

More DP: Edit Distance and Independent Sets in Trees

Lecture 14

October 16, 2018

How many subproblems?

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

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

How many distinct subproblems when computing $f(n, n)$?

- (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.

What is the running time for each subproblem?

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 worst-case time to evaluate the output of a subproblem given values for its recursive subproblems when computing $f(n, n)$ is:

- (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.

What is the total running time?

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 I of size n
 - 1 Estimate number of different subproblems generated as a function of n . Is it polynomial or exponential in 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.

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 *English*¹?

Recursive Solution

When is $w \in L^k$?

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$k = 0$: $w \in L^k$ 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 $v \in L^{k-1}$

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$k > 1$: $w \in L^k$ 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 ($k = 0$)

 If ($n = 0$) Output YES

 Else Output NO

If ($k = 1$)

 Output **IsStringinL**($A[1..n]$)

Else

 For ($i = 1$ to $n - 1$) do

 If (**IsStringinL**($A[1..i]$) and **IsStringinLk**($A[i + 1..n]$, $k - 1$))

 Output YES

Output NO

Analysis

IsStringinLk($A[1..n]$, k):

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Output NO

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

Analysis

IsStringinLk($A[1..n]$, k):

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 If (**IsStringinL**($A[1..i]$) and **IsStringinL**($A[i + 1..n]$, $k - 1$))

 Output YES

Output NO

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

Analysis

IsStringinLk(A[1..n], k):

If ($k = 0$)

 If ($n = 0$) Output YES

 Else Output NO

If ($k = 1$)

 Output **IsStringinL(A[1..n])**

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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?

Analysis

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 Output YES

Output NO

- How many distinct sub-problems are generated by **IsStringinL**($A[1..n]$, k)? $O(nk)$
- How much space? $O(nk)$ pause
- Running time? $O(n^2k)$

Naming subproblems and recursive equation

ISLk(i, h): a boolean which is **1** if $A[i..n]$ is in L^h , **0** otherwise

Base case: **ISLk**($n + 1, 0$) = **1** interpreting $A[n + 1..n]$ as ϵ

Naming subproblems and recursive equation

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Recursive relation:

- **ISLk**(i, h) = **1** if $\exists i < j \leq n + 1$ such that
(**ISLk**($j, h - 1$) = **1** and **IsStringinL**($A[i..(j - 1)]$) = **1**)
- **ISLk**(i, h) = **0** otherwise

Alternately:

ISLk(i, h) = $\max_{i < j \leq n + 1}$ (**ISLk**($j, h - 1$) **IsStringinL**($A[i..(j - 1)]$))

Output: **ISLk**($1, k$)

Another variant

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

Exercise

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

Exercise

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

$LPS(i, j)$: length of longest palindromic subsequence of $A[i..j]$.

Recursive expression/code?

Part I

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

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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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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 $M = \{(1, 1), (2, 2), (3, 3), (4, 5)\}$.

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 $\delta = \alpha_{pq} = 1$.

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?

374

473

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

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

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- (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 $X = \alpha x$ and $Y = \beta y$

α, β : 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
β	y

or

α	x
βy	

or

αx	
β	y

Observation

Prefixes must have optimal alignment!

Problem Structure

Observation

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

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 .

$EDIST(A[1..m], B[1..n])$

If ($m = 0$) return $n\delta$

If ($n = 0$) return $m\delta$

$m_1 = \delta + EDIST(A[1..(m - 1)], B[1..n])$

$m_2 = \delta + EDIST(A[1..m], B[1..(n - 1)])$

$m_3 = COST[A[m], B[n]] + EDIST(A[1..(m - 1)], B[1..(n - 1)])$

return $\min(m_1, m_2, m_3)$

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

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

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 $\text{Opt}(0, j) = \delta \cdot j$

Example

DEED and 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 ( $m = 0$ )
```

```
     $M[i][j] = n\delta$ 
```

```
  ElseIf ( $n = 0$ )
```

```
     $M[i][j] = m\delta$ 
```

```
  Else
```

```
     $m_1 = \delta + \text{EDIST}(A[1..(m-1)], B[1..n])$ 
```

```
     $m_2 = \delta + \text{EDIST}(A[1..m], B[1..(n-1)])$ 
```

```
     $m_3 = \text{COST}[A[m], B[n]] + \text{EDIST}(A[1..(m-1)], B[1..(n-1)])$ 
```

```
     $M[i][j] = \min(m_1, m_2, m_3)$ 
```

```
  return  $M[i][j]$ 
```

Removing Recursion to obtain Iterative Algorithm

EDIST(*A*[1..*m*], *B*[1..*n*])

int *M*[0..*m*][0..*n*]

for *i* = 1 to *m* do *M*[*i*, 0] = *i*δ

for *j* = 1 to *n* do *M*[0, *j*] = *j*δ

for *i* = 1 to *m* do

for *j* = 1 to *n* do

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

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] = iδ  
  for j = 1 to n 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

Removing Recursion to obtain Iterative Algorithm

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

Analysis

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

Matrix and DAG of Computation

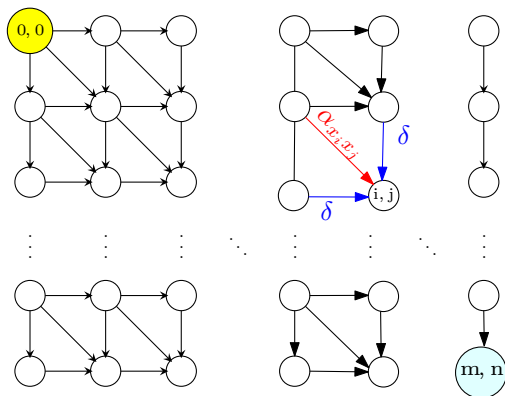


Figure: Iterative algorithm in previous slide computes values in row order.

Example

DEED and DREAD

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

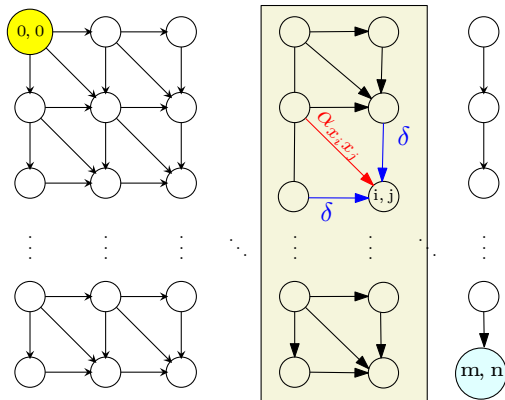


Figure: $M(i, j)$ only depends on previous column values. Keep only two columns and compute in column order.

Space Efficient Algorithm

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

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Derive a dynamic programming algorithm for the problem.

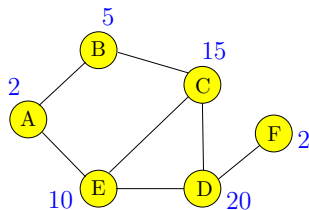
Part III

Maximum Weighted Independent Set in Trees

Maximum Weight Independent Set Problem

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

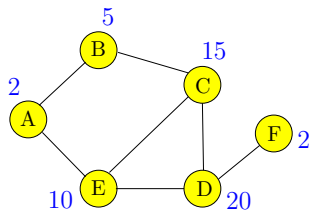
Goal Find maximum weight independent set in G



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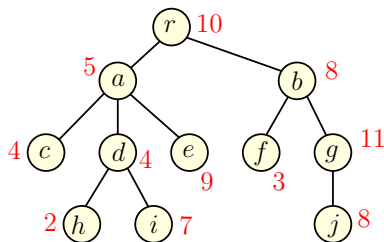


Maximum weight independent set in above graph: $\{B, D\}$

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

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.

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What about a tree?

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- 3 Saw that if graph G is arbitrary there was no good ordering that resulted in a small number of subproblems.

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

Towards a Recursive Solution

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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 T hanging at a grandchild of r .

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Subproblems? Subtrees of T rooted at nodes in T .

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Subproblems? Subtrees of T rooted at nodes in T .

How many of them?

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.

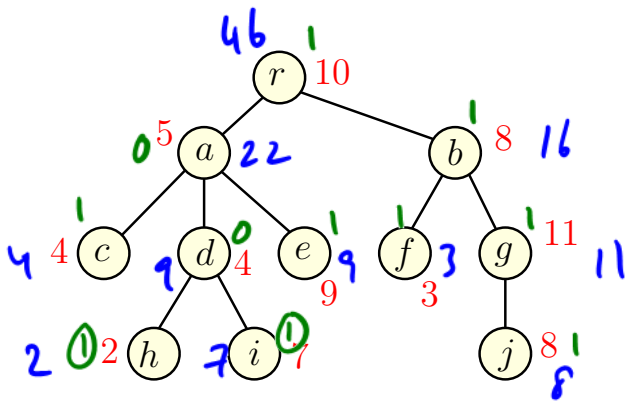
Case $r \notin \mathcal{O}$: Then \mathcal{O} contains an optimum solution for each subtree of T hanging at a child of r .

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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 u

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

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

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

Running time:

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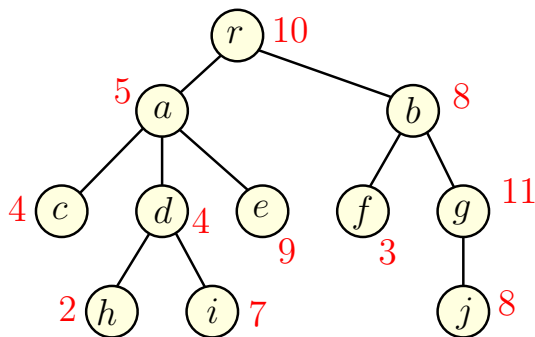
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Running time:

- 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



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.