

# More Dynamic Programming

Lecture 10

February 19, 2015

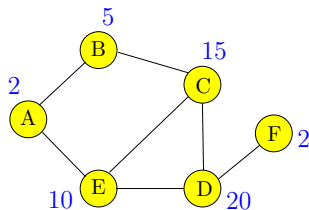
# Part I

## Maximum Weighted Independent Set in Trees

# Maximum Weight Independent Set Problem

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

Goal Find maximum weight independent set in  $G$

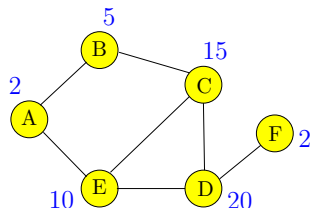


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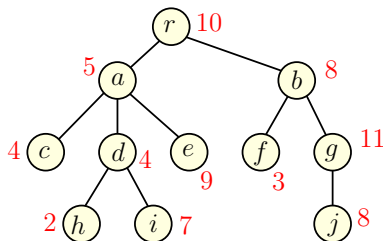


Maximum weight independent set in above graph:  $\{B, D\}$

# Maximum Weight Independent Set in a Tree

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

Goal Find maximum weight independent set in  $T$



Maximum weight independent set in above tree: ??

# Towards a Recursive Solution

- 1 For an arbitrary graph  $G$ :
  - 1 Number vertices as  $v_1, v_2, \dots, v_n$
  - 2 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  $v_n$ ).
  - 3 Saw that if graph  $G$  is arbitrary there was no good ordering that resulted in a small number of subproblems.
- 2 What about a tree?
- 3 Natural candidate for  $v_n$  is root  $r$  of  $T$ ?

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- 2 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  $r$ .
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# A Recursive Solution

- 1  $T(u)$ : subtree of  $T$  hanging at node  $u$ .
- 2  $OPT(u)$ : max weighted independent set value in  $T(u)$ .
- 3  $OPT(u) = \max \begin{cases} \sum_{v \text{ child of } u} OPT(v), \\ w(u) + \sum_{v \text{ grandchild of } u} OPT(v) \end{cases}$



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# Iterative Algorithm

- 1 Compute  $OPT(u)$  bottom up. To evaluate  $OPT(u)$  need to have computed values of all children and grandchildren of  $u$
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- 3 Post-order traversal of a tree.

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### 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  $T$  \*)

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

## 3 Running time:

- 1 Naive bound:  $O(n^2)$  since each  $M[v_i]$  evaluation may take  $O(n)$  time and there are  $n$  evaluations.
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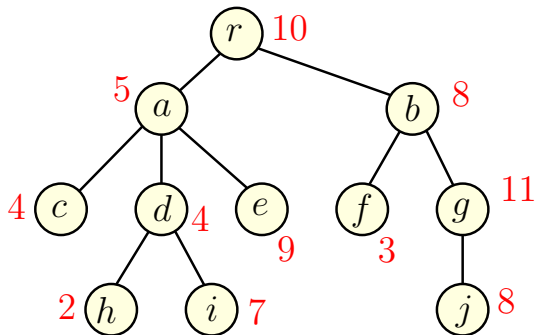
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# Example

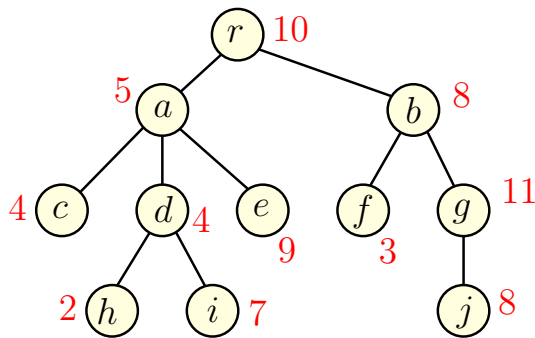




# Dominating set

## Definition

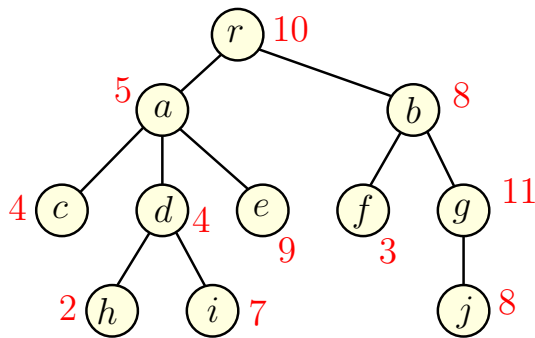
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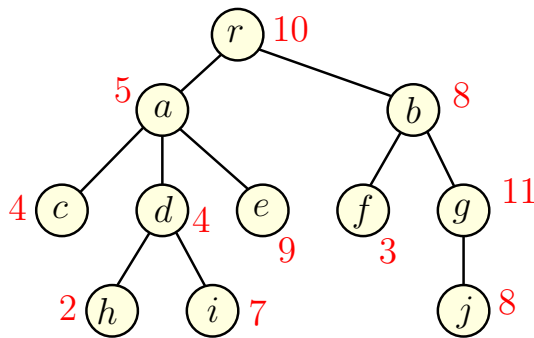
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**Dominating Set** is **NP-Hard!**

## Part II

# DAGs and Dynamic Programming

# Recursion and DAGs

## Observation

Let  $A$  be a recursive algorithm for problem  $\Pi$ . For each instance  $I$  of  $\Pi$  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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*For each instance  $I$  of  $\Pi$ , 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  $G(I)$
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# Iterative Algorithm for...

## Dynamic Programming and DAGs

### Observation

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# 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  $v_i$ .
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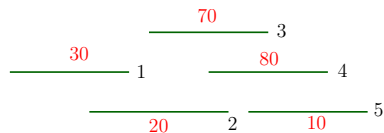
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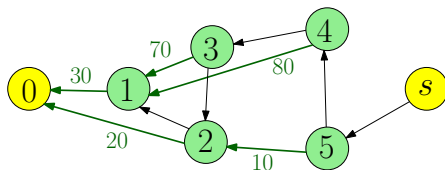
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# Example



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



# Relating Optimum Solution

- 1 Given interval problem instance  $I$  let  $G(I)$  denote the DAG constructed as described.
- 2 We have...

## Claim

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

- 3 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.

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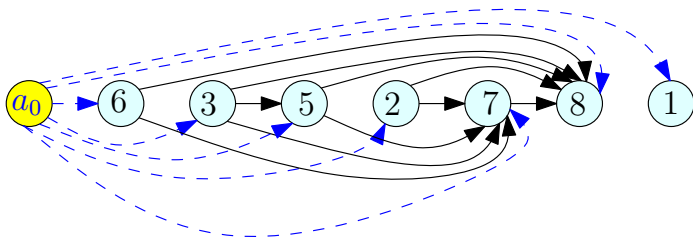
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Given sequence  $a_1, a_2, \dots, a_n$  create DAG as follows:

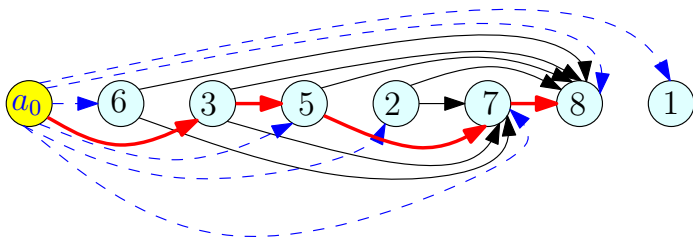
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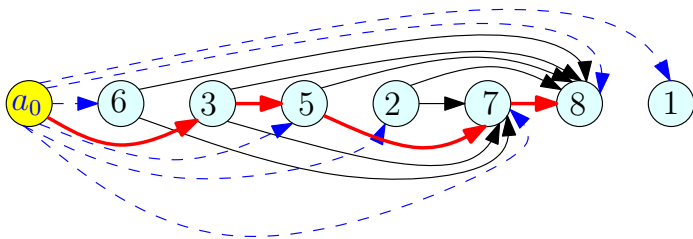
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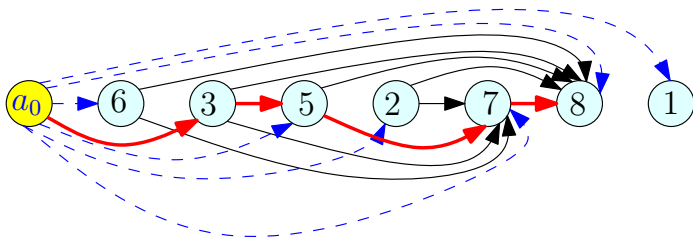
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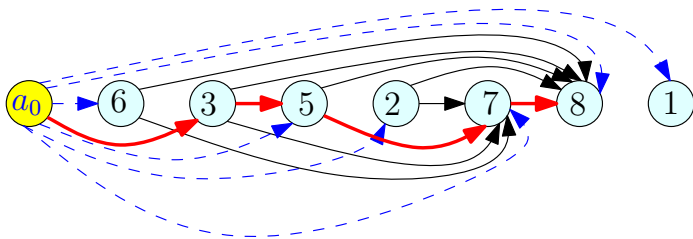
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## Part III

# Edit Distance and Sequence Alignment

# Spell Checking Problem

- 1 Given a string “exponen” that is not in the dictionary, how should a spell checker suggest a *nearby* string?
- 2 What does nearness mean?
- 3 **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** 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:

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# 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  $\delta$ .
- 2 [Mismatch cost] For each pair  $p$  and  $q$  that have been matched in  $M$ , we incur cost  $\alpha_{pq}$ ; typically  $\alpha_{pp} = 0$ .

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# Sequence Alignment

**Input** Given two words  $X$  and  $Y$ , and gap penalty  $\delta$  and mismatch costs  $\alpha_{pq}$

**Goal** Find alignment of minimum cost

# Edit distance

## Basic observation

- 1 Let  $X = \alpha x$  and  $Y = \beta y$ .
- 2  $\alpha, \beta$ : strings.  $x$  and  $y$  single characters.
- 3 Optimal edit distance between  $X$  and  $Y$  as alignment. Consider last column of alignment of the two strings:

$\alpha$	$x$	or	$\alpha$	$x$	or	$\alpha x$	
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Let  $X = x_1x_2 \cdots x_m$  and  $Y = y_1y_2 \cdots y_n$ . If  $(m, n)$  are not matched then either the  $m$ th position of  $X$  remains unmatched or the  $n$ th position of  $Y$  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  $y_1 \cdots y_n$
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# 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

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for all  $i$  do  $M[i, 0] = i\delta$ 
```

```
for all  $j$  do  $M[0, j] = j\delta$ 
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for  $i = 1$  to  $m$  do
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- 1 Running time is  $O(mn)$ .

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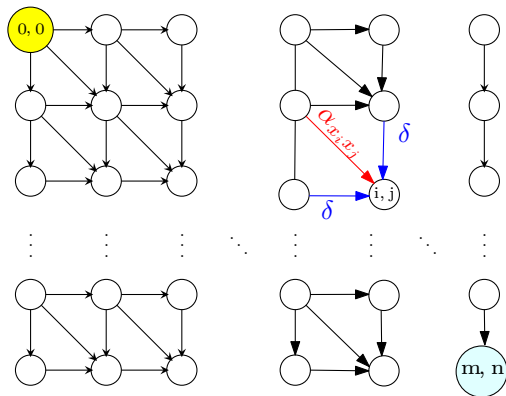
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## Analysis

- 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  $M(i, j)$

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# Computing in column order to save space

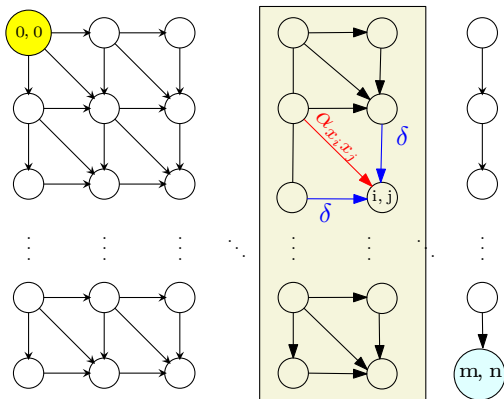


Figure:  $M(i, j)$  only depends on previous column values. Keep only two columns and compute in column order.

# Space Efficient Algorithm

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

  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  $M$  we can construct the actual alignment (exercise)
- 2 Matrix  $N$  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.



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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.









