them? O(n)

Example



A Recursive Solution

T(u): subtree of T hanging at node uOPT(u): max weighted independent set value in T(u)

OPT(u) =

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$$OPT(u) = \max \begin{cases} \sum_{v \text{ child of } u} OPT(v), \\ w(u) + \sum_{v \text{ grandchild of } u} OPT(v) \end{cases}$$

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- Compute OPT(u) bottom up. To evaluate OPT(u) need to have computed values of all children and grandchildren of u
- What is an ordering of nodes of a tree *T* to achieve above? Post-order traversal of a tree.

 $\begin{aligned} \mathsf{MIS-Tree}(T): \\ & \text{Let } \mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_n \text{ be a post-order traversal of nodes of T} \\ & \text{for } i = 1 \text{ to } n \text{ do} \\ & M[\mathbf{v}_i] = \max \begin{pmatrix} \sum_{v_i \text{ child of } v_i} M[v_i], \\ & w(v_i) + \sum_{v_j \text{ grandchild of } v_i} M[v_j] \end{pmatrix} \\ & \text{return } M[\mathbf{v}_n] \text{ (* Note: } \mathbf{v}_n \text{ is the root of } T \text{ *)} \end{aligned}$

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Space:

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Space: O(n) to store the value at each node of TRunning time:

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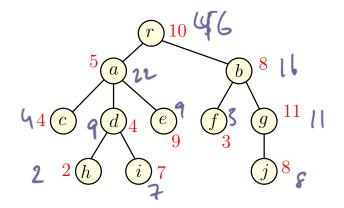
Naive bound: O(n²) since each M[v_i] evaluation may take O(n) time and there are n evaluations.

 $\begin{aligned} \mathsf{MIS-Tree}(T): \\ & \text{Let } \mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_n \text{ be a post-order traversal of nodes of T} \\ & \text{for } i = 1 \text{ to } n \text{ do} \\ & M[\mathbf{v}_i] = \max \begin{pmatrix} \sum_{v_i \text{ child of } v_i} M[v_i], \\ & w(v_i) + \sum_{v_j \text{ grandchild of } v_i} M[v_j] \end{pmatrix} \\ & \text{return } M[\mathbf{v}_n] \text{ (* Note: } \mathbf{v}_n \text{ is the root of } T \text{ *)} \end{aligned}$

Space: O(n) to store the value at each node of TRunning time:

- Naive bound: O(n²) since each M[v_i] evaluation may take O(n) time and there are n evaluations.
- Better bound: O(n). A value M[v_j] is accessed only by its parent and grand parent.

Example



Takeaway Points

- Oynamic programming is based on finding a recursive way to solve the problem. Need a recursion that generates a small number of subproblems.
- Given a recursive algorithm there is a natural DAG associated with the subproblems that are generated for given instance; this is the dependency graph. An iterative algorithm simply evaluates the subproblems in some topological sort of this DAG.
- The space required to evaluate the answer can be reduced in some cases by a careful examination of that dependency DAG of the subproblems and keeping only a subset of the DAG at any time.