

Chapter 4: Divide and Conquer

Divide each difficulty into as many parts as is
feasible and necessary to resolve it.

(Rene Descartes)

Introduction	2
Illustration	3
General Divide-and-Conquer Recurrence	4
The Master Theorem	5
Mergesort	6
Mergesort Algorithm	6
Merge Algorithm	7
Illustration.	8
Comments on Mergesort	9
Analysis of Mergesort	10
Quicksort	11
Quicksort Algorithm	11
Partition Algorithm.	12
Illustration.	13
Comments on Quicksort	14
Binary Search	15
Binary Search Algorithm	15
Illustration.	16
Comments on Binary Search	17
Closest Pair	18
Idea for Closest Pair	18

Closest Pair Algorithm	19
Comments on Closest Pair.	20
Convex Hull	21
First Idea for Convex Hull	21
Second Idea for Convex Hull	22
Quickhull Algorithm	23
Quickhull Algorithm Continued.	24
Comments on Quickhull	25

Introduction

Divide-and-conquer is an approach to solving a problem by:

- Divide an instance into smaller instances of the problem.
- Solve the smaller instances.
- Combine the solutions of the smaller instances into a solution for the larger instance.

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Chapter 2: Slide – 2

Illustration

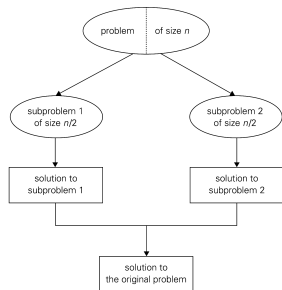


FIGURE 4.1 Divide-and-conquer technique (typical case)

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Chapter 2: Slide – 3

General Divide-and-Conquer Recurrence

The general case is to divide an instance of size n into a instances of size n/b , taking $f(n)$ time to divide and combine.

$$T(n) = \begin{matrix} & a & T(n/b) & + & f(n) \\ \text{time for} & \text{number of} & \text{time for} & & \text{time for} \\ \text{size } n & \text{subinstances} & \text{size } n/b & & \text{divide \& combine} \end{matrix}$$

Some examples are:

$$\begin{array}{ll} \text{Mergesort} & T(n) = 2T(n/2) + \Theta(n) \\ \text{Binary Search} & T(n) = T(n/2) + \Theta(1) \\ \text{Strassen's} & T(n) = 7T(n/2) + \Theta(n^2) \end{array}$$

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Chapter 2: Slide – 4

The Master Theorem

For the recurrence $T(n) = aT(n/b) + f(n)$, if $f(n) \in \Theta(n^d)$, then

$$T(n) \in \begin{cases} \Theta(n^d) & \text{if } a < b^d \\ \Theta(n^d \log n) & \text{if } a = b^d \\ \Theta(n^{\log_b a}) & \text{if } a > b^d \end{cases}$$

Note: $T(n) \geq f(n)$. Also $T(n) \geq a^{\log_b n} = n^{\log_b a}$ because recurrence tree has at least $a^{\log_b n}$ nodes. Examples:

$$\text{Mergesort} \quad T(n) = 2T(n/2) + \Theta(n) \quad \Theta(n \log n)$$

$$\text{Binary Search} \quad T(n) = T(n/2) + \Theta(1) \quad \Theta(\log n)$$

$$\text{Strassen's} \quad T(n) = 7T(n/2) + \Theta(n^2) \quad \Theta(n^{\log_2 7})$$

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Chapter 2: Slide – 5

Mergesort

6

Mergesort Algorithm

```

algorithm Mergesort( $A[0..n-1]$ )
    // Sorts a given array by mergesort
    // Input: An array  $A$  of orderable elements
    // Output: Array  $A[0..n-1]$  in ascending order
    if  $n \leq 1$  then
        copy  $A[0..\lfloor n/2 \rfloor - 1]$  to  $B[0..\lfloor n/2 \rfloor - 1]$ 
        copy  $A[\lfloor n/2 \rfloor..n-1]$  to  $C[0..\lfloor n/2 \rfloor - 1]$ 
        Mergesort( $B$ )
        Mergesort( $C$ )
        Merge( $B, C, A$ )
    
```

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Chapter 2: Slide – 6

Merge Algorithm

```

algorithm Merge( $B[0..p-1], C[0..q-1],$ 
                 $A[0..p+q-1]$ )
    // Merges two sorted arrays into one array
    // Input: Sorted arrays  $B$  and  $C$ 
    // Output: Sorted array  $A$ 
     $i \leftarrow 0; j \leftarrow 0$ 
    for  $k \leftarrow 0$  to  $p+q-1$  do
        if  $i < p$  and ( $j = q$  or  $B[i] \leq C[j]$ ) then
             $A[k] \leftarrow B[i]; i \leftarrow i + 1$ 
        else
             $A[k] \leftarrow C[j]; j \leftarrow j + 1$ 

```

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Chapter 2: Slide – 7

Illustration

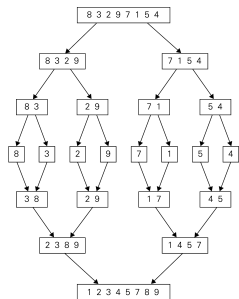


FIGURE 4.2 Example of mergesort operation

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Chapter 2: Slide – 8

Comments on Mergesort

- Note that $n = \lfloor n/2 \rfloor + \lceil n/2 \rceil$ in *Mergesort*.
- Note that $0 \leq i \leq p$, $0 \leq j \leq q$, and $i + j = k$ in *Merge* algorithm. Comparison only happens if both $i < p$ and $j < q$.
- 2 recursive calls to *Mergesort* on instances of size $n/2$. Copy and merge is $\Theta(n)$.
- Constant factor can be improved by more efficient use of arrays.

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Chapter 2: Slide – 9

Analysis of Mergesort

- The number of comparisons $C(n)$ is:
- Base cases for $n =$ a power of 2: $C(1) = 0$, $C(2) = 1$, $C(4) \leq 5$, $C(8) \leq 17$.
- Induction assuming $C(n/2) \leq (n/2) \log_2(n/2)$

$$\begin{aligned}
 C(n) &\leq 2C(n/2) + n - 1 \\
 &\leq 2(n/2) \log_2(n/2) + n - 1 \\
 &\leq n \log_2(n/2) + n - 1 \\
 &\leq n \log_2 n - n \log_2 2 + n - 1 \\
 &\leq n \log_2 n - n + n - 1 \\
 &\leq n \log_2 n
 \end{aligned}$$

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Chapter 2: Slide – 10

Quicksort

11

Quicksort Algorithm

```
algorithm Quicksort( $A[l..r]$ )  
  // Sorts a subarray by quicksort  
  // Input: An subarray of  $A$   
  // Output: Array  $A[l..r]$  in ascending order  
  if  $l < r$  then  
     $p \leftarrow \text{Partition}(A[l..r])$  //  $p$  is index of pivot  
    Quicksort( $A[l..p-1]$ )  
    Quicksort( $A[p+1..r]$ )
```

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Chapter 2: Slide – 11

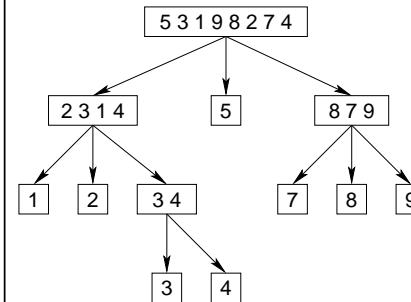
Partition Algorithm

```
algorithm Partition( $A[l..r]$ )  
  // Partitions a subarray using  $A[l]$  as pivot  
  // Input: Subarray of  $A$   
  // Output: Final position of pivot  
   $pivot \leftarrow A[l]$   
   $i \leftarrow l; j \leftarrow r + 1$   
  repeat  
    repeat  $i \leftarrow i + 1$  until  $A[i] \geq pivot$   
    repeat  $j \leftarrow j - 1$  until  $A[j] \leq pivot$   
    if  $i < j$  then swap  $A[i]$  and  $A[j]$   
  until  $i \geq j$   
  swap  $A[l]$  and  $A[j]$   
  return  $j$ 
```

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Chapter 2: Slide – 12

Illustration



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Chapter 2: Slide – 13

Comments on Quicksort

- *Partition* ensures that if p is the final position of the pivot, then $i < p$ implies $A[i] \leq A[p]$ and $i > p$ implies $A[i] \geq A[p]$.
- Best Case: If *Partition* always splits subarray in half, then $T(n) = 2T(n/2) + \Theta(n) \in \Theta(n \log n)$.
- Worst Case: If pivot is always picks min or max, then $T(n) = T(n-1) + \Theta(n) \in \Theta(n^2)$.
- Average Case: Randomly chosen pivot is $\Theta(n \log n)$.
- In median-of-three partitioning, pivot is median of leftmost, rightmost, and middle element.
- Finding median is $\Theta(n)$ with a modification.

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Chapter 2: Slide – 14

Binary Search

15

Binary Search Algorithm

```

algorithm BinarySearch( $A[0..n-1], K$ )
    // Searches a sorted array using binary search
    // Input: An sorted array  $A$  and a key  $K$ 
    // Output: The index of  $K$  or  $-1$ 
     $l \leftarrow 0; r \leftarrow n - 1$ 
    while  $l \leq r$  do
         $m \leftarrow \lfloor (l + r) / 2 \rfloor$ 
        if  $K = A[m]$  then return  $m$ 
        else if  $K < A[m]$  then  $r \leftarrow m - 1$ 
        else  $l \leftarrow m + 1$ 
    return  $-1$ 
    
```

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Chapter 2: Slide – 15

Illustration

	K	31											
values in A	3	14	27	31	39	42	55	70	74	81	85	93	98
indices of A	0	1	2	3	4	5	6	7	8	9	10	11	12
iteration 1	l					m							r
iteration 2	l		m					r					
iteration 3				l	m	r							
iteration 4					l, m, r								

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Chapter 2: Slide – 16

Comments on Binary Search

- *BinarySearch* ensures that $A[l-1] < K$ (or $l = 0$) and $K < A[r+1]$ (or $r = n-1$).
- The number of possible indices is $r - l + 1$, which reduces by half or more each iteration.
- The number of times n can be divided by 2 until it becomes < 1 is about $\log_2 n$ (exact $1 + \lfloor \log_2 n \rfloor$)

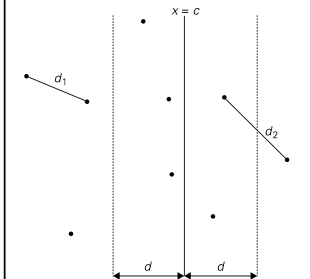
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Chapter 2: Slide – 17

Closest Pair

18

Idea for Closest Pair



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Chapter 2: Slide – 18

Closest Pair Algorithm

```

algorithm ClosestPair( $P$ )
    // Finds closest pair of points
    // Input: A set of  $n$  points sorted by coordinates
    // Output: Distance between closest pair
    if  $n < 2$  then return  $\infty$ 
    else if  $n = 2$  then return distance between pair
    else
         $c \leftarrow$  median value for  $x$  coordinate
         $d_1 \leftarrow \text{ClosestPair}(\text{points with } x < c)$ 
         $d_2 \leftarrow \text{ClosestPair}(\text{points with } x > c)$ 
         $d \leftarrow \min(d_1, d_2)$ 
         $d_3 \leftarrow$  process points with  $c - d < x < c + d$ 
        return  $\min(d, d_3)$ 
    
```

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Chapter 2: Slide – 19

Comments on Closest Pair

- Preprocessing can sort the points by their x and y coordinates (use two lists). Recursive calls can create sorted lists for subsets of points.
- The calculation of d_3 is $\Theta(n)$.
 - Select points with $c - d < x < c + d$.
 - Examine points in order of y values.
 - For any point (x, y) , there can only be a few points within $(x \pm d, y \pm d)$.
- Preprocessing is $\Theta(n \log n)$, and recurrence is $T(n) = 2T(n/2) + \Theta(n) \in \Theta(n \log n)$.

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Chapter 2: Slide – 20

Convex Hull

21

First Idea for Convex Hull

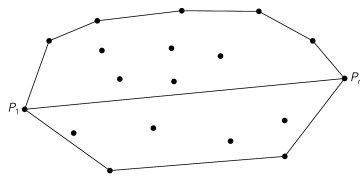


FIGURE 4.8 Upper and lower hulls of a set of points

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Chapter 2: Slide – 21

Second Idea for Convex Hull

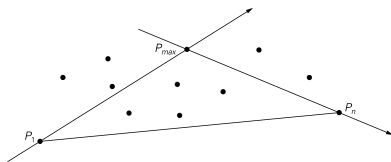


FIGURE 4.9 The idea of quickhull

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Chapter 2: Slide – 22

Quickhull Algorithm

algorithm *Quickhull*(P)

```
// Finds convex hull
// Input: A set of  $n$  points
// Output: A list of points, all of convex hull
 $P_1 \leftarrow$  point in  $P$  with lowest  $x$  value
 $P_2 \leftarrow$  point in  $P$  with highest  $x$  value
 $A \leftarrow$  points in  $P$  "above"  $\overline{P_1P_2}$ 
 $B \leftarrow$  points in  $P$  "below"  $\overline{P_1P_2}$ 
return  $P_1$ ,  $QHWalk(P_1, P_2, A)$ ,
 $P_2$ , and  $QHWalk(P_2, P_1, B)$ 
```

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Chapter 2: Slide – 23

Quickhull Algorithm Continued

algorithm *QHWalk*(P_1, P_2, P)

```
// Finds convex hull clockwise from  $P_1$  to  $P_2$ 
// Input: Two points and a set of  $n$  points
// Output: A list of points, part of convex hull
if  $n \leq 1$  then return  $P$ 
else
     $P_3 \leftarrow$  point in  $P$  farthest from  $\overline{P_1P_2}$ 
     $A \leftarrow$  points in  $P$  "left" of  $\overline{P_1P_3}$ 
     $B \leftarrow$  points in  $P$  "left" of  $\overline{P_3P_2}$ 
    return  $QHWalk(P_1, P_3, A)$ ,  $P_3$ ,
    and  $QHWalk(P_3, P_1, B)$ 
```

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Chapter 2: Slide – 24

Comments on Quickhull

- *Quickhull* and *QHWalk* finds extreme points.
 - Points with lowest and highest x values are extreme.
 - Points farthest from any line are extreme.
 - P_3 is on the left of $\overline{P_1P_2}$ if $x_1y_2 + x_3y_1 + x_2y_3 - x_3y_2 - x_2y_1 - x_1y_3 > 0$. Farthest point on left maximizes this value.
- Best Case: If *QHWalk* always split points in thirds (A , B , and other), then $T(n) = 2T(n/3) + \Theta(n) \in \Theta(n)$.
- Worst Case: If split is bad, then $T(n) = T(n-1) + \Theta(n) \in \Theta(n^2)$.