

More Dynamic Programming

Lecture 15

March 12, 2015

What is the running time of the following?

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

$$f(x, y) = \sum_{i=1}^{x+y-1} x * f(x + y - i, i - 1),$$
$$f(0, y) = y \quad f(x, 0) = x.$$

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

- 1 Develop a recursive backtracking style algorithm \mathcal{A} for given problem.
- 2 Identify *structure* of subproblems generated by \mathcal{A} on an instance \mathbf{l} of size \mathbf{n}
 - 1 Estimate number of different subproblems generated as a function of \mathbf{n} . Is it polynomial or exponential in \mathbf{n} ?
 - 2 If the number of problems is “small” (polynomial) then they typically have some “clean” structure.
- 3 Rewrite subproblems in a compact fashion.
- 4 Rewrite recursive algorithm in terms of notation for subproblems.
- 5 Convert to iterative algorithm by bottom up evaluation in an appropriate order.
- 6 Optimize further with data structures and/or additional ideas.

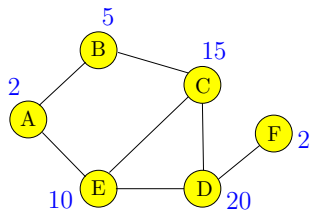
Part I

Maximum Weighted Independent Set in Trees

Maximum Weight Independent Set Problem

Input Graph $\mathbf{G} = (\mathbf{V}, \mathbf{E})$ and weights $\mathbf{w}(\mathbf{v}) \geq \mathbf{0}$ for each $\mathbf{v} \in \mathbf{V}$

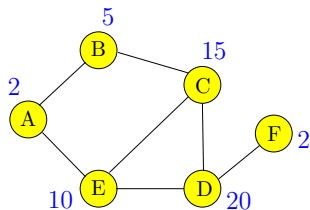
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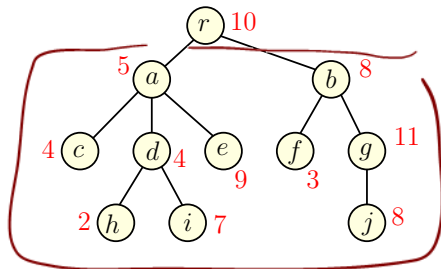


Maximum weight independent set in above graph: $\{\mathbf{B}, \mathbf{D}\}$

Maximum Weight Independent Set in a Tree

Input Tree $\mathbf{T} = (\mathbf{V}, \mathbf{E})$ and weights $\mathbf{w}(\mathbf{v}) \geq \mathbf{0}$ for each $\mathbf{v} \in \mathbf{V}$

Goal Find maximum weight independent set in \mathbf{T}



Maximum weight independent set in above tree: ??

Independent set in a tree...

In a tree with n nodes, there is always an independent set of size (bigger is better [this is America!])

(A) $\Omega(1)$

(B) $\Omega(\log n)$

(C) $\Omega(\sqrt{n})$

(D) $n/2$

(E) $n - 5$

Towards a Recursive Solution

For an arbitrary graph \mathbf{G} :

- 1 Number vertices as $\mathbf{v}_1, \mathbf{v}_2, \dots, \mathbf{v}_n$
- 2 Find recursively optimum solutions without \mathbf{v}_n (recurse on $\mathbf{G} - \mathbf{v}_n$) and with \mathbf{v}_n (recurse on $\mathbf{G} - \mathbf{v}_n - \mathbf{N}(\mathbf{v}_n)$ & include \mathbf{v}_n).
- 3 Saw that if graph \mathbf{G} is arbitrary there was no good ordering that resulted in a small number of subproblems.

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What about a tree? Natural candidate for \mathbf{v}_n is root \mathbf{r} of \mathbf{T} ?

Towards a Recursive Solution

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

Case $r \notin \mathcal{O}$: Then \mathcal{O} contains an optimum solution for each subtree of T hanging at a child of r .

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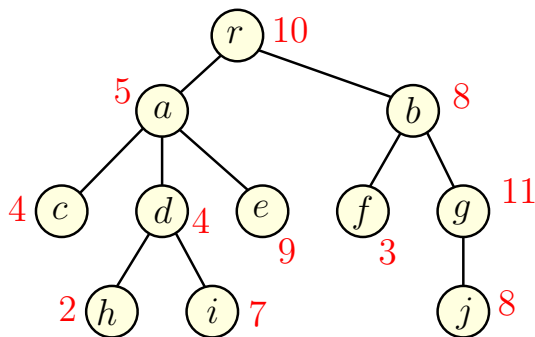
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Subproblems? Subtrees of T hanging at nodes in T .

How many of them? $O(n)$

Example



A Recursive Solution

$T(u)$: subtree of T hanging at node u

$OPT(u)$: max weighted independent set value in $T(u)$

$$OPT(u) =$$

A Recursive Solution

$\mathbf{T(u)}$: subtree of \mathbf{T} hanging at node \mathbf{u}

$\mathbf{OPT(u)}$: max weighted independent set value in $\mathbf{T(u)}$

$$\mathbf{OPT(u)} = \max \left\{ \begin{array}{l} \sum_{\mathbf{v} \text{ child of } \mathbf{u}} \mathbf{OPT(v)}, \\ \mathbf{w(u)} + \sum_{\mathbf{v} \text{ grandchild of } \mathbf{u}} \mathbf{OPT(v)} \end{array} \right.$$

Iterative Algorithm

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

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- 2 What is an ordering of nodes of a tree **T** to achieve above?
Post-order traversal of a tree.

Iterative Algorithm

MIS-Tree(**T**):

Let v_1, v_2, \dots, v_n be a post-order traversal of nodes of **T**
for $i = 1$ **to** n **do**

$$M[v_i] = \max \left(\begin{array}{l} \sum_{v_j \text{ child of } v_i} M[v_j], \\ w(v_i) + \sum_{v_j \text{ grandchild of } v_i} M[v_j] \end{array} \right)$$

return $M[v_n]$ (* Note: v_n is the root of **T** *)

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Space:

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Space: $O(n)$ to store the value at each node of **T**

Running time:

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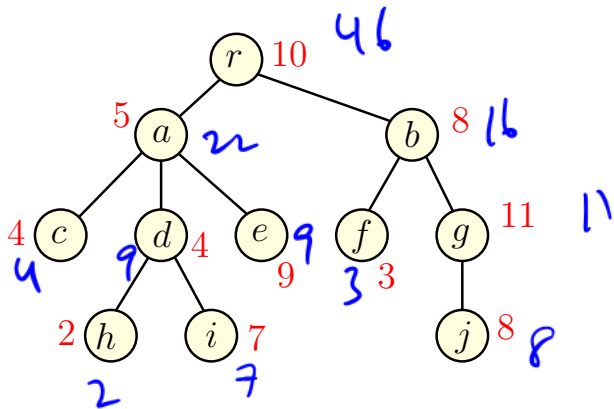
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Space: $O(n)$ to store the value at each node of **T**

Running time:

- 1 Naive bound: $O(n^2)$ since each $M[v_i]$ evaluation may take $O(n)$ time and there are n evaluations.
- 2 Better bound: $O(n)$. A value $M[v_j]$ is accessed only by its parent and grand parent.

Example



Part II

Edit Distance and Sequence Alignment

Spell Checking Problem

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What does nearness mean?

Question: Given two strings $x_1x_2 \dots x_n$ and $y_1y_2 \dots y_m$ 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**:

FOOD → MOOD → MONOD → MONED → 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

Edit Distance: Alternate View

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Formally, an **alignment** is a set \mathbf{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 $\mathbf{M} = \{(1, 1), (2, 2), (3, 3), (4, 5)\}$.

Edit Distance: Alternate View

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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

- 1 Spell-checkers and Dictionaries
- 2 Unix `diff`
- 3 DNA sequence alignment ... but, we need a new metric

Similarity Metric

Definition

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

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

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

An Example

Example

o | c | c | u | r | r | e | n | c | e
o | c | c | u | r | r | a | n | c | e

$$\text{Cost} = \delta + \alpha_{ae}$$

Alternative:

o | c | c | u | r | r | e | n | c | e
o | c | c | u | r | r | a | n | c | e

$$\text{Cost} = 3\delta$$

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

o | c | c | u | r | r | e | n | c | e
o | c | c | u | r | r | e | n | c | e

$$\text{Cost} = 19\delta.$$

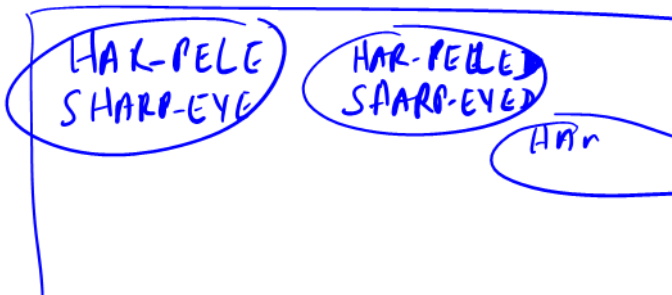
What is the edit distance between...

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

HAR-PELED

SHARP_EYED

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



Sequence Alignment

Input Given two words X and Y , and gap penalty δ and mismatch costs α_{pq}

Goal Find alignment of minimum cost

Edit distance

Basic observation

Let $\mathbf{X} = \alpha x$ and $\mathbf{Y} = \beta y$

α, β : strings.

x and y single characters.

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

α	x
β	y

or

α	x
βy	

or

αx	
β	y

Observation

Prefixes must have optimal alignment!

Problem Structure

Observation

Let $\mathbf{X} = x_1x_2 \cdots x_m$ and $\mathbf{Y} = y_1y_2 \cdots y_n$. If (m, n) are not matched then either the m th position of \mathbf{X} remains unmatched or the n th position of \mathbf{Y} remains unmatched.

- ① **Case** x_m and y_n are matched.
 - ① Pay mismatch cost $\alpha_{x_my_n}$ plus cost of aligning strings $x_1 \cdots x_{m-1}$ and $y_1 \cdots y_{n-1}$
- ② **Case** x_m is unmatched.
 - ① Pay gap penalty plus cost of aligning $x_1 \cdots x_{m-1}$ and $y_1 \cdots y_n$
- ③ **Case** y_n is unmatched.
 - ① 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 $\text{Opt}(i, j)$ be optimal cost of aligning $x_1 \cdots x_i$ and $y_1 \cdots y_j$. Then

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

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

Dynamic Programming Solution

```
for all i do M[i, 0] = iδ
```

```
for all j do M[0, j] = jδ
```

```
for i = 1 to m do
```

```
  for j = 1 to n do
```

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

Analysis

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Analysis

- 1 Running time is $O(mn)$.

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Analysis

- 1 Running time is $O(mn)$.
- 2 Space used is $O(mn)$.

Matrix and DAG of Computation

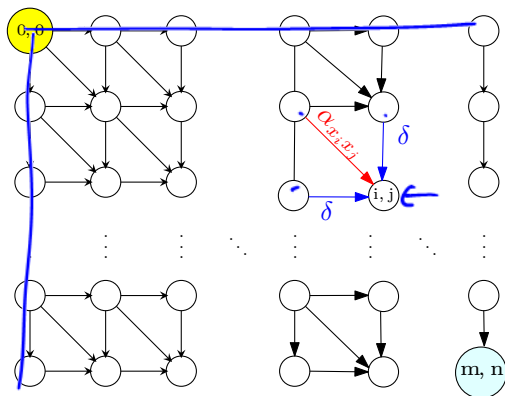


Figure : Iterative algorithm in previous slide computes values in row order. Optimal value is a shortest path from $(0,0)$ to (m,n) in .

Sequence Alignment in Practice

- 1 Typically the DNA sequences that are aligned are about 10^5 letters long!
- 2 So about 10^{10} operations and 10^{10} bytes needed
- 3 The killer is the 10GB storage
- 4 Can we reduce space requirements?

Optimizing Space

1 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
- 3 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

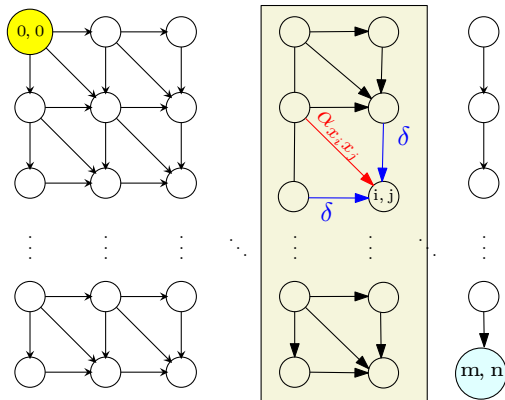


Figure : $M(i, j)$ only depends on previous column values. Keep only two columns and compute in column order.

Space Efficient Algorithm

```
for all i do N[i, 0] = iδ
for j = 1 to n do
  N[0, j] = jδ (* corresponds to M(0, j) *)
  for i = 1 to m do
    N[i, j] = min {
      αxi,yj + N[i - 1, 0]
      δ + N[i - 1, j]
      δ + N[i, 0]
    }
  for i = 1 to m do
    Copy N[i, 0] = N[i, j]
```

Analysis

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

Analyzing Space Efficiency

- 1 From the $m \times n$ matrix \mathbf{M} we can construct the actual alignment (exercise)
- 2 Matrix \mathbf{N} computes cost of optimal alignment but no way to construct the actual alignment
- 3 Space efficient computation of alignment? More complicated algorithm — see notes and Kleinberg-Tardos book.

Part III

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



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LCS between ABAZDC and BACBAD is 4 via ABAD

ABAZDC
BACBAD

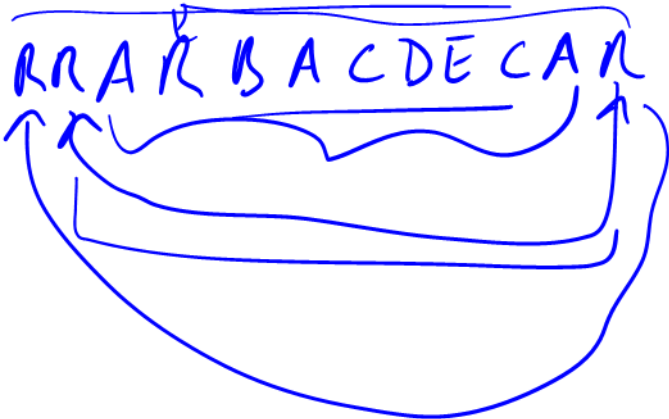
$OPT(i, j) = \text{LCS for } x_1 \dots x_i \text{ and } y_1 \dots y_j$

$$\text{OPT}(i, j) = \max \begin{cases} \text{OPT}(i, j-1) \\ \text{OPT}(i-1, j) \\ \text{OPT}(i-1, j-1) \\ 1 + \text{OPT}(i-1, j-1) \end{cases} \text{ if } x_i = y_j$$

$$\text{OPT}(i, 0) = 0$$

$$\text{OPT}(0, j) = 0$$

RACECAR



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Example

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Derive a dynamic programming algorithm for the problem.