

More Dynamic Programming

Lecture 9

February 18, 2014

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.

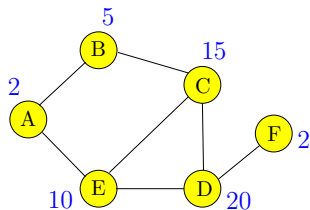
Part I

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 $\mathbf{v} \in \mathbf{V}$

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

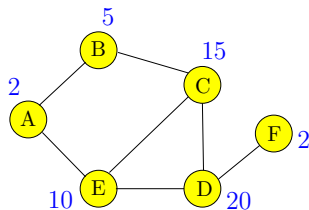


Maximum weight independent set in above graph: $\{\mathbf{B}, \mathbf{D}\}$

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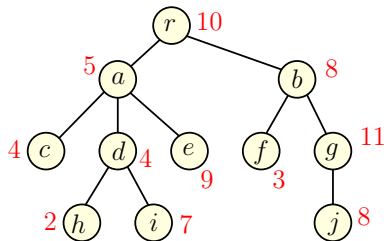


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Maximum Weight Independent Set in a Tree

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.

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

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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 .

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 T hanging at a grandchild of r .

Subproblems? Subtrees of T hanging at nodes in T .

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A Recursive Solution

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

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

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

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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?
Post-order traversal of a tree.

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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 *)

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.

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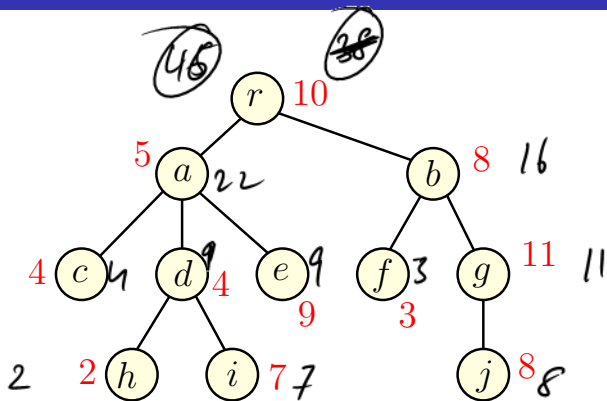
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Example



Part II

Edit Distance and Sequence Alignment

Spell Checking Problem

Given a string “exponen” that is not in the dictionary, how should a spell checker suggest a *nearby* string?

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?

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

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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 |

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 $M = \{(1, 1), (2, 2), (3, 3), (4, 5)\}$. Cost of an alignment is the number of mismatched columns plus number of unmatched indices in both strings.

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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 $\alpha_{pp} = 0$.

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

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An Example

Example

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

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

Alternative:

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

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

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

| | | | | | | | | | | | | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|--|--|
| o | c | u | r | r | a | 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 | or | α | x | or | αx | |
| β | y | | βy | | | 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

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

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

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

Analysis

- 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

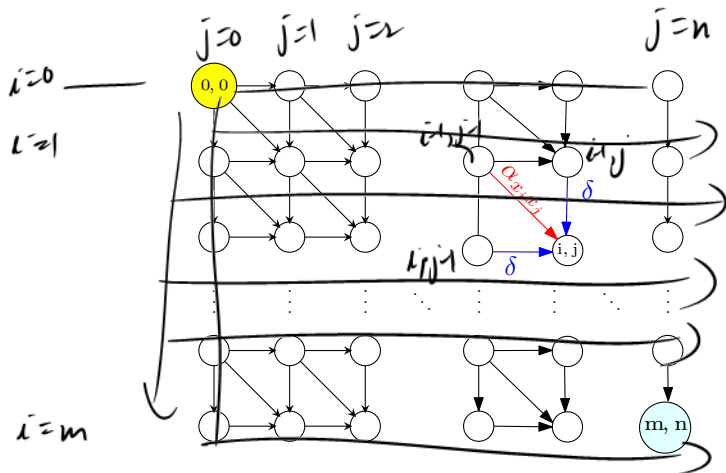


Figure : Iterative algorithm in previous slide computes values in row order. Optimal value is a shortest path from $(0,0)$ to (m,n) in .

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

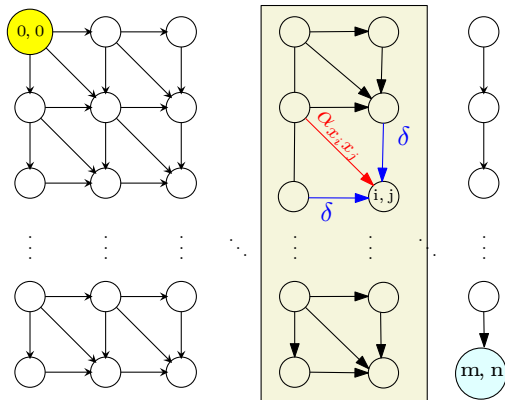


Figure : $M(i, j)$ only depends on previous column values. Keep only two columns and compute in column order.

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 {
      αxiyj + 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 text book.

Part III

DAGs and Dynamic Programming

Recursion and DAGs

Observation

Let A be a recursive algorithm for problem Π . For each instance I of Π there is an associated DAG $G(I)$.

- 1 Create directed graph $G(I)$ as follows...
- 2 For each sub-problem in the execution of A on I create a node.
- 3 If sub-problem v depends on or *recursively calls* sub-problem u add *directed edge* (u, v) to graph.
- 4 $G(I)$ is a DAG. Why? If $G(I)$ has a cycle then A will not terminate on I .

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Iterative Algorithm for...

Dynamic Programming and DAGs

Observation

*An iterative algorithm **B** obtained from a recursive algorithm **A** for a problem Π does the following:*

*For each instance **I** of Π , it computes a topological sort of **G(I)** and evaluates sub-problems according to the topological ordering.*

- 1 Sometimes the DAG **G(I)** can be obtained directly without thinking about the recursive algorithm **A**
- 2 In some cases (**not all**) the computation of an optimal solution reduces to a shortest/longest path in DAG **G(I)**
- 3 Topological sort based shortest/longest path computation is dynamic programming!

A quick reminder...

A Recursive Algorithm for weighted interval scheduling

Let O_i be value of an optimal schedule for the first i jobs.

```
Schedule( $n$ ):  
  if  $n = 0$  then return 0  
  if  $n = 1$  then return  $w(v_1)$   
   $O_{p(n)} \leftarrow$  Schedule( $p(n)$ )  
   $O_{n-1} \leftarrow$  Schedule( $n - 1$ )  
  if ( $O_{p(n)} + w(v_n) < O_{n-1}$ ) then  
     $O_n = O_{n-1}$   
  else  
     $O_n = O_{p(n)} + w(v_n)$   
  return  $O_n$ 
```

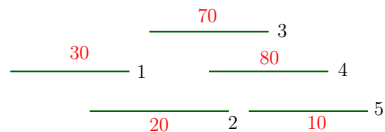
Weighted Interval Scheduling via...

Longest Path in a DAG

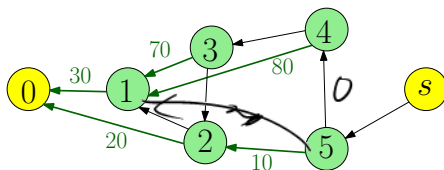
Given intervals, create a **DAG** as follows:

- 1 Create one node for each interval, plus a dummy sink node **0** for interval **0**, plus a dummy source node **s**.
- 2 For each interval **i** add edge **(i, p(i))** of the length/weight of **v_i**.
- 3 Add an edge from **s** to **n** of length **0**.
- 4 For each interval **i** add edge **(i, i - 1)** of length **0**.

Example



$$p(5) = 2, p(4) = 1, p(3) = 1, p(2) = 0, p(1) = 0$$



Relating Optimum Solution

Given interval problem instance I let $G(I)$ denote the DAG constructed as described.

Claim

Optimum solution to weighted interval scheduling instance I is given by longest path from s to t in $G(I)$.

Assuming claim is true,

- 1 If I has n intervals, DAG $G(I)$ has $n + 2$ nodes and $O(n)$ edges. Creating $G(I)$ takes $O(n \log n)$ time: to find $p(i)$ for each i . How?
- 2 Longest path can be computed in $O(n)$ time — recall $O(m + n)$ algorithm for shortest/longest paths in DAGs.

Relating Optimum Solution

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Claim

Optimum solution to weighted interval scheduling instance I is given by longest path from s to 0 in $G(I)$.

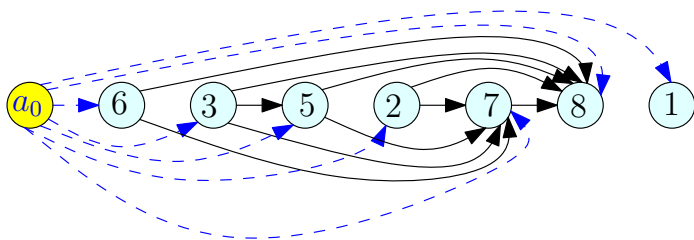
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DAG for Longest Increasing Sequence

Given sequence a_1, a_2, \dots, a_n create DAG as follows:

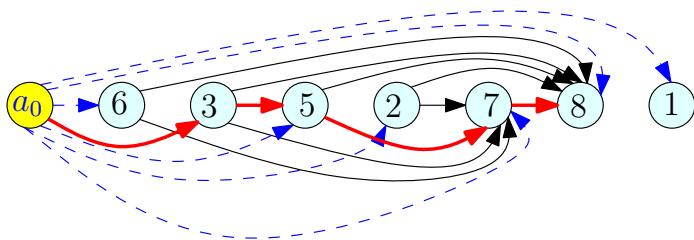
- 1 add sentinel a_0 to sequence where a_0 is less than smallest element in sequence
- 2 for each i there is a node v_i
- 3 if $i < j$ and $a_i < a_j$ add an edge (v_i, v_j)
- 4 find longest path from v_0



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Takeaway Points

- 1 Dynamic programming is based on finding a recursive way to solve the problem. Need a recursion that generates a small number of subproblems.
- 2 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**.
- 3 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.

Notes

