

Design by Induction

(Based on [Manber 1989])

Yih-Kuen Tsay

Department of Information Management National Taiwan University

Introduction



- It is not necessary to design the steps required to solve a problem from scratch.
- It is sufficient to guarantee the following:
 - 1. It is possible to solve one small instance or a few small instances of the problem. (base case)
 - 2. A solution to every problem/instance can be constructed from solutions to smaller problems/instances. (inductive step)

Evaluating Polynomials



Problem

Given a sequence of real numbers a_n , a_{n-1} , \cdots , a_1 , a_0 , and a real number x, compute the value of the polynomial

$$P_n(x) = a_n x^n + a_{n-1} x^{n-1} + \cdots + a_1 x + a_0.$$

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Motivation: different approaches to the inductive step may result in algorithms of very different time complexities.



- Let $P_{n-1}(x) = a_{n-1}x^{n-1} + \cdots + a_1x + a_0$.
- Induction hypothesis (first attempt)
 We know how to evaluate a polynomial represented by the input a_{n-1}, \dots, a_1, a_0 , at the point x, i.e., we know how to compute $P_{n-1}(x)$.
- $P_n(x) = a_n x^n + P_{n-1}(x).$



- Induction hypothesis (second attempt)
 We know how to compute $P_{n-1}(x)$, and we know how to compute x^{n-1} .
- $P_n(x) = a_n x(x^{n-1}) + P_{n-1}(x).$



- Let $P'_{n-1}(x) = a_n x^{n-1} + a_{n-1} x^{n-2} + \cdots + a_1$.
- Induction hypothesis (final attempt)
 We know how to evaluate a polynomial represented by the coefficients a_n , a_{n-1} , \cdots , a_1 , at the point x, i.e., we know how to compute $P'_{n-1}(x)$.
- $P_n(x) = P'_n(x) = P'_{n-1}(x) \cdot x + a_0.$



More generally,

$$\begin{cases} P'_0(x) = a_n \\ P'_i(x) = P'_{i-1}(x) \cdot x + a_{n-i}, \text{ for } 1 \le i \le n \end{cases}$$



Algorithm Polynomial_Evaluation (\bar{a}, x) ; begin $P := a_n$; for i := 1 to n do $P := x * P + a_{n-i}$ end

This algorithm is known as Horner's rule.

Maximal Induced Subgraph



Problem

Given an undirected graph G = (V, E) and an integer k, find an induced subgraph H = (U, F) of G of maximum size such that all vertices of H have degree $\geq k$ (in H), or conclude that no such induced subgraph exists.

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Design Idea: in the inductive step, we try to remove one vertex (that cannot possibly be part of the solution) to get a smaller instance.

One-to-One Mapping



Problem

Given a finite set A and a mapping f from A to itself, find a subset $S \subseteq A$ with the maximum number of elements, such that (1) the function f maps every element of S to another element of S (i.e., f maps S into itself), and (2) no two elements of S are mapped to the same element (i.e., f is one-to-one when restricted to S).

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Design Idea: similar to the previous problem; in the inductive step, we try to remove one element (that cannot possibly be part of the solution) to get a smaller instance.

One-to-One Mapping (cont.)



```
Algorithm Mapping (f, n);
begin
   S:=A:
   for j := 1 to n do c[j] := 0;
   for i := 1 to n do increment c[f[i]];
    for i := 1 to n do
       if c[j] = 0 then put j in Queue;
    while Queue not empty do
        remove i from the top of Queue;
       S := S - \{i\};
       decrement c[f[i]];
       if c[f[i]] = 0 then put f[i] in Queue
end
```

Celebrity



Problem

Given an $n \times n$ adjacency matrix, determine whether there exists an i (the "celebrity") such that all the entries in the i-th column (except for the ii-th entry) are 1, and all the entries in the i-th row (except for the ii-th entry) are 0.

Note: A celebrity corresponds to a sink of the directed graph.

Note: Every directed graph has at most one sink.

Celebrity



Problem

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Note: Every directed graph has at most one sink.

Motivation: the trivial solution has a time complexity of $O(n^2)$. Can we do better, in O(n)?

Celebrity (cont.)



```
Algorithm Celebrity (Know);
begin
   i := 1:
   i := 2;
    next := 3:
    while next \le n+1 do
       if Know[i, j] then i := next
                    else i := next;
        next := next + 1:
   if i = n + 1 then candidate := j
                else candidate := i:
```

Celebrity (cont.)



```
wrong := false;
    k := 1:
    Know[candidate, candidate] := false;
    while not wrong and k < n do
       if Know[candidate, k] then wrong := true;
       if not Know[k, candidate] then
          if candidate \neq k then wrong := true;
        k := k + 1:
    if not wrong then celebrity := candidate
                 else celebrity := 0;
end
```

The Skyline Problem



Problem

Given the exact locations and shapes of several rectangular buildings in a city, draw the skyline (in two dimension) of these buildings, eliminating hidden lines.

Motivation: different approaches to the inductive step may result in algorithms of very different time complexities.

The Skyline Problem



Problem

Given the exact locations and shapes of several rectangular buildings in a city, draw the skyline (in two dimension) of these buildings, eliminating hidden lines.

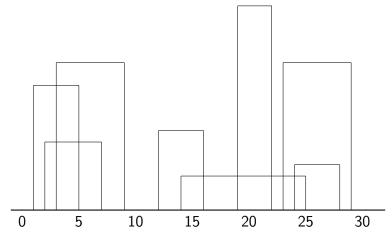
Motivation: different approaches to the inductive step may result in algorithms of very different time complexities.

Compare: adding buildings one by one to an existing skyline vs. merging two skylines of about the same size

Representation of a Skyline



(1,**11**,5), (2,**6**,7), (3,**13**,9), (12,**7**,16), (14,**3**,25), (19,**18**,22), (23,**13**,29), and (24,**4**,28).

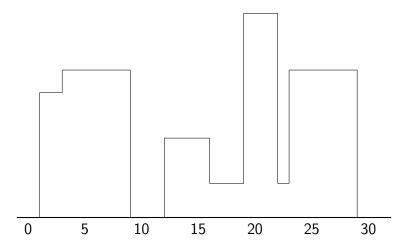


Algorithms 2012

Representation of a Skyline (cont.)



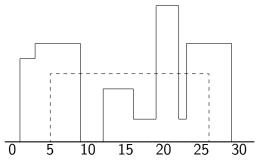
(1,**11**,3,**13**,9,**0**,12,**7**,16,**3**,19,**18**,22,**3**,23,**13**,29).



Adding a Building



• Add (5,**9**,26) to (1,**11**,3,**13**,9,**0**,12,**7**,16,**3**,19,**18**,22,**3**,23,**13**,29).



The skyline becomes (1,**11**,3,**13**,9,**9**,19,**18**,22,**9**,23,**13**,29).

Merging Two Skylines



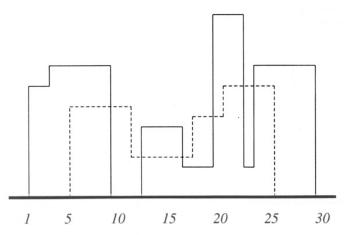


Figure 5.7 Merging two skylines.

Source: [Manber 1989].

Balance Factors in Binary Trees



Problem

Given a binary tree T with n nodes, compute the balance factors of all nodes.

The balance factor of a node is defined as the difference between the height of the node's left subtree and the height of the node's right subtree.

Balance Factors in Binary Trees



Problem

Given a binary tree T with n nodes, compute the balance factors of all nodes.

The balance factor of a node is defined as the difference between the height of the node's left subtree and the height of the node's right subtree.

Motivation: an example of why we must strengthen the hypothesis (and hence the problem to be solved).

Balance Factors in Binary Trees (cont.)



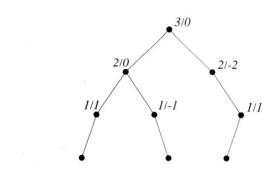


Figure 5.8 A binary tree. The numbers represent h/b, where h is the height and b is the balance factor.

Source: [Manber 1989].

Balance Factors in Binary Trees (cont.)



Induction hypothesis

We know how to compute balance factors of all nodes in trees that have < n nodes.

Balance Factors in Binary Trees (cont.)



- Induction hypothesis
 We know how to compute balance factors of all nodes in trees that have < n nodes.</p>
- Stronger induction hypothesis
 We know how to compute balance factors and heights of all nodes in trees that have < n nodes.</p>

Maximum Consecutive Subsequence



Problem

Given a sequence x_1, x_2, \dots, x_n of real numbers (not necessarily positive) find a subsequence x_i, x_{i+1}, \dots, x_j (of consecutive elements) such that the sum of the numbers in it is maximum over all subsequences of consecutive elements.

Example:

In the sequence (2, -3, 1.5, -1, 3, -2, -3, 3), the maximum subsequence is (1.5, -1, 3).

Maximum Consecutive Subsequence



Problem

Given a sequence x_1, x_2, \dots, x_n of real numbers (not necessarily positive) find a subsequence x_i, x_{i+1}, \dots, x_j (of consecutive elements) such that the sum of the numbers in it is maximum over all subsequences of consecutive elements.

Example:

In the sequence (2, -3, 1.5, -1, 3, -2, -3, 3), the maximum subsequence is (1.5, -1, 3).

Motivation: another example of strengthening the hypothesis.

Maximum Consecutive Subsequence (cont.)



Induction hypothesis

We know how to find the maximum subsequence in sequences of size < n.

Maximum Consecutive Subsequence (cont.)



- Induction hypothesis
 We know how to find the maximum subsequence in sequences of size < n.
 </p>
- Stronger induction hypothesis
 We know how to find, in sequences of size < n, the maximum subsequence overall and the maximum subsequence that is a suffix.
 </p>

Maximum Consecutive Subsequence (cont.)



```
Algorithm Max_Consec_Subseq (X, n);
begin
    Global\_Max := 0:
    Suffix_Max := 0:
    for i := 1 to n do
       if x[i] + Suffix\_Max > Global\_Max then
          Suffix\_Max := Suffix\_Max + x[i];
          Global Max = Suffix Max
       else if x[i] + Suffix\_Max > 0 then
               Suffix\_Max := Suffix\_Max + x[i]
       else Suffix Max := 0
end
```

The Knapsack Problem



Problem

Given an integer K and n items of different sizes such that the i-th item has an integer size k_i , find a subset of the items whose sizes sum to exactly K, or determine that no such subset exists.

The Knapsack Problem



Problem

Given an integer K and n items of different sizes such that the i-th item has an integer size k_i , find a subset of the items whose sizes sum to exactly K, or determine that no such subset exists.

Design Idea: use strong induction so that solutions to all smaller instances may be used.



- Let P(n, K) denote the problem where n is the number of items and K is the size of the knapsack.
- Induction hypothesis
 We know how to solve P(n-1, K).



- Let P(n, K) denote the problem where n is the number of items and K is the size of the knapsack.
- Induction hypothesis
 We know how to solve P(n-1, K).
- Stronger induction hypothesis
 We know how to solve P(n-1, k), for all $0 \le k \le K$.



An example of the table constructed for the knapsack problem:

	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
$k_1 = 2$	0	-	ı	-	-	-	-	-	-	-	-	-	-	-	-	-	-
$k_2 = 3$	0	-	0	ı	-	-	-	-	-	-	-	-	-	-	-	-	-
$k_3 = 5$	0	-	0	0	-	0	-	1	1	-	I	-	-	-	-	-	-
$k_4 = 6$	0	-	0	0	-	0		0	0	1	0	I	-	1	1	-	I

"I": a solution containing this item has been found.

"O": a solution without this item has been found.

"-": no solution has yet been found.



```
Algorithm Knapsack (S, K);
    P[0,0].exist := true;
    for k := 1 to K do
        P[0, k].exist := false;
    for i := 1 to n do
        for k := 0 to K do
            P[i, k].exist := false;
            if P[i-1,k].exist then
              P[i, k].exist := true;
              P[i, k].belong := false
            else if k - S[i] > 0 then
                   if P[i-1, k-S[i]].exist then
                      P[i, k].exist := true;
                      P[i, k]. belong := true
```