Designing Algorithms with Divide-and-Conquer

Lecture 06.03 by Marina Barsky

Main algorithm design strategies

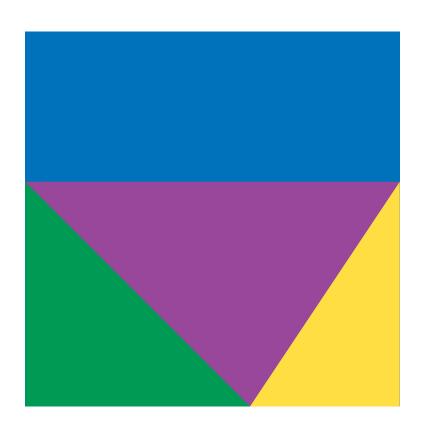
- ✓ Exhaustive Computation. Generate every possible candidate solution and select an optimal solution.
- ✓ Greedy. Create next candidate solution one step at a time by using some greedy choice.
- Divide and Conquer. Divide the problem into non-overlapping subproblems of the same type, solve each subproblem with the same algorithm, and combine sub-solutions into a solution to the entire problem.
- **Dynamic Programming.** Start with the smallest subproblem and combine optimal solutions to smaller subproblems into optimal solution for larger subproblems, until the optimal solution for the entire problem is constructed.
- *Iterative Improvement.* Perform multiple iterations of the algorithm, at each iteration moving closer to the optimal solution, until no further improvement is possible.

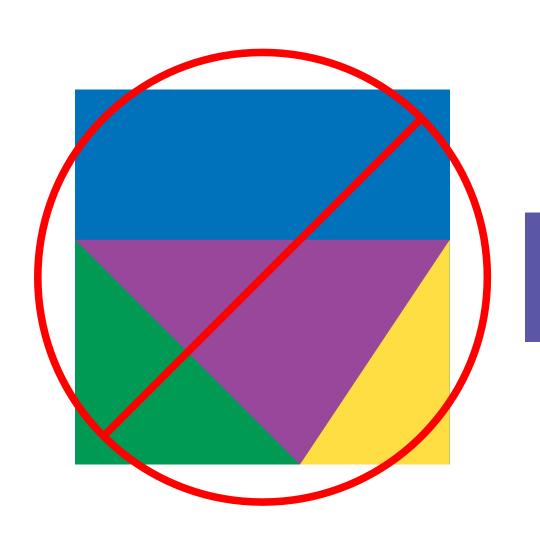
Big problem to be solved

Divide: Break into <u>non-overlapping</u> subproblems of <u>the same type</u>



Problem

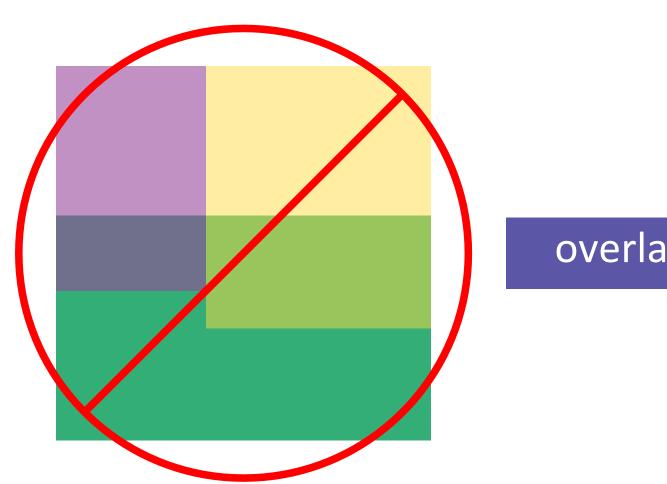




not the same type

Problem





overlapping

Divide-and-conquer steps

- 1. Break into *non-overlapping* subproblems of the same type
- 2. Solve subproblems
- 3. Combine results Most difficult!

Two examples:

- Counting inversions
- Closest pair

Counting inversions

Motivation

- Music site tries to match user song preferences with others.
- □ I rank *n* songs.
- Music site consults database to find people with similar tastes.

songs

	А	В	С	D	Е	F
me	1	2	3	4	5	6

How similar are me and you?

you 1	3	4	2	5	6
-------	---	---	---	---	---

Similarity of rankings

Similarity metric:
 number of *inversions* between two rankings.

□ My rank: 1,2,3,4,5,6

□ Your rank: 1,3,4,2,5,6

- for the same songs

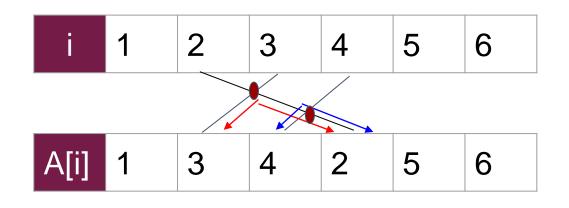
songs

	А	В	С	D	Е	F	
me	1	2	3	4	5	6	
you	1	3	4	2	5	6	

For a perfect match you should have ranked D at 4, but you ranked it at 2

Definition

An *inversion* is a pair (A[i], A[j]) of array elements such that index i<j and A[i] > A[j]



2 inversions in total: (3,2) and (4,2)

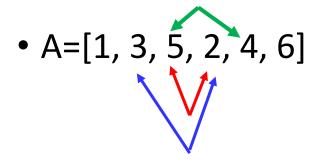
Problem: counting inversions

Input: an array A of length n with numbers 1,2,...n in some order

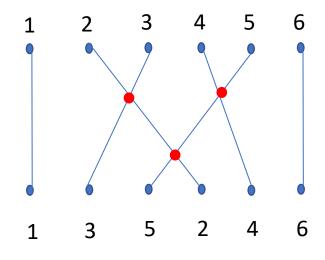
Output: number of *inversions*: number of pairs A[i],A[j] of array elements with i < j and A[i] > A[j]

- If A is sorted what is the number of inversions?
- What is the number of inversions if A is reversed?
- What is the number of inversions in A=[1,3,5,2,4,6]?

Example



• Inversions:



What is the largest-possible number of inversions that a 6-element array can have?

Brute-force algorithm for counting inversions

```
Algorithm count_naive (array A of n integers)
```

```
\begin{aligned} &\text{count:= 0} \\ &\text{for i from 1 to n-1:} \\ &\text{for j from i+1 to n:} \\ &\text{if A[j] < A[i]} \\ &\text{count:= count + 1} \end{aligned}
```

Complexity?

Can we do better?

But how can we do better if total number of inversions is O(n²)???

Idea 1: Divide + Conquer

After dividing array into 2 halves, n/2 each: For each (i,j) recursively determine if (A[i],A[j]) is an inversion

There are 3 possible cases (3 types of inversions):

```
Left inversions: if i, j \le n/2
Right inversions: if i, j > n/2
These two can be computed recursively
```

Split inversions: if i <= n/2 and j > n/2 But how to compute these?

 $\frac{n}{2}$ 2, 1

Developing recursive algorithm

```
count (array A of length n)
if n=1
       return 0
Else
       x = count (1<sup>st</sup> half of A, n/2)
       y = count (2^{nd} half of A, n/2)
                                            We do not know how
       z = count_split_inv(A, n)
                                                 to do that
return x+y+z
```

If we manage to do *CountSplitInv* in O(n) time then *Count* will run in O(n log n) - just like Merge Sort

Idea 2. What if we use *merge* from merge sort?

- □ Have recursive calls both count inversions and sort
- ☐ It turns out that the *merge* subroutine automatically recovers inversions!

Recursive Algorithm (in progress)

```
sort_count (array A of length n)
            if n=1
                     return (A,0)
             Else
                     (B, x) = sort count (1<sup>st</sup> half of A, n/2)
B- sorted 1st half of A
                     (C, y) = sort count (2<sup>nd</sup> half of A, n/2)
C- sorted 2<sup>nd</sup> half of A
                     (D, z) = count_split_inv(B,C)
                                                        We still do not know
             return (D, x+y+z)
                                                           how to do that
```

If we manage to do count_split_inv in O(n) time then sort_count will run in O(n log n) - just like Merge Sort

merge subroutine: from Merge Sort

```
D = will contain sorted array
```

 $B = 1^{st}$ sorted subarray [1:n/2]

 $C = 2^{nd}$ sorted subarray [n/2:n]

$$i = 1$$

$$j = 1$$



```
D

k
```

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

Stop and think

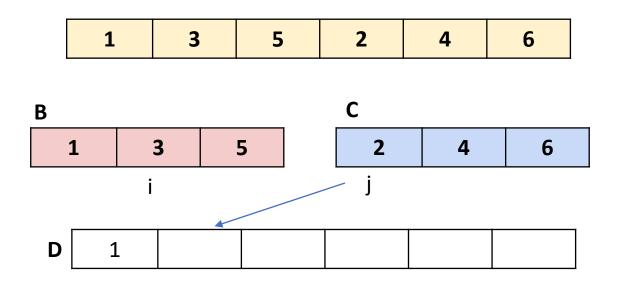
Suppose the input array A has no split inversions.

В

What is the relationship between the sorted subarrays B and C?

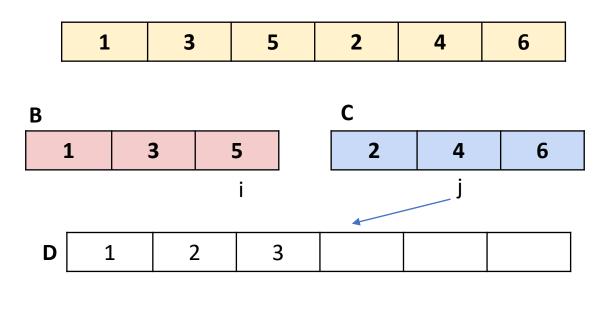
- A. B has the smallest element of A, C has the second--smallest, B has the third-smallest, and so on.
- A. All elements of B are less than all elements of C.
- A. There is not enough information to answer this question.

Sample merge



Discovered 2 inversions: (3,2) and (5,2)

Sample merge



Discovered inversion (5,4)

General claim

The split inversions involving an element y of the 2nd 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.

- \Box If x copied to output D before y, then x < y
 - => no inversions involving x and y
- \Box If y copied to output D before x, then y < x
 - \Rightarrow x and all elements after it are (split) inversions.

Recursive Algorithm (revised)

```
sort_count_inv (array A of length n)

if n=1
    return (A, 0)

Else

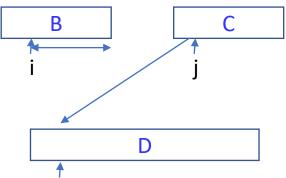
    (B, x) = sort_count_inv(1<sup>st</sup> half of A)
     (C, y) = sort_count_inv(2<sup>nd</sup> half of A)
     (D, z) = merge_count_split_inv(B,C)

return (D, x+y+z)
```

Split inversions are recovered during the merge of the sorted sub-arrays

Merge and count

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



• When element of 2^{nd} array C gets copied to output D, increment total by number of elements remaining in 1^{st} array B

Runtime of merge_count_split_inv: O(n) + O(n) =

O(n)

sort_count_inv runs in O(n log n) time
just like Merge Sort

Closest pair

Motivation

The closest-pair is a subroutine for:

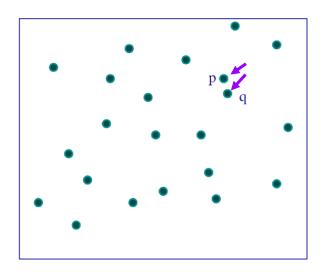
- Dynamic minimum spanning trees
- Straight skeletons and roof design
- Ray-intersection diagram
- Collision detection applications
- Hierarchical clustering
- Traveling salesman heuristics
- Greedy matching

• ...

"A pair of the closest points, the one lying on a robot and the other on its obstacles, yields the most important information for generation of obstacle-avoiding robot motions." ref

Closest Pair Problem

- **Input**: *n* points in *d* dimensions
- Output: two points p and q whose mutual distance is smallest



A naive algorithm takes $O(dn^2)$ time.

(Number of dimensions *d* can be assumed a constant for a given problem)

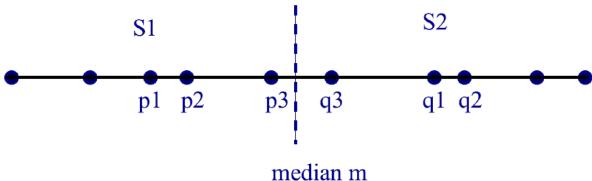
Can we do better?

Closest pair in one dimension

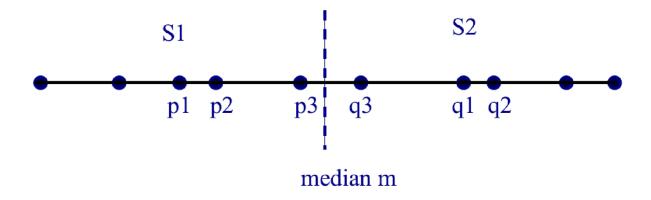
Can be solved in $O(n \log n)$ via sorting, and then linear scanning.

Let's develop a recursive solution to find the closest pair

- If the points are sorted by their coordinate:
- Divide the points set S into 2 sets S_1 , S_2 , by median x-coordinate m such that p < q for all $p \in S_1$ and $q \in S_2$
- Recursively compute closest pair (p₁,p₂) in S₁ and (q₁,q₂) in S₂

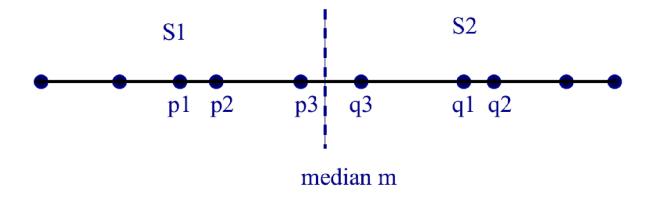


Closest pair in one dimension: combine step



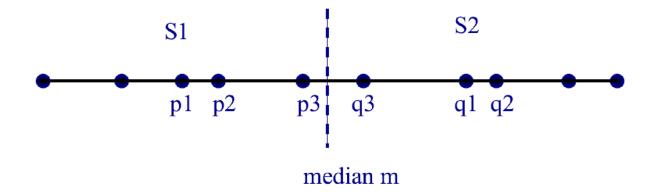
- Let δ be the smallest pairwise distance found in 2 partitions $\delta = \min(|p_2 p_1|, |q_2 q_1|)$
- The closest pair is either (p_1,p_2) , or (q_1,q_2) , or some (p_3,q_3) where $p_3 \in S_1$ and $q_3 \in S_2$
- Can we find (p_3,q_3) in a constant time?

Closest pair in 1 dimension



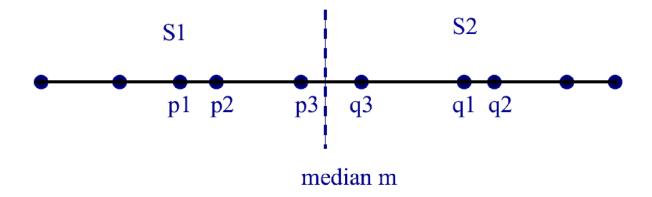
- The closest pair is either (p_1,p_2) , or (q_1,q_2) , or some (p_3,q_3) where $p_3 \in S_1$ and $q_3 \in S_2$
- Key observation: If m is the dividing coordinate, then both p_3 and q_3 have to be within δ of m

Closest pair in 1 dimension



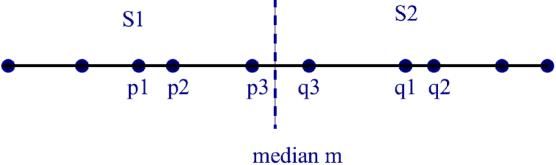
- Key observation: If m is the dividing coordinate, then both p_3 and q_3 have to be within δ of m
- How many such pairs exist?

Closest pair in 1 dimension



- Key observation: If m is the dividing coordinate, then both p_3 and q_3 have to be within δ of m
- How many points of S1 can lie in the interval $(m \delta, m]$?
- So we need to check one pair only constant time

Closest pair 1D: recursive algorithm



closest_pair (S – set of sorted points $p_i...p_n$, n>=2)

if
$$|S| = 2$$

return $\delta = |p_2 - p_1|$

Here we only compute the shortest distance, but it is easy to modify to return 2 points which produced this distance

```
Divide S into S_1 and S_2 at m = value[n/2] \delta_1 = closest\_pair(S_1) \delta_2 = closest\_pair(S_2) \delta_3 = closest\_pair\_across(S_1, S_2, min(\delta_1, \delta_2)) Constant time return \delta = min(\delta_1, \delta_2, \delta_3)
```

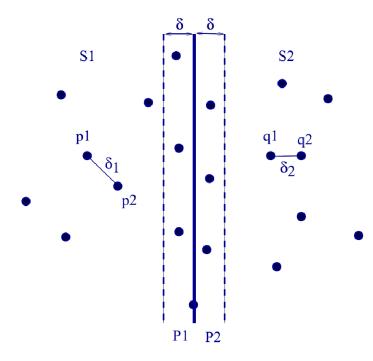
Closest pair in 1 dimension: time complexity

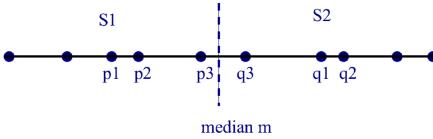
```
closest_pair (S – set of sorted points p_i...p_n, n>=2)
     if |S| = 2
          return \delta = |p_2 - p_1|
     Divide S into S_1 and S_2 at m = value[n/2]
     \delta_1 = \text{closest\_pair}(S_1)
     \delta_2 = closest_pair (S_2)
     \delta_3 = \text{closest\_pair\_across} (S_1, S_2, \min(\delta_1, \delta_2))
                                                                     Constant time
     return \delta = \min(\delta_1, \delta_2, \delta_3)
                  T(n) = 2T(n/2) + O(1)
                                                      We will learn why later
                  Which solves into O(n)
```

Together with sorting: O(n log n)

Closest pair in 2 dimensions

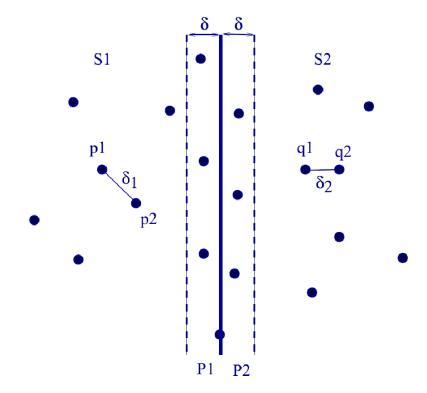
The previous algorithm does not generalize to higher dimensions, **or does it**?





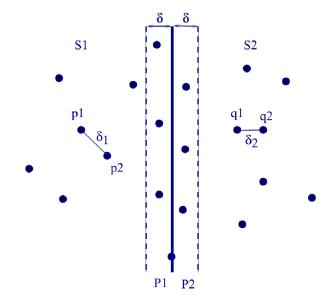
2D closest pair: divide

- Taking sorting as a free O(n log n) invariant, we sort all points in S by x coordinate
- Partition S into S₁, S₂ by vertical line l defined by median x-coordinate in S



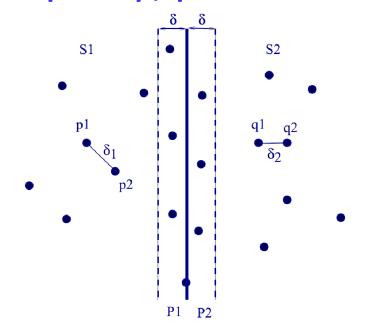
2D closest pair: conquer

- Recursively compute closest pair distances δ_1 and δ_2 in S_1 and S_2
- Set δ =min(δ_1 , δ_2)



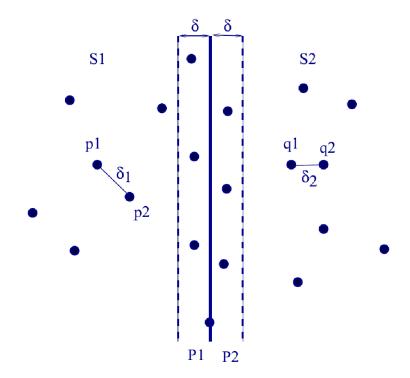
2D closest pair: combine

- Closest pair distances in S_1 and S_2 are δ_1 and δ_2 . δ =min(δ_1 , δ_2)
- Now need to combine: compute the closest pair across dividing line \boldsymbol{l}
- In each candidate pair (p,q), where $p \in S_1$ and $q \in S_2$, the only candidate points p, q must both lie within δ of ℓ .



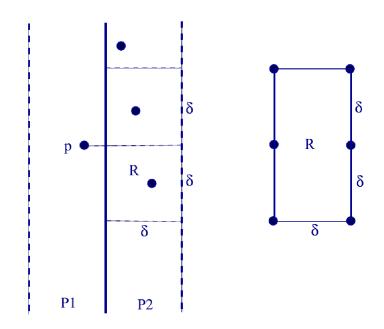
2D closest pair combine: complications

- At this point, complications arise, which were not present in 1D
- It is entirely possible that all n/2 points of S_1 (and S_2) lie within δ of l
- Naïvely, this would require n²/4 comparisons



Combining split points

- Consider a point $p \in S_1$.
- All points of S_2 within distance δ of p must lie in a $\delta x 2\delta$ rectangle R
- How many points can be inside
 R if we know that each pair is at least δ apart?
- In 2D, this number is at most 6!



So we only need to perform (n/2)*6 distance calculations during the combine step!

We do not have the $O(n \log n)$ algorithm yet. Why?

Combine in linear time

- In order to determine at most 6 potential mates of p, project p and all points of S₂ into y axis
- Pick out points whose projection is within δ of p: at most δ
- If we pre-sort S_1 and S_2 by the y coordinate
- Then we can do our check for all $p \in S_1$, by walking sorted lists S_{1v} and S_{2v} , in total O(n) time

The entire solution then runs in $O(n \log n)$