Week 2

3mg 17, 18

(5)

Divide and Conquer

Counting Inversions



Design and Analysis of Algorithms I

Tim Roughgarder

(3) conquer via secursive calls.
(3) contine sol of subproblem, the most warmed problem regeneity.

Odivel into Amake subprob.

wither and a makeum non man

The Problem

Input: array A containing the numbers 1,2,3,..,n in some arbitrary order

Output : number of inversions = number of pairs (i,j) Pet of invession

ed. 16 A is sorted # Inversion =0

of array indices with i<j and A[i] > A[j]

To find the # of inversions, we will the musulus in sorted order once and the second time we write them as they given. Went, we connect nowbers. The # of wassing lives = # unusions

Examples and Motivation

Example (1,3,5,2,4,6)

<u>Inversions</u>: (3,2), (5,4)

Motivation: numerical similarity measure

between two ranked lists eg: for collaborative filtering

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see section 3-2.3 for the olef of (to suggest stend forome to lung).

111/11-2

The worst case is that the array is in the brokeward archer

N(4-1) W (My) (Mar) (n - cn - n) = 1 AW (J-W) N. L. What is the largest-possible number of inversions that a 6element array can have?

0 21

036

0 64

7000 loops.

High-Level Approach

Brute-force: $\theta(n^2)$ time = Asymptotic winning time.

Can we do better? Yes!

Divide + Conquer KEY IDEA # 1: classity the vivessions of an array A of laugth in into one of three lynes. Call an inversion (i,j) [with i<j]

<u>Left</u>: if i,j ≤ n/2 ←

<u>Right</u> : if i,j > n/2 [≪]

<u>Split</u>: if i<=n/2 < j[≮]

combine styr of algo

Note: can compute these recursively need separate subroutine for

these

High-Level Algorithm

Count (array A, length n)

if n=1, return 0 (600)

X = Count (1st half of A, n/2)

 $Y = Count (2^{nd} half of A, n/2_$

Z = CountSplitInv(A,n) <-- CURRENTLY UNIMPLEMENTED

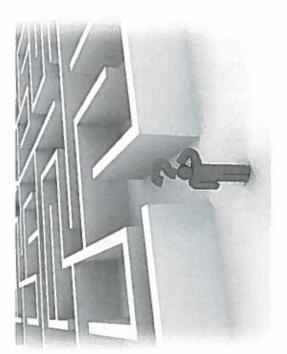
return x+y+z

Goal: implement CountSplitInv in linear (O(n)) time then count will run in O(nlog(n)) time [just like Merge Sort]

July 17, 18

Divide and Conquer

Counting Inversions II



Design and Analysis of Algorithms I

Piggybacking on Merge Sort

KEY IDEA # 2 : have recursive calls both count

inversions and sort.

[i.e., piggy back on Merge Sort]

Motivation: Merge subroutine naturally

uncovers split inversions [as we'll see]

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High-Level Algorithm (revised)

Sort-and-Count (array A, length n)

if n=1, return 0

else

(D,Z) = CountSplitInv(A,n) CURRENTED + Norge UNIMPLEMENTED (C,Y) = Sort-and-Count(2nd half of A, n/2) (B,X) = Sort-and-Count(1st half of A, n/2)return X+Y+Z Sorted version — Sorted version Sorted version of 1st half

=> then Count will run in O(nlog(n)) time [just like Merge Sort] Goal: implement CountSplitInv in linear (O(n)) time



Pseudocode for Merge:

D = output [length = n]
B = 1st sorted array [n/2]
C = 2nd sorted array [n/2]
i = 1



for k = 1 to n if B(i) < C(j) k D(k) = B(i) k i++ else [C(j) < B(i)] k D(k) = C(j)

end

(ignores end cases)

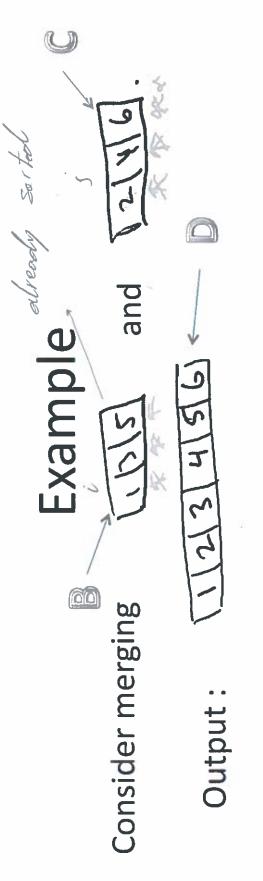
Suppose the input array A has no split inversions. What is the relationship between the sorted subarrays B and C?

 $_{
m O}$ B has the smallest element of A, C the second-smallest, B, the thirdsmallest, and so on.

O All elements of B are less than all elements of C. \checkmark

O All elements of B are greater than all elements of C.

O There is not enough information to answer this question.



⇒When 2 copied to output, discover the split ⇒ when 4 copied to output, discover (5,4) After copying 2 i is greater than I inversions (3,2) and (5,2)

General Claim

Claim the split inversions involving an element y of the 2ndd array C are precisely the numbers left in the 1st array B when y is copied to the output D.

Proof: Let x be an element of the 1st array B.

 $\sqrt{1}$. If x copied to output D before y, then x < y

⇒ no inversions involving x and y

2. If y copied to output D before x, then y < x

=> 🕱 and y are a (split) inversion.

Q.E.D

Merge_and_CountSplitInv

-- while merging the two sorted subarrays, keep running total of number of split inversions



copied to output D, increment total by number of elements -- when element of 2nd array C gets

remaining in 1st array B merge

running total

Run time of subroutine : $O(\vec{n}) + O(\vec{n}) = O(n)$

=> Sort_and_Count runs in O(nlog(n)) time [just like Merge Sort]

If you wild own MX then the total cost is Tim Roughgarden but if you odd it contstant times then it is our

mutin-1

Strassen's algo-, not milial , purclamental prob.

200 81/8 (15

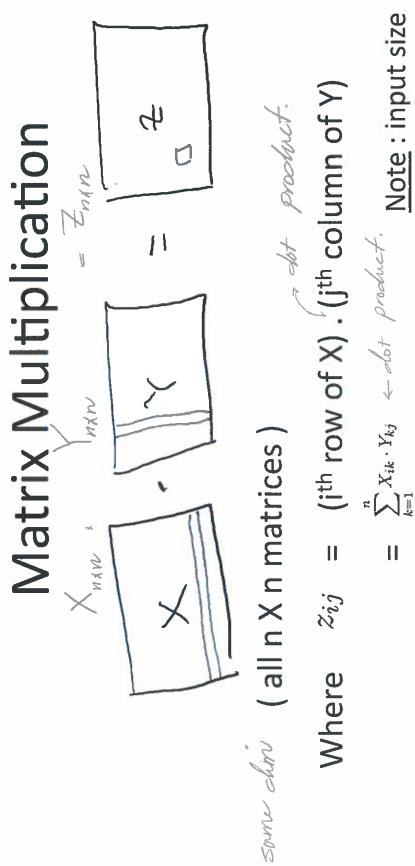


Matrix

Design and Analysis of Algorithms I

Divide and Conquer

Multiplication



 $= \theta(n^2) \xrightarrow{not} th$

The best we can hape, but there are if entires in the methin Tim Roughgarden · runding time: (no tention) *n = O (no) milt som. 15 Och?

Tim Roughgarden

Example (n=2)

$$z_{ij} = \sum_{k=1}^{n} X_{ik} \cdot Y_{kj}$$

$$\theta(n)$$

What is the asymptotic running time of the straightforward iterative algorithm for matrix multiplication?

- $\bigcirc \theta(n \log n)$
- $\bigcirc \theta(n^2)$
- $> O \theta(n^3)$
- $O \theta(n^4)$

We assume that the time to access tack while

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The whole idea of divide and congress.

The Divide and Conquer Paradigm

- 1. DIVIDE into smaller subproblems
- 2. CONQUER subproblems recursively.
- 3. COMBINE solutions of subproblems into one for the original problem.

Applying Divide and Conquer to method

Write X = (C) and (- (C) H

[where A through H are all n/2 by n/2 matrices]

multiplication 12 12 matrices we express the original I problem in town of the (CE + DG | CF + DH Then: (you check) RE + TSG AF + BH

L. KITTING

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Recursive Algorithm #1

AE+BG AF+BH GE+DG CF+DH

AC, BG, MF, BM, CE, DC, CF, DH

Step 1: recursively compute the 8 hecessary products. of 2 2 2 methos

 $\overline{\text{Step 2}}: \text{do the necessary additions } (\theta(n^2) \ time) \ \checkmark$

 $\overline{\operatorname{Fact}}:\operatorname{runtime}$ is $\theta(n^3)$ [follows from the master method]

Strassen's Algorithm (1969)

products 7 vs 8 secondly not very significant but since Step 1: recursively compute only 7 (cleverly chosen)

Step 2: do the necessary (clever) additions + subtractions (still $\theta(n^2)$ time)

Fact: better than cubic time!

[see Master Method lecture]

Tim Boughparden

O See the book

15 cest of surmotion and monthly theolion the same?

15 14 Zearly

(a 2) 11 X

The Details

The Seven Products: P(-A(F-H), 12- (A+1)H, 8-1-(LAD)E, 8-1-016-E), 8-1-(A+D) (E+H) (3+3) (7-4) - 4) - (4-1) (C-6) - 3

Claim: X. \ - (ce +00 ar +34) - (Ps +84-82+80) P,+8,

*ハロナガガーアンータイ F RE+BC Proof: AC+AC+OC+DC+DC+OC-OC-OC-ACH

Question: where did this come from? Open!)

COSt 6+ 1 and

×

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Design and Analysis of Algorithms I

Divide and Conquer

Closest Pair I

from comparticul goon solvaned material

Joan - 1

The Closest Pair Problem

Input : a set $P = \{p_1, ..., p_n\}$ of n points in the plane \mathbb{R}^2 .

Notation: $d(p_i, p_j) = \text{Euclidean distance}$

So if
$$p_i = (x_i, y_i)$$
 and $p_j = (x_j, y_j)$

$$d(p_i, p_j) = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2}$$

Output : a pair $p*, q* \in P$ of distinct points that minimize d(p,q) over p,q in the set P

Initial Observations

/ Assumption : (for convenience) all points have distinct

x-coordinates, distinct y-coordinates.

two leaps of one neno of paints)

no Hos.

Brute-force search : takes $\theta(n^2)$ time.

1-D Version of Closest Pair: For now, the easy problems.

1. Sort points (O(nlog(n)) time)

2. Return closest pair of adjacent points (O(n) time)

Gan we get the same runging time for the 2D version.

Goal: O(nlog(n)) time algorithm for 2-D version.

High-Level Approach

1. Make copies of points sorted by x-coordinate (P_x) and by y-[O(nlog(n)) time] coordinate (P_y)

closest in if

(but this is not enough!)

but not closest

close in &

2. Use Divide+Conquer

The Divide and Conquer Paradigm

1. DIVIDE into smaller subproblems.

2. CONQUER subproblems recursively.

The germity is here 3. COMBINE solutions of subproblems into one for the original problem.

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ExION in the slides and his tracking but the book is force

, 4

We sort in the a wol there ast in tooth.

ClosestPair(P_x , P_v)

BASE CASE

Lusion stop

Qx, Qy, Rx, Ry [takes O(n) time] sorted by a coordinate and by coordinate 1. Let Q = left half of P, R = right half of P. Form

Terminate 2. $(p_1,q_1)=ClosestPair(Q_x,Q_y)$ The bucky one is that the closed punity of 3 extra in Q or R

 $\rightarrow 3. (p_2, q_2) = ClosestPair(R_x, R_y)$

4. $(p_3,q_3) = ClosestSplitPair(P_x,P_y)$ while p_3

5. Return best of (p_1,q_1) , (p_2,q_2) , (p_3,q_3)

and then conjuse it with the closust of the left is right a But we go through it couly if the closest split pain is closes than the other two cases: Very idea wounds that we don't weed to find the closest split power

we don't need full blown syllet subroutine in O(n) time. What will be the overall running time of the Closest Pair algorithm? (Choose the smallest upper Suppose we can correctly implement the ClosestSplitPair bound that applies.)

 \bigcirc 0(n)

 $\sim 0(n\log n)$

 \bigcirc $O(n(\log n)^2)$

 \bigcirc $O(n^2)$

recursive call V Result of 1st its distance is less than $d(p_1,q_1)$ Result of 2nd and d(p₂,q₂).

recursive call

the closest split pair in "unlucky case" where

Key Idea: only need to bother computing

andy in the unducky case we go

through the April care.

I says if we although care about the split pain on not.

ClosestPair(P_x , P_v)

Let Q = left half of P, R = right half of P. Form

BASE CASE

 Q_x , Q_y , R_x , R_y [takes O(n) time]

- 2. $(p_1,q_1) = ClosestPair(Q_x,Q_v)$
- 3. $(p_2,q_2) = ClosestPair(R_x,R_y)$
- Let $\delta = min\{d(p_1, q_1), d(p_2, q_2)\}$
- 5. $(p_3,q_3) = ClosestSplitPair(P_x,P_v,\delta)$
- 6. Return best of (p_1,q_1) , (p_2,q_2) , (p_3,q_3) WILL DESCRIBE NEXT

pair (we go through The cast it we cloud go through spalit. This only It we closest pair of P is a split O(n) time Requirements whenever Correct

(bust stat poin) Tim Roughgarden harvo the

The cast is Engitty ran abricus.

If a splutjais is the closest it should be in

ClosestSplitPair(P_x , P_v , δ

Let $\bar{x}=$ biggest x-coordinate in left of P. (O(1) time)

5 is the

Let S_v = points of P with x-coordinate to be x-8 and x+6, (O(n) time) Sorted by y-coordinate

best pair = (p,q), best = d(p,q)Let p,q = i^{th} , $(i+j)^{th}$ points of S_v Initialize best $= \delta$, best pair = NULL For j = 1 to # min(7, 2-i) If d(p,q) < best For i = 1 to $|S_v| - ||w||$ time

O(n) Gine

At end return best pair

Correctness Claim

 $min\{d(p_1,q_1),d(p_2,q_2)\}$

Claim : Let $p \in Q, q \in R$ be a split pair with $d(p,q) < \delta$

Then: (A) p and q are members of S_v

(B) p and q are at most $\frac{7}{7}$ positions apart in S_v .



Corollary1: If the closest pair of P is a split pair, then the ClosestSplitPair finds it. claim is true! Corollary2 ClosestPair is correct, and runs in O(nlog(n)) time

Assuming

July 20, 2018 (17)

Divide and Conquer

Closest Pair II



Design and Analysis of Algorithms I

Correctness Claim

 $min\{d(p_1,q_1),d(p_2,q_2)\}$

Claim : Let $p \in Q, q \in R$ be a split pair with $d(p,q) < \delta$

Then: (A) p and q are members of S_v

(B) p and q are at most 7 positions apart in S_v .

We want to wove this

Tim Roughbarder

Fim Roughgarden

Proof of Correctness Claim (A) Let $p = (x_1, y_1) \in Q, \ q = (x_2, y_2) \in R, \ d(p,q) \le \delta$

Note: Since $d(p,q) \leq \delta, \; |x_1-x_2| \leq \delta \; and \; |y_1-y_2| \leq \delta$ Proof of (A) [p and q are members of S_v i.e. $x_1, x_2 \in [\bar{x} - \delta, \bar{x} + \delta]$]

ネータ トス ネット Strip that defines Sy

Note: $p \in Q => x_1 \le \bar{x}$ and $q \in R => x_2 \ge \bar{x}$. $=> x_1, x_2 \in [\bar{x} - \delta, \bar{x} + \delta]$ nouth where we assume droig 151

Proof of Correctness Claim (B)

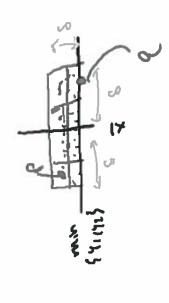
(B): $p = (x_1, y_1)$ and $q = (x_2, y_2)$ are at most 7 positions apart in Sy white in Sy differ by 7 positions.

Mysumythow starts point. We asstyrise early box has zero or one point + Siz them in y-coordinate, two nockes are alrest only to points (That's why we sort why? bus it we have more than one pourt in a bou, it contraddicts - assume of has the Tim Roughgarden O with the assumption that O 8=min, del, b), det, yes? lower y Key Picture : draw $\delta/2 imes \delta/2$ boxes with center $ar{x}$ and bottom min{y₁,y₂}

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Proof of Correctness Claim (B)

Lemma 1: all points of S_y with y-coordinate between those of p and q, inclusive, lie in one of these 8 boxes.



 $\overline{\text{Proof}}$: First, recall y-coordinates of p,q differ by < δ x-coordinates between $\,\bar{x}-\delta\,$ and $\,\bar{x}+\delta\,$ Second, by definition of S_v, all have

O.E.D

dea/b) = 1/2/2 / (2) 2

Proof of Correctness Claim (B) and the smokest

Lemma 2 : At most one point of P in each box.



Proof: by contradiction

Suppose a,b lie in the same box. Then:

I. a,b are either both in Q or both in R \vee

II.
$$d(a,b) \le \frac{\delta}{2} \cdot \sqrt{2} \le \delta$$

But (i) and (ii) contradict the definition of $\,\delta\,$ (as smallest distance between pairs of points in Q or in R)

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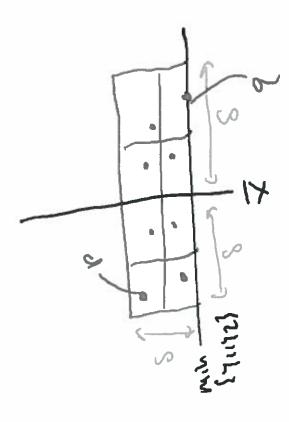
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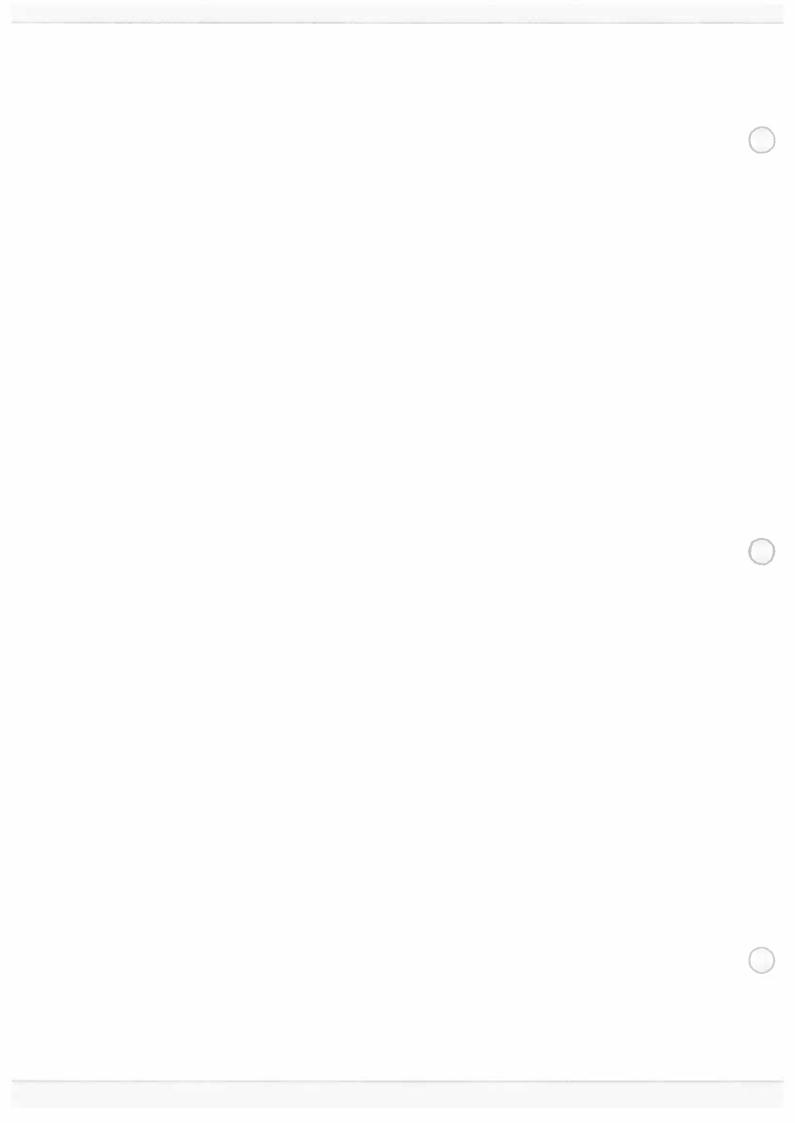
Final Wrap-Up

Lemmas 1 and 2 => at most 8 points in this picture (including p and q)

=> Positions of p,q in S_y differ by at most 7



O.F. D



Master Method

Motivation



Integer Multiplication Revisited

Motivation: potentially useful algorithmic ideas often need mathematical analysis to evaluate Recall: grade-school multiplication algorithm uses $\theta(n^2)$ operation to multiply two n-digit numbers

A Recursive Algorithm

lets assum

Write $x = 10^{n/2}a + b$ $y = 10^{n/2}c + d$ [where a,b,c,d are n/2 – digit numbers] Recursive approach

 $x \cdot y = 10^n ac + 10^{n/2} (ad + bc) + bd$

then compute (*) in the obvious way. Algorithm#1: recursively compute ac, ad, bc, bd,

2

A Recursive Algorithm

T(n) = maximum number of operations this algorithm needs to multiply two n-digit numbers Recurrence: express T(n) in terms of running time of

recursive calls.

Every reactions has two rarts.

| Base Case: T(1) <= a constant.

 $\overline{\text{For all } n > 1}: \quad T(n) \leq 4T(n/2) + O(n)$

Work done by recursive calls

Zecurssive Cills)

here coutside

Work done

(3) wears for examples grading the numbers and although them up. In case of integer mustifiliation there are 4 Terrebuse calls.

A Better Recursive Algorithm

 $(a+b)(c+d)^{(3)}$ [recall ad+bc = (3)-(1)-(2)] $\longrightarrow \text{only } 3 \text{ quantite alls.}$ Algorithm #2 (Gauss): recursively compute ac, bd, bd,

New Recurrence:

Base Case: T(1) <= a constant

Which recurrence best describes the running time of Gauss's algorithm for integer multiplication?

$$\bigcirc \ T(n) \leq 2T(n/2) + O(n^2)$$

$$\rightarrow O3T(n/2) + O(n)$$

$$O 4T(n/2) + O(n)$$

$$O 4T(n/2) + O(n^2)$$

but outside of the recursive calle, the mondy part, how livined our companision with Mage Sort We couly have a recursite calls.

A Better Recursive Algorithm

Algorithm #2 (Gauss): recursively compute ac, bd, $(a+b)(c+d)^{(2)}$ [recall ad+bc = (3) - (1) - (2)]

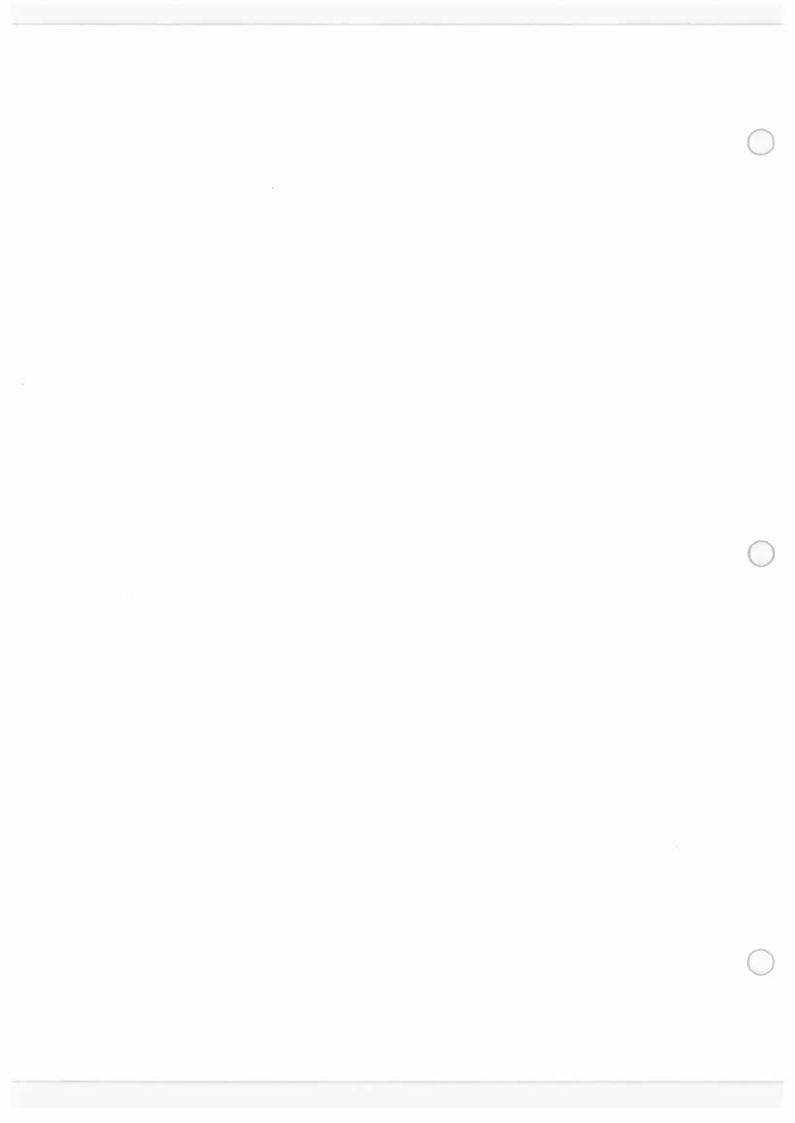
New Recurrence:

For all n>1: $T(n) \leq 3T(n/2) + O(n)$ here characters. Base Case: T(1) <= a constant

Work done

better than 41(2) of This would be

Work done by recursive calls anly 3 recursive calls

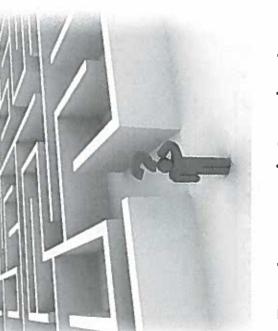


July 29 18 (A)

Master Method

The Precise Statement

= upper bound on the recursiveably. Led to analyse recursive algo. = summed fine



Design and Analysis of Algorithms I

The Master Method

Cool Feature : a "black box" for solving recurrences.

Assumption: all subproblems have equal size.

If the sizes are different, you amont we muster method.

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TWO ingredients of cash maske algo.

Recurrence Format

1. Base Case: T(n) <= a constant for all sufficiently small n

For all larger n :

Ir all larger
$$n$$
:
$$\int_{V} We \text{ ignore constants in big on } T(n) \leq a T(n/b) + O(n^d_{\mathbb{A}}) \text{ notation that not}$$

where

b = input size shrinkage factor (>1) between to be greate then one d = exponent in running time of "combine step" (>=0) a = number of recursive calls (>= 1)[a,b,d independent of n]

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Should be

(Or muster theorem

gives only bigger bounch (worse case).

changes leading constants) Base doesn't matter (only

T(n) =time is hound

 $O(n^d \log n)$

if $a < b^d$ (Case 2) if $a > b^d$ (Case 3) if $a = b^d$ (Case 1) 0(nd) doministed but october

 $O(n^{\log p a})$

Base matters (because that on the experient. makes a

of different by a compant.

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WIKIPEDIA

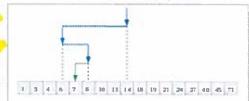
Binary search algorithm

In computer science, binary search, also known as half-interval search, [1] logarithmic search, [2] or binary chop, [3] is a search algorithm that finds the position of a target value within a sorted array. [4][5] Binary search compares the target value to the middle element of the array. If they are not equal, the half in which the target cannot lie is eliminated and the search continues on the remaining half, again taking the middle element to compare to the target value, and repeating this until the target value is found. If the search ends with the remaining half being empty, the target is not in the array. Even though the idea is simple, implementing binary search correctly requires attention to some subtleties about its exit conditions and midpoint calculation.

Binary search runs in at worst logarithmic time, making $O(\log n)$ comparisons, where n is the number of elements in the array, the O is Big O notation, and \log is the logarithm. Binary search takes constant (O(1)) space, meaning that the space taken by the algorithm is the same for any number of elements in the array. Binary search is faster than linear search except for small arrays, but the array must be sorted first. Although specialized data structures designed for fast searching—such as hash tables—can be searched more efficiently, binary search applies to a wider range of problems.

There are numerous variations of binary search. In particular, fractional cascading speeds up binary searches for the same value in multiple arrays, efficiently solving a series of search problems in computational geometry and numerous other fields. Exponential search extends binary search to unbounded lists. The binary search tree and B-tree data structures are based on binary search.

Binary search algorithm



Visualization of the binary search algorithm where 7 is the target value.

Class	Search algorithm
Data structure	Array
Worst-case performance	O(log n)
Best-case performance	O(1)
Average performance	O(log n)
Worst-case space complexity	O(1)

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n > lgn

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Interpolation search
Fractional cascading
Noisy binary search
Quantum binary search

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Algorithm

Binary search works on sorted arrays. Binary search begins by comparing the middle element of the array with the target value. If the target value matches the middle element, its position in the array is returned. If the target value is less than or greater than the middle element, the search continues in the lower or upper half of the array, respectively, eliminating the other half from consideration.^[7]

Procedure

Given an array A of n elements with values or records A_0 , A_1 , ..., A_{n-1} , sorted such that $A_0 \le A_1 \le ... \le A_{n-1}$, and target value T, the following subroutine uses binary search to find the index of T in A. [7]

- 1. Set L to 0 and R to n-1.
- 2. If L > R, the search terminates as unsuccessful.
- 3. Set m (the position of the middle element) to the floor, or the greatest integer less than (L + R)/2.
- 4. If $A_m < T$, set L to m + 1 and go to step 2.
- 5. If $A_m > T$, set R to m 1 and go to step 2.
- 6. Now $A_m = T$, the search is done; return m.

This iterative procedure keeps track of the search boundaries with the two variables. The procedure may be expressed in <u>pseudocode</u> as follows, where the variable names and types remain the same as above, floor is the floor function, and unsuccessful refers to a specific variable that conveys the failure of the search.^[7]

```
function binary_search(A, n, T):
    L := 0
    R := n - 1
    while L <= R:
        m := floor((L + R) / 2)
        if A[m] < T:
            L := m + 1
        else if A[m] > T:
            R := m - 1
        else:
            return m
    return unsuccessful
```

Alternative procedure

In the above procedure, the algorithm checks whether the middle element (m) is equal to the target (T) in every iteration. Some implementations leave out this check during each iteration. The algorithm would perform this check only when one element is left (when L = R). This results in a faster comparison loop, as one comparison is eliminated per iteration. However, it requires one more iteration on average. [8]

The first implementation to leave out this equality check was published by <u>Hermann Bottenbruch</u> in 1962. However, Bottenbruch's version assumes that there is more than one element in the array. The subroutine for Bottenbruch's implementation is as follows, assuming that n > 1:^{[9][8]}

- 1. Set L to 0 and R to n-1.
- 2. If $L \ge R$, go to step 6.
- 3. Set m (the position of the middle element) to the ceiling, or the least integer greater than (L + R)/2.
- 4. If $A_m > T$, set R to m 1 and go to step 2.
- 5. Otherwise, if $A_m \le T$, set L to m and go to step 2.
- 6. Now L = R, the search is done. If $A_L = T$, return L. Otherwise, the search terminates as unsuccessful.

Duplicate elements

The procedure may return any index whose element is equal to the target value, even if there are duplicate elements in the array. For example, if the array to be searched was [1, 2, 3, 4, 4, 5, 6, 7] and the target was 4, then it would be correct for the algorithm to either return the 4th (index 3) or 5th (index 4) element. The regular procedure would return the 4th element (index 3). However, it is sometimes necessary to find the leftmost element or the rightmost element if the target value is duplicated in the array. In the above example, the 4th element is the leftmost element of the value 4, and the 5th element is the rightmost element. The alternative procedure above will always return the index of the rightmost element if an element is duplicated in the array. [9]

Procedure for finding the leftmost element

To find the leftmost element, the following procedure can be used:^[10]

- 1. Set L to 0 and R to n.
- 2. If $L \ge R$, go to step 6.
- 3. Set m (the position of the middle element) to the floor, or the greatest integer less than (L + R)/2.
- 4. If $A_m < T$, set L to m + 1 and go to step 2.
- 5. Otherwise, if $A_m \ge T$, set R to m and go to step 2.
- 6. Now L = R, the search is done, return L.

If L < n and $A_L = T$, then A_L is the leftmost element that equals T. Even if T is not in the array, L is the <u>rank</u> of T in the array, or the number of elements in the array that are less than T.

Where floor is the floor function and == checks for equality, the pseudocode for this version is:

```
function binary_search_leftmost(A, n, T):
    L := 0
    R := n
    while L < R:
        m := floor((L + R) / 2)
        if A[m] < T:
            L := m + 1
        else:
            R := m
    return L</pre>
```

Procedure for finding the rightmost element

To find the rightmost element, the following procedure can be used: [10]

- 1. Set L to 0 and R to n.
- 2. If $L \ge R$, go to step 6.
- 3. Set m (the position of the middle element) to the floor, or the greatest integer less than (L + R)/2.
- 4. If $A_m > T$, set R to m and go to step 2.
- 5. Otherwise, if $A_m \le T$, set L to m + 1 and go to step 2.
- 6. Now L = R, the search is done, return L.

If $L \le n$ and $A_L = T$, then A_L is the rightmost element that equals T. Even if T is not in the array, L is the number of elements in the array that are greater than T.

Where floor is the floor function and == checks for equality, the pseudocode for this version is:

```
function binary_search_rightmost(A, n, T):
    L := 0
    R := n
    while L < R:
        m := floor((L + R) / 2)
        if A[m] > T:
            R := m
        else:
        L := m + 1
    return L
```

Approximate matches

The above procedure only performs *exact* matches, finding the position of a target value. However, due to the ordered nature of sorted arrays, it is trivial to extend binary search to perform approximate matches. For example, binary search can be used to compute, for a given value, its rank (the number of smaller elements), predecessor (next-smallest element), successor (next-largest element), and nearest neighbor. Range queries seeking the number of elements between two values can be performed with two rank queries.^[11]

 Rank queries can be performed using a modified version of binary search. By returning m on a successful search, and L on an unsuccessful search, the number of elements less than the target value is returned instead.^[11] If there are Predecessor / Nearest neighbor

1 2 3 4 7 8 10 11 13 14 15

Rank = 4 Successor

Binary search can be adapted to compute approximate matches. In the example above, the rank, predecessor, successor, and nearest neighbor are shown for the target value 5, which is not in the array.

duplicate elements in the array, then the procedure for finding the leftmost element must be used.

- Predecessor and successor queries can be performed with rank queries. Once the rank of the target value is known, its predecessor is the element at the position given by its rank (as it is the largest element that is smaller than the target value). Its successor is the element after it (if it is present in the array) or at the next position after the predecessor (otherwise). [12] The procedure for finding the rightmost element must be used for computing the successor if there are duplicate elements in the array. The nearest neighbor of the target value is either its predecessor or successor, whichever is closer.
- Range queries are also straightforward. Once the ranks of the two values are known, the number of elements greater
 than or equal to the first value and less than the second is the difference of the two ranks. This count can be adjusted
 up or down by one according to whether the endpoints of the range should be considered to be part of the range and
 whether the array contains keys matching those endpoints.^[13]

Performance

The performance of binary search can be analyzed by reducing the procedure to a binary comparison tree, where the root node is the middle element of the array. The middle element of the lower half is the left child node of the root and the middle element of the upper half is the right child node of the root. The rest of the tree is built in a similar fashion. This model represents binary search; starting from the root node, the left or right subtrees are traversed depending on whether the target value is less or more than the node under consideration, representing the successive elimination of elements. [6][14]

The worst case is iterations of the comparison loop, where the notation denotes the floor function that rounds its argument to the next-smallest integer and is the binary logarithm. The worst case is reached when the search reaches the deepest level of the tree, equivalent to a binary search that has reduced to one element and, in each iteration, always eliminates the smaller subarray out of the two if they are not of equal size. [a][14]

20 40 80 100

A tree representing binary search.

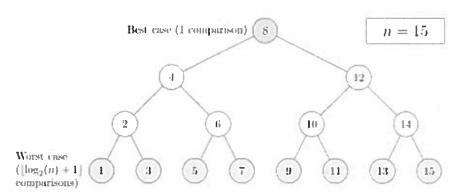
50

A tree representing binary search. The array being searched here is [20, 30, 40, 50, 80, 90, 100], and the target value is 40.

The worst case may also be reached when the target element is not in the array. If is one less than a power of two, then this is always the case. Otherwise, the search may perform , the worst case, or

iterations, one less than the worst case, depending on whether the search reaches the deepest or second-deepest level of the tree.[15]

On average, assuming that each element is equally likely to be searched, binary search makes



The worst case is reached when the search reaches the deepest level of the tree, while the best case is reached when the target value is the middle element.

iterations when the target element is in the array. This is approximately equal to

iterations. When the target element is not in the array, the average case is iterations, assuming that the range between and outside elements is equally likely to be searched.^[14]

In the best case, where the target value is the middle element of the array, its position is returned after one iteration. [16]

In terms of iterations, no search algorithm that works only by comparing elements can exhibit better average and worst-case performance than binary search. This is because the comparison tree representing binary search has the fewest levels possible as each level is filled completely with nodes if there are enough nodes.^[b] Otherwise, the search algorithm can eliminate few elements in an iteration, increasing the number of iterations required in the average and worst case. This is the case for other search algorithms based on comparisons, as while they may work faster on some target values, the average performance over *all* elements is affected. By dividing the array in half, binary search ensures that the size of both subarrays are as similar as possible.^[14]

Performance of alternative procedure

Each iteration of the binary search procedure defined above makes one or two comparisons, checking if the middle element is equal to the target in each iteration. Assuming that each element is equally likely to be searched, each iteration makes 1.5 comparisons on average. A variation of the algorithm checks whether the middle element is equal to the target at the end of the search, eliminating on average half a comparison from each iteration. This slightly cuts the time taken per iteration on most computers, while guaranteeing that the search takes the maximum number of iterations, on average adding one iteration to the search. Because the comparison loop is performed only times in the worst case, the slight increase in comparison loop efficiency does not compensate for the extra iteration for all but enormous .[c][17][18]

Binary search versus other schemes

Sorted arrays with binary search are a very inefficient solution when insertion and deletion operations are interleaved with retrieval, taking time for each such operation, and complicating memory use.^[19] There are other data structures that support much more efficient insertion and deletion. Binary search can be used to perform exact matching and set membership (determining whether a target value is in a collection of values), but there are data structures that support faster exact matching and set membership. However, unlike many other searching schemes, binary search can be used for efficient approximate matching, usually performing such matches in time regardless of the type or structure of the values themselves.^[20] In addition, there are some operations, like finding the smallest and largest element, that can be performed efficiently on a sorted array.^[11]

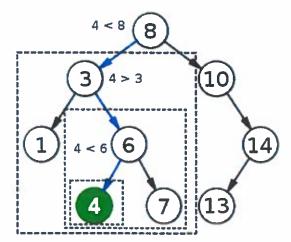
Hashing

For implementing associative arrays, hash tables, a data structure that maps keys to records using a hash function, are generally faster than binary search on a sorted array of records; [21] most implementations require only amortized constant time on average. [d][23] However, hashing is not useful for approximate matches, such as computing the next-smallest, next-largest, and nearest key, as the only information given on a failed search is that the target is not present in any record. [24] Binary search is ideal for such matches performing them in logarithmic time. Binary search also supports approximate matches. Some operations, like finding the smallest and largest element, can be done efficiently on sorted arrays but not on hash tables. [20]

Trees

A binary search tree is a binary tree data structure that works based on the principle of binary search. The records of the tree are arranged in sorted order, and each record in the tree can be searched using an algorithm similar to binary search, taking on average logarithmic time. Insertion and deletion also require on average logarithmic time in binary search trees. This can be faster than the linear time insertion and deletion of sorted arrays, and binary trees retain the ability to perform all the operations possible on a sorted array, including range and approximate queries. [20][25]

However, binary search is usually more efficient for searching as binary search trees will most likely be imperfectly balanced, resulting in slightly worse performance than binary search. This even applies to balanced binary search trees, binary search trees that balance their own nodes, because they rarely produce optimally-balanced trees. Although unlikely, the tree may be severely imbalanced with few internal nodes with two children, resulting in the average and worst-case search time approaching comparisons. [e] Binary search trees take more space than sorted arrays. [27]



Binary search trees are searched using an algorithm similar to binary search.

Binary search trees lend themselves to fast searching in external memory stored in hard disks, as binary search trees can efficiently be structured in filesystems. The B-tree generalizes this method of tree organization; B-trees are frequently used to organize long-term storage such as databases and filesystems. [28][29]

Linear search

Linear search is a simple search algorithm that checks every record until it finds the target value. Linear search can be done on a linked list, which allows for faster insertion and deletion than an array. Binary search is faster than linear search for sorted arrays except if the array is short, although the array needs to be sorted beforehand. [f][31] All sorting algorithms based on comparing elements, such as quicksort and merge sort, require at least comparisons in the worst case. [32] Unlike linear search, binary search can be used for efficient approximate matching. There are operations such as finding the smallest and largest element that can be done efficiently on a sorted array but not on an unsorted array. [33]

Set membership algorithms

A related problem to search is <u>set membership</u>. Any algorithm that does lookup, like binary search, can also be used for set membership. There are other algorithms that are more specifically suited for set membership. A <u>bit array</u> is the simplest, useful when the range of keys is limited. It compactly stores a collection of <u>bits</u>, with each bit representing a single key within the range of keys. Bit arrays are very fast, requiring only time.^[34] The Judy1 type of <u>Judy</u> array handles 64-bit keys efficiently.^[35]

For approximate results, Bloom filters, another probabilistic data structure based on hashing, store a set of keys by encoding the keys using a bit array and multiple hash functions. Bloom filters are much more space-efficient than bit arrays in most cases and not much slower: with hash functions, membership queries require only time. However, Bloom filters suffer from false positives. [9][h][37]

Other data structures

There exist data structures that may improve on binary search in some cases for both searching and other operations available for sorted arrays. For example, searches, approximate matches, and the operations available to sorted arrays can be performed more efficiently than binary search on specialized data structures such as van Emde Boas trees, fusion trees, tries, and bit arrays. However, while these operations can always be done at least efficiently on a sorted array regardless of the keys, such data structures are usually only faster because they exploit the properties of keys with a certain attribute (usually keys that are small integers), and thus will be time or space consuming for keys that lack that attribute. [20] Some structures, such as Judy arrays, use a combination of approaches to mitigate this while retaining efficiency and the ability to perform approximate matching. [35]

Variations

Uniform binary search

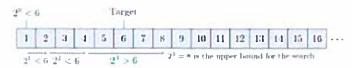
Uniform binary search stores, instead of the lower and upper bounds, the index of the middle element and the change in the middle element from the current iteration to the next iteration. Each step reduces the change by about half. For example, if the array to be searched was [1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11], the middle element would be 6. Uniform binary search works on the basis that the difference between the index of middle element of the array and the left and right subarrays is the same. In this case, the middle element of the left subarray ([1, 2, 3, 4, 5]) is 3 and the middle element of the right subarray ([7, 8, 9, 10, 11]) is 9. Uniform binary search would store the value of 3 as both indices differ from 6 by this same amount. To reduce the search space, the algorithm either adds or subtracts this change from the middle element. The main advantage of uniform binary search is that the

procedure can store a table of the differences between indices for each iteration of the procedure. Uniform binary search may be faster on systems where it is inefficient to calculate the midpoint, such as on decimal computers.^[39]

Uniform binary search stores the difference between the current and the two next possible middle elements instead of specific bounds.

Exponential search

Exponential search extends binary search to unbounded lists. It starts by finding the first element with an index that is both a power of two and greater than the target value. Afterwards, it sets that index as the upper bound, and switches to binary search. A search takes iterations of the exponential search and at most iterations of the binary search, where \boldsymbol{x} is the position of the target value. Exponential search works on bounded lists, but becomes an improvement over binary search only if the target value lies near the beginning of the array. [40]



Visualization of exponential searching finding the upper bound for the subsequent binary search.

Interpolation search

Instead of calculating the midpoint, interpolation search estimates the position of the target value, taking into account the lowest and highest elements in the array as well as length of the array. This is only possible if the array elements are numbers. It works on the basis that the midpoint is not the best guess in many cases. For example, if the target value is close to the highest element in the array, it is likely to be located near the end of the array. When the distribution of the array elements is uniform or near uniform, it makes comparisons. [41][42][43]



Visualization of interpolation search. In this case, no searching is needed because the estimate of the target's location within the array is correct. Other implementations may specify another function for estimating the target's location.

In practice, interpolation search is slower than binary search for small arrays, as interpolation search requires extra computation. Although its time complexity grows more slowly than binary search, this only compensates for the extra computation for large arrays.^[41]

Fractional cascading

Fractional cascading is a technique that speeds up binary searches for the same element in multiple sorted arrays. Searching each array separately requires time, where is the number of arrays. Fractional cascading reduces this to by storing specific information in each array about each element and its position in the other arrays. [44][45]

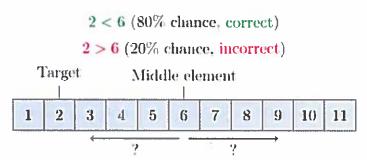
Fractional cascading was originally developed to efficiently solve various <u>computational geometry</u> problems, but it also has been applied elsewhere, in domains such as data mining and Internet Protocol routing.^[44]

Noisy binary search

Noisy binary search algorithms solve the case where the algorithm cannot reliably compare elements of the array. For each pair of elements, there is a certain probability that the algorithm makes the wrong comparison. Noisy binary search can find the correct position of the target with a given probability that controls the reliability of the yielded position. [NIII][49][50]

Quantum binary search

Classical computers are bounded to the worst case of exactly iterations when performing binary search. Quantum algorithms for binary search are still bounded to a proportion of queries (representing iterations of the classical procedure), but the constant factor is less than one, providing for faster performance on



Goal: Search for the index of the target value so that there is a probability of p that the index is correct. p is specified before the search.

In noisy binary search, there is a certain probability that a comparison is incorrect.

quantum computers. Any *exact* quantum binary search procedure—that is, a procedure that always yields the correct result—requires at least — queries in the worst case, where is the natural logarithm.^[51] There is an exact quantum binary search procedure that runs in queries in the worst case.^[52] In comparison, Grover's algorithm is the optimal quantum algorithm for searching an unordered list of elements, and it requires — queries.^[53]

History

In 1946, John Mauchly made the first mention of binary search as part of the Moore School Lectures, a seminal and foundational college course in computing. [9] In 1957, William Wesley Peterson published the first method for interpolation search. [9][54] Every published binary search algorithm worked only for arrays whose length is one less than a power of two [k] until 1960, when Derrick Henry Lehmer published a binary search algorithm that worked on all arrays. [56] In 1962, Hermann Bottenbruch presented an ALGOL 60 implementation of binary search that placed the comparison for equality at the end, increasing the average number of iterations by one, but reducing to one the number of comparisons per iteration. [8] The uniform binary search was developed by A. K. Chandra of Stanford University in 1971. [9] In 1986, Bernard Chazelle and Leonidas J. Guibas introduced fractional cascading as a method to solve numerous search problems in computational geometry. [44][57][58]

Implementation issues

Although the basic idea of binary search is comparatively straightforward, the details can be surprisingly tricky ... — Donald Knuth^[2]

When Jon Bentley assigned binary search as a problem in a course for professional programmers, he found that ninety percent failed to provide a correct solution after several hours of working on it, mainly because the incorrect implementations failed to run or returned a wrong answer in rare edge cases. [59] A study published in 1988 shows that accurate code for it is only found in five out of twenty textbooks. [60] Furthermore, Bentley's own implementation of binary search, published in his 1986 book *Programming Pearls*, contained an overflow error that remained undetected for over twenty years. The Java programming language library implementation of binary search had the same overflow bug for more than nine years. [61]

In a practical implementation, the variables used to represent the indices will often be of fixed size, and this can result in an arithmetic overflow for very large arrays. If the midpoint of the span is calculated as (L + R)/2, then the value of L + R may exceed the range of integers of the data type used to store the midpoint, even if L and R are within the range. If L and R are nonnegative, this can be avoided by calculating the midpoint as L + (R - L)/2. [62]

If the target value is greater than the greatest value in the array, and the last index of the array is the maximum representable value of L, the value of L will eventually become too large and overflow. A similar problem will occur if the target value is smaller than the least value in the array and the first index of the array is the smallest representable value of R. In particular, this means that R must not be an unsigned type if the array starts with index 0.160[62]

An infinite loop may occur if the exit conditions for the loop are not defined correctly. Once *L* exceeds *R*, the search has failed and must convey the failure of the search. In addition, the loop must be exited when the target element is found, or in the case of an implementation where this check is moved to the end, checks for whether the search was successful or failed at the end must be in place. Bentley found that most of the programmers who incorrectly implemented binary search made an error in defining the exit conditions.^[8][63]

Library support

Many languages' standard libraries include binary search routines:

- C provides the function bsearch() in its standard library, which is typically implemented via binary search, although
 the official standard does not require it so.^[64]
- C++'s Standard Template Library provides the functions binary_search(), lower_bound(), upper_bound()
 and equal_range().^[65]
- COBOL provides the SEARCH ALL verb for performing binary searches on COBOL ordered tables. [66]
- Go's sort standard library package contains the functions Search, SearchInts, SearchFloat64s, and SearchStrings, which implement general binary search, as well as specific implementations for searching slices of integers, floating-point numbers, and strings, respectively.^[67]
- Java offers a set of overloaded binarySearch() static methods in the classes Arrays (https://docs.oracle.com/javase/10/docs/api/java/util/Arrays.html) and Collections (https://docs.oracle.com/javase/10/docs/api/java/util/Collections.html) in the standard java.util package for performing binary searches on Java arrays and on Lists, respectively.[68][69]
- Microsoft's .NET Framework 2.0 offers static generic versions of the binary search algorithm in its collection base classes. An example would be System. Array's method BinarySearch<T>(T[] array, T value).^[70]
- For Objective-C, the Cocoa framework provides the NSArray indexOfObject:inSortedRange:options:usingComparator: (https://developer.apple.com/library/mac/documentation/Cocoa/Reference/Foundation/Classes/NSArray_Class/NSArray.html#//apple_ref/occ/instm/NSArray/indexOfObject:inSortedRange:options:usingComparator:) method in Mac OS X 10.6+. [71] Apple's Core Foundation C framework also contains a CFArrayBSearchValues() (https://developer.apple.com/library/mac/documentation/CoreFoundation/Reference/CFArrayRef/Reference/reference.html#//apple_ref/c/func/CFArrayBSearchValues) function. [72]
- Python provides the bisect module.^[73]
- Ruby's Array class includes a bsearch method with built-in approximate matching.^[74]

See also

- Bisection method the same idea used to solve equations in the real numbers
- Multiplicative binary search binary search variation with simplified midpoint calculation

Notes and references

Notes

- a. This happens as binary search will not always divide the array perfectly. Take for example the array [1, 2, ..., 16]. The first iteration will select the midpoint of 8. On the left subarray are eight elements, but on the right are nine. If the search takes the right path, there is a higher chance that the search will make the maximum number of comparisons.^[14]
- b. Any search algorithm based solely on comparisons can be represented using a binary comparison tree. An internal path is any path from the root to an existing node. Let I be the internal path length, the sum of the lengths of all internal paths. If each element is equally likely to be searched, the average case is - or simply one plus the average of all the internal path lengths of the tree. This is because internal paths represent the elements that the search algorithm compares to the target. The lengths of these internal paths represent the number of iterations after the root node. Adding the average of these lengths to the one iteration at the root yields the average case. Therefore, to minimize the average number of comparisons, the internal path length must be minimized. It turns out that the tree for binary search minimizes the internal path length. Knuth 1998 proved that the external path length (the path length over all nodes where both children are present for each already-existing node) is minimized when the external nodes (the nodes with no children) lie within two consecutive levels of the tree. This also applies to internal paths as internal path length Iis linearly related to external path length . For any tree of nodes, . When each subtree has a similar number of nodes, or equivalently the array is divided into halves in each iteration, the external nodes as well as their interior parent nodes lie within two levels. It follows that binary search minimizes the number of average comparisons as its comparison tree has the lowest possible internal path length. [14]
- c. Knuth 1998 showed on his MIX computer model, intended to represent an ordinary computer, that the average running time of this variation for a successful search is units of time compared to units for regular binary search. The time complexity for this variation grows slightly more slowly, but at the cost of higher initial complexity. [17]
- d. It is possible to perform hashing in guaranteed constant time. [22]
- e. The *worst* binary search tree for searching can be produced by inserting the values in sorted order or in an alternating lowest-highest key pattern. [26]
- f, Knuth 1998 performed a formal time performance analysis of both of these search algorithms. On Knuth's MIX computer, which Knuth designed as a representation of an ordinary computer, binary search takes on average units of time for a successful search, while linear search with a sentinel node at the end of the list takes—units. Linear search has lower initial complexity because it requires minimal computation, but it quickly outgrows binary search in complexity. On the MIX computer, binary search only outperforms linear search with a sentinel if [14][30]
- g. This is because simply setting all of the bits which the hash functions point to for a specific key can affect queries for other keys which have a common hash location for one or more of the functions.^[36]
- h. There exist improvements of the Bloom filter which improve on its complexity or support deletion; for example, the cuckoo filter exploits **cuckoo hashing** to gain these advantages.^[36]

entropy function and is the probability that the procedure yields the wrong position. [46]

- j. The noisy binary search problem can be considered as a case of the <u>Rényi-Ulam game</u>, ^[47] a variant of <u>Twenty</u> <u>Questions</u> where the answers may be wrong. ^[48]
- k. That is, arrays of length 1, 3, 7, 15, 31 ... [55]

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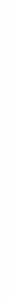
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This page was last edited on 18 July 2018, at 19:51 (UTC).

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Master Method

Examples

Design and Analysis of Algorithms I

The Master Method

If
$$T(n) \leq aT\left(\frac{n}{b}\right) + O(n^d)$$
 were outside the recussion

factor sub problem Size is smaller than the creasinal problem size then

$$\lceil O(n^d \log n) \rceil$$

$$T(n) = \begin{cases} o(n^d) \\ o(n^{\log b a}) \end{cases}$$

if
$$a = b^d$$
 (Case 1)
if $a < b^d$ (Case 2)

if
$$a > b^d$$
 (Case 3)

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Example #1

Merge Sort

$$\begin{vmatrix} \mathbf{a} = 2 \\ \mathbf{b} = 2 \\ \mathbf{d} = 1 \end{vmatrix} b^d = \begin{vmatrix} \mathbf{a} = \mathbf{b} \\ \mathbf{d} = 1 \end{vmatrix} > Case 1$$

$$\mathbf{d} = \mathbf{1}$$

$$\mathbf{T}(n) = O(n^d \log n) = O(n \log n)$$

Where are the respective values of a, b, d for a binary search of a sorted array, and which case of the Master Method does this correspond to?

$$a = b^d = > T(n) = O(n^d \log n) = O(\log n)$$

O 1, 2, 1 [Case 2]

O 2, 2, 0 [Case 3]

O 2, 2, 1 [Case 1]

y multiplications: ou ad be bot

Example #3

Integer Multiplication Algorithm # 1 (10)10 Gauss mich

$$\sqrt{b} = 2$$
 $\sqrt{b} = 2$ $\sqrt{case 3}$

$$=> T(n) = O(n^{\log_b a}) = O(n^{\log_2 4})$$

adding and mulaphahor

 $=O(n^2)$

Where are the respective values of a,b,d for Gauss's recursive integer multiplication algorithm, and which case of the Master Method does this correspond to?

O 2, 2, 1 [Case 1]

O 3, 2, 1 [Case 1]

algorithm!!!

Better than

the grade-

school

O 3, 2, 1 [Case 2]

 $a = 3, b^d = 2 a > b^d (Case 3)$ O 3, 2, 1 [Case 3] $=> T(n) = O(n^{\log_2 3}) = O(n^{1.59})$

Example #5

Strassen's Matrix Multiplication Algorithm

Matrix of size 10- matrix of singe 4

$$a=7$$
 recursive only:
 $b=2$ why
 $d=2$ $b^d=4< a$ (Case 3)

$$=> T(n) = O(n^{\log_2 7}) = O(n^{2.81})$$

=> beats the naïve iterative algorithm!

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Example #6

Fictitious Recurrence

$$T(n) \le 2T(n/2) + O(n^2)$$

$$\Rightarrow a = 2$$

$$\Rightarrow b = 2 \Rightarrow d = 2 \Rightarrow Case 2)$$

$$\Rightarrow T(n) = O(n^2)$$





Master Method

Proof (Part I)

Design and Analysis of Algorithms I

The conall each case is more important than the proof well.

The Master Method

If
$$T(n) \le aT\left(\frac{n}{b}\right) + O(n^d)$$

then

$$T(n) = \begin{cases} O(n^d \log n) & \text{if } a = b^d \text{ (Case 1)} \\ O(n^d) & \text{if } a < b^d \text{ (Case 2)} \\ O(n^{\log b a}) & \text{if } a > b^d \text{ (Case 3)} \end{cases}$$

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Preamble

Assume: recurrence is

I.
$$T(1) \leq c$$

II.
$$T(n) \leq aT(n/b) + cn^d$$
 constant c) And n is a power of b.

/// And n is a power of b.

(general case is similar, but more tedious)

<u>Idea</u>: generalize MergeSort analysis.

(i.e., use a recursion tree

statement: at each level j = 0,1,2,..., logon, there are
blank> What is the pattern? Fill in the blanks in the following subproblems, each of size <blank> # of times you can divide n by b before reaching 1 a's subproblems, of sign in

O a^j and n/a^j, respectively.

○ a^j and n/b^j, respectively.

O b^j and n/a^j, respectively.

O b^j and n/b^j, respectively.

The Recursion Tree

input stor

अंदर इच्छा

Level 0

a braches

Level 1

. .

Level log_bn

0

Base cases (size 1)

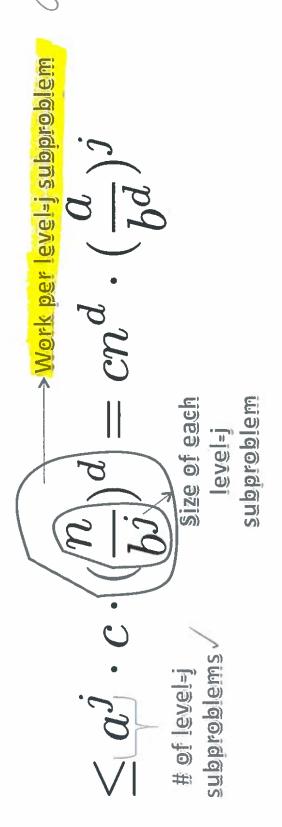
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Cames

Poof I-3

Work at a Single Level

Total work at level j [ignoring work in recursive calls]

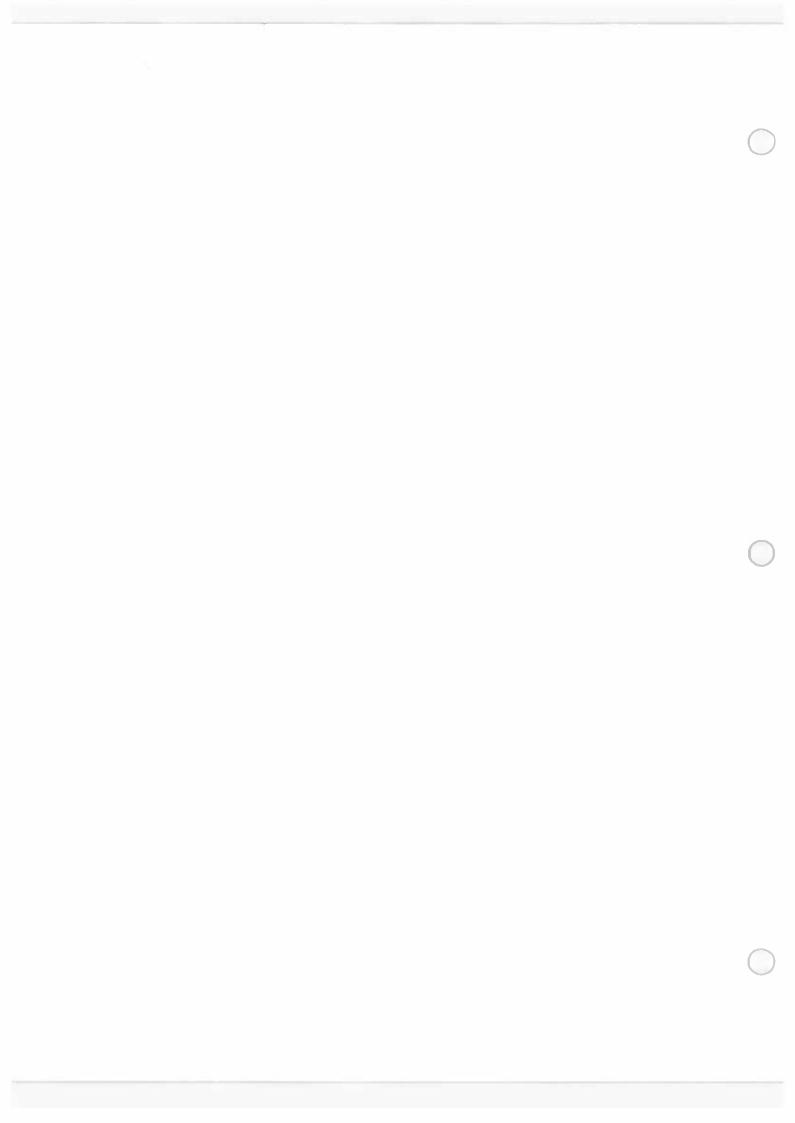


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Total Work

Summing over all levels $j = 0,1,2,..., log_b n$:

Total
$$\leq cn^d \cdot \sum_{j=0}^{\log_b n} (\frac{a}{b^d})^j$$
 (*)



July 21, 2018 (2)

Master Method

Intuition for the 3 Cases

Design and Analysis of Algorithms I



How To Think About (*)

Our upper bound on the work at level i:

 $cn^d \times (\frac{a}{hd})^j$

The cores are alreally saying bear or change with I

Interpretation

bd = rate of work shrinkage (RWS) good south for each problem with: bes as we so down the tree there are more a = rate of subproblem proliferation (RSP)

(ber subproblem)

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Which of the following statements are true? (Check all that apply.)

- recursion levelj. If RSP < RWS, then the amount of work is decreasing with the
 - ☐ If RSP > RWS, then the amount of work is increasing with the recursion level j.
- No conclusions can be drawn about how the amount of work varies with the recursion level j unless RSP and RWS are equal.
- If RSP and RWS are equal, then the amount of work is the same at every recursion level j.

a = 1 = (a) remains the some

Down to is decreasing with the level (2001) (3) work is intreasing in the level In fact the three case are: (1) work done pur wow is the same (laves are more)

Intuition for the 3 Cases

Upper bound for level j: $cn^d \times (\frac{a}{b^d})^j$

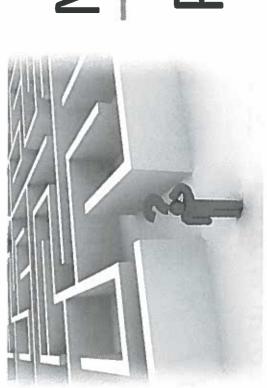
1. RSP = RWS => Same amount of work each level (like [expect O(ndlog(n)] Merge Sort) 2. RSP < RWS => less work each level => most work at the root for the [might expect O(n^d)]

[might expect O(# leaves)] 3. RSP > RWS => more work each level => most work at the leaves

0 (n Gra

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Master Method

Proof (Part II)

Design and Analysis of Algorithms I



Total work: $\leq cn^d \times \sum_{j=0}^{\log_b n} \binom{a}{b^d}$

case I.I.f $a=b^d$, then

$$\leq cn^a \times$$

$$\leq cn^a \times \sum_j$$

*

$$(*) = cn^d(\log_b n + 1)$$

$$= O(n^d \log n)$$

 $[\ \mathsf{end}\ \mathsf{Case}\ 1\]$

geometric serus

Basic Sums Fact

For r
eq 1 , we have

$$1 + r + r^2 + r^3 + \dots + r^k = \frac{r^{k+1} - 1}{r^{k+1} - 1}$$

Proof: by induction (you check)

Upshot:

- = a constant wey hist 1. If r<1 is constant, RHS is <= $\frac{1}{1-r}$ i.e., 1st term of sum dominates
- 2. If r>1 is constant, RHS is <= $\langle r^k \rangle / (1 + \frac{1}{2})$ i.e., last term of sum dominates

Independent of k

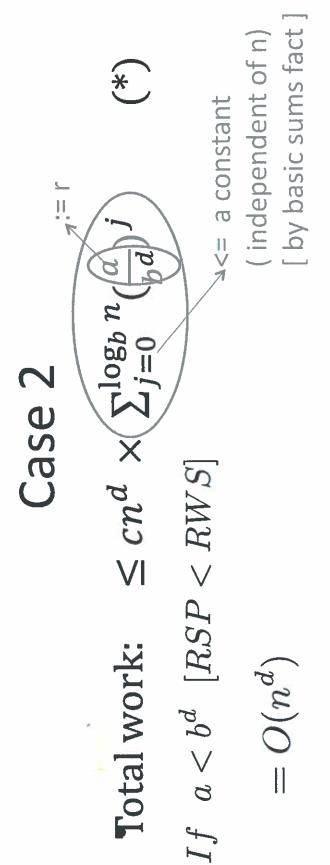
" the control to the tent

Istant dominate ((1))

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at most lets say

Poot I 2



[total work dominated by top level]

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a := r > 1

Total work:
$$\leq cn^d \times (\sum_{j=0}^{\log_b n} \binom{a}{b^d}$$

$$\log_{b} n \binom{a}{bq} j$$

largest term

$$(*) = O(n^d \cdot (\frac{a}{b^d})^{\log_b n})$$

 $If \ a > b^d \ [RSP > RWS]$

Note:
$$b^{-d\log_b n} = (b^{\log_b n})^{-d} = n^{-d}$$

$$So : (*) = O(a^{\log_b n})$$

Level 0

Level 1



of leaves = $a^{\log_b n}$

Which of the following quantities is equal to $a^{\log_b n}$?

Level log_bn

- O The number of levels of the recursion tree.
- O The number of nodes of the recursion tree.
- O The number of edges of the recursion tree.
- → The number of leaves of the recursion tree.

Case 3 continued

Total work: $\leq cn^d \times \sum_{j=0}^{\log b} {n \left(\frac{a}{b^d}\right)^j}$

 $So: (*) = O(a^{\log_b n}) = O(\# leaves)$

-Simpler to apply - More intuitive Note: $(a^{\log_b n}) = (n^{\log_b a}) \leftarrow$

 $[Since (\log_b n)(\log_b a) = (\log_b a)(\log_b n)]$

[End Case 3]

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The Master Method

If
$$T(n) \le aT\left(\frac{n}{b}\right) + O(n^d)$$

then

$$T(n) = \begin{cases} O(n^d \log n) & \text{if } a \\ O(n^d) & \text{if } a \\ O(n^{\log b a}) & \text{if } a \end{cases}$$

$$if a = b^d \text{ (Case 1)}$$

if
$$a < b^d$$
 (Case 2) if $a > b^d$ (Case 3)