CS 580: Algorithm Design and Analysis

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Announcement: Homework 2 due on Tuesday, February 6th at 11:59PM

Recap: Minimum Weight Spanning Trees

Cut Property: Minimum weight edge crossing a cut must be in the MST (assume edge weights are distinct)

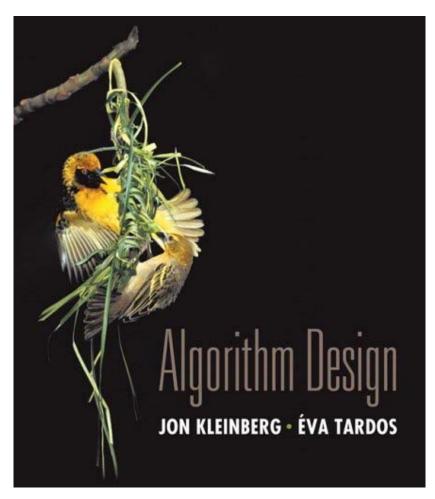
Cycle Property: Maximum weight edge in a cycle must not be in the MST (assuming edge weights are distinct)

Prim's Algorithm

- Repeatedly applies cut property to expand tree
- O(m log n) time with Binary Heap
- O(m+n log n) time with Fibonacci Héap

Prim's Algorithm

- · Consider edges in increasing order of weight
- For each edge we can either
 - Discard via Cycle Property, or
 - Add via Cut Property
- O(m log n) running time.

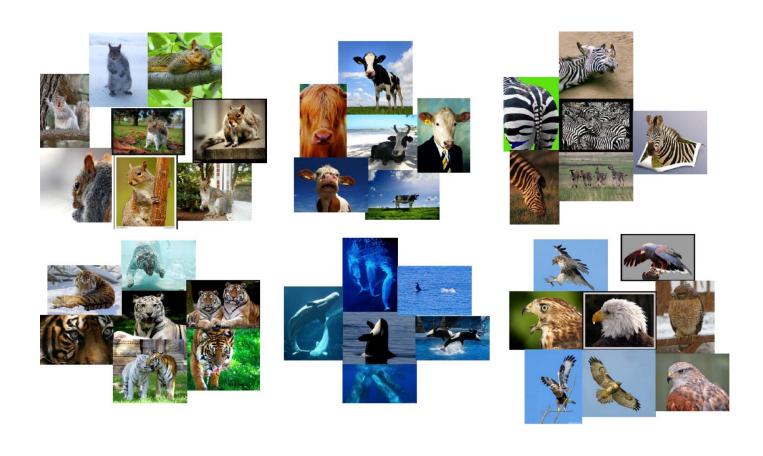


Greedy Algorithms



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4.7 Clustering



Clustering

Clustering. Given a set U of n objects labeled $p_1, ..., p_n$, classify into coherent groups.

photos, documents. micro-organisms

Distance function. Numeric value specifying "closeness" of two objects.

number of corresponding pixels whose intensities differ by some threshold

Fundamental problem. Divide into clusters so that points in different clusters are far apart.

- Routing in mobile ad hoc networks.
- Identify patterns in gene expression.
- Document categorization for web search.
- Similarity searching in medical image databases
- Skycat: cluster 109 sky objects into stars, quasars, galaxies.

Clustering of Maximum Spacing

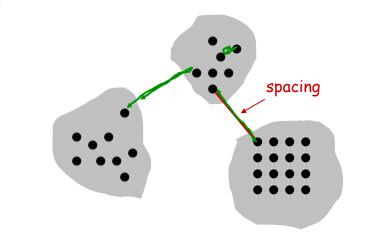
k-clustering. Divide objects into k non-empty groups.

Distance function. Assume it satisfies several natural properties.

- $d(p_i, p_j) = 0$ iff $p_i = p_j$ (identity of indiscernibles)
- $d(p_i, p_j) \ge 0$ (nonnegativity)
- $d(p_i, p_j) = d(p_j, p_i)$ (symmetry)

Spacing. Min distance between any pair of points in different clusters.

Clustering of maximum spacing. Given an integer k, find a k-clustering of maximum spacing.



k = 4

Greedy Clustering Algorithm

Single-link k-clustering algorithm.



- Form a graph on the vertex set U, corresponding to <u>n clusters</u>.
- Find the closest pair of objects such that each object is in a different cluster, and add an edge between them.
- Repeat n-k times until there are exactly k clusters.

Key observation. This procedure is precisely Kruskal's algorithm (except we stop when there are k connected components).

Remark. Equivalent to finding an MST and deleting the <u>k-1</u> most expensive edges.

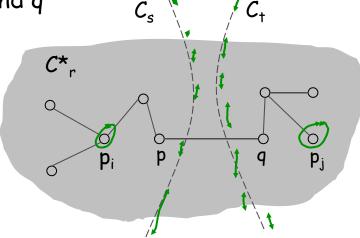
Graph 6 has O(n²) edges O(n² 109n)

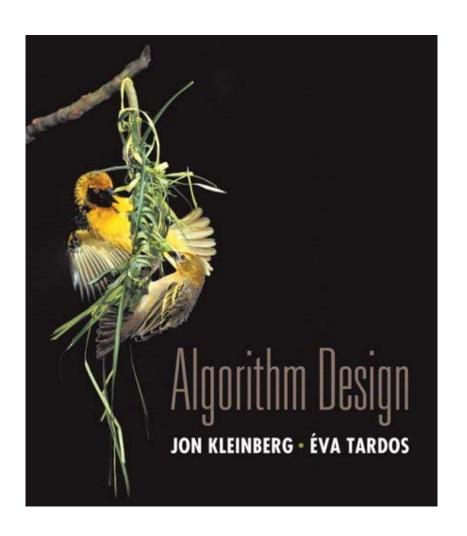
Greedy Clustering Algorithm: Analysis

Theorem. Let C^* denote the clustering $C^*_1, ..., C^*_k$ formed by deleting the k-1 most expensive edges of a MST. C^* is a k-clustering of max spacing.

Pf. Let C denote some other clustering $C_1, ..., C_k$.

- The spacing of C^* is the length d^* of the $(k-1)^{st}$ most expensive edge.
- Let p_i , p_j be in the same cluster in C^* , say C^*_r , but different clusters in C, say C_s and C_t .
- Some edge (p, q) on p_i - p_j path in C^*_r spans two different clusters in C.
- All edges on p_i - p_j path have length $\leq d^*$ since Kruskal chose them.
- Spacing of C is ≤ d* since p and q
 are in different clusters.





Divide and Conquer



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Divide-and-Conquer

Divide-and-conquer.

- Break up problem into several parts.
- Solve each part recursively.
- Combine solutions to sub-problems into overall solution.

Most common usage.

- Break up problem of size n into two equal parts of size $\frac{1}{2}$ n.
- Solve two parts recursively.
- Combine two solutions into overall solution in linear time.

Consequence.

- Brute force: n².
- Divide-and-conquer: n log n.

Veni, vidi, vici.

- Julius Caesar

5.1 Mergesort

Sorting

Sorting. Given n elements, rearrange in ascending order.

Applications.

- Sort a list of names.
- Organize an MP3 library.
- Display Google PageRank results.
- List RSS news items in reverse chronological order.
- Find the median.
- Find the closest pair.
- Binary search in a database.
- Identify statistical outliers.
- Find duplicates in a mailing list.
- Data compression.
- Computer graphics.
- Computational biology.
- Supply chain management.
- Book recommendations on Amazon.
- Load balancing on a parallel computer.

. . .

obvious applications

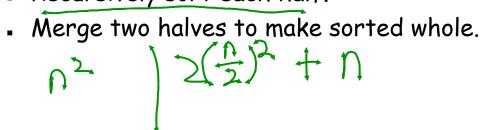
problems become easy once items are in sorted order

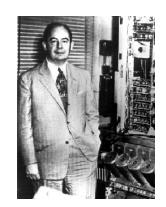
non-obvious applications

Mergesort

Mergesort.

- Divide array into two halves.
- Recursively sort each half.





Jon von Neumann (1945)

A L G O R I T H M S

A L G O R I T H M S divide
$$Q(1)$$

A G L O R H I M S T sort $2T(n/2)$

A G H I L M O R S T merge $Q(n)$

Merging

Merging. Combine two pre-sorted lists into a sorted whole.

How to merge efficiently?

- Linear number of comparisons.

 05demo-merge.ppt
- Use temporary array.



Challenge for the bored. In-place merge. [Kronrud, 1969]

using only a constant amount of extra storage

A Useful Recurrence Relation

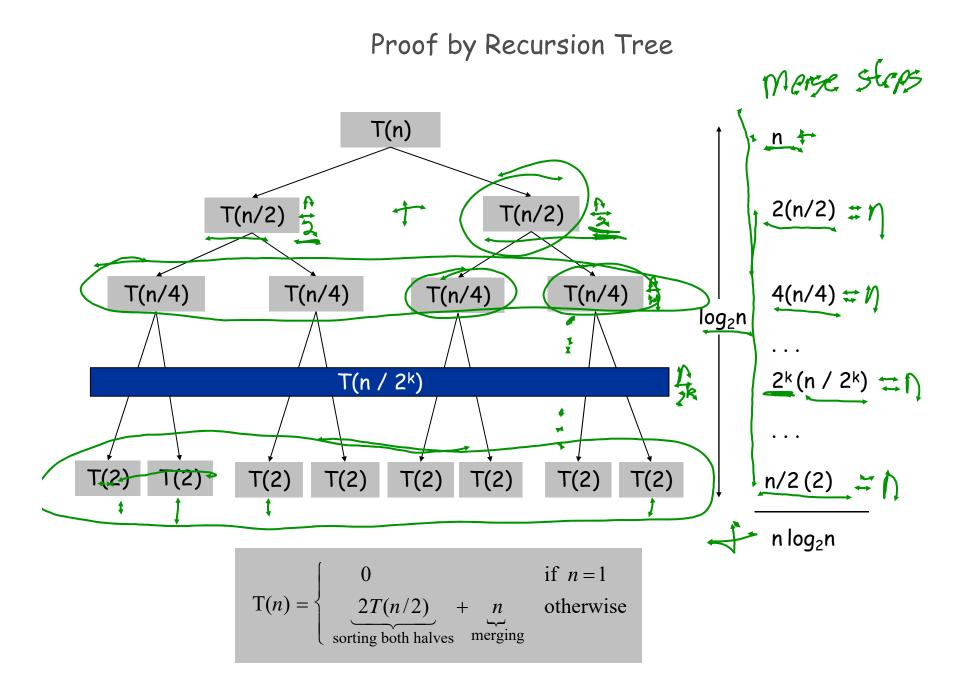
Def. T(n) = number of comparisons to mergesort an input of size n.

Mergesort recurrence.

$$T(n) \leq \begin{cases} 0 & \text{if } n = 1 \\ T(\lceil n/2 \rceil) + T(\lceil n/2 \rceil) + n & \text{otherwise} \end{cases}$$
solve left half solve right half

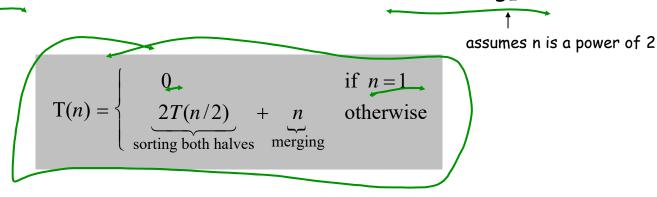
Solution.
$$T(n) = O(n \log_2 n)$$
.

Assorted proofs. We describe several ways to prove this recurrence. Initially we assume n is a power of 2 and replace \leq with =.



Proof by Telescoping

Claim. If T(n) satisfies this recurrence, then $T(n) = n \log_2 n$.



2T(n/2)

$$= \underbrace{\frac{T(n/2)}{n/2}} + 1$$

$$= \underbrace{\frac{T(n/4)}{n/4}} + \underbrace{1 + 1}$$

$$= \underbrace{\frac{T(n/n)}{n/n}} + \underbrace{1 + \dots + 1}_{\log_2 n}$$

$$= \log_2 n$$

T(n)

Proof by Induction

Claim. If T(n) satisfies this recurrence, then $T(n) = n \log_2 n$.

Guess This Recult ence, then
$$T(n)$$
 if $n = 1$

$$T(n) = \begin{cases} 0 & \text{if } n = 1 \\ 2T(n/2) + n & \text{otherwise} \\ \text{sorting both halves merging} \end{cases}$$

Pf. (by induction on n)

Inductive hypothesis:
$$T(n) = n \log_2 n$$
.

• Goal: show that
$$T(2n) = 2n \log_2 (2n)$$
.

$$T(2n) = 2T(n) + 2n$$

$$= 2n \log_2 n + 2n$$

$$= 2n(\log_2(2n) - 1) + 2n$$

$$= 2n \log_2(2n)$$

assumes n is a power of 2

T(n)=下(含)+丁(等)中的

Analysis of Mergesort Recurrence

Claim. If T(n) satisfies the following recurrence, then $T(n) \le n \lceil \lg n \rceil$.

$$T(n) \leq \begin{cases} 0 & \text{if } n = 1 \\ T(\lceil n/2 \rceil) + T(\lceil n/2 \rceil) + n & \text{otherwise} \end{cases}$$
solve left half solve right half merging

Pf. (by induction on n)

- Base case: n = 1.
- Define $n_1 = \lfloor n/2 \rfloor$, $n_2 = \lceil n/2 \rceil$.
- Induction step: assume true for 1, 2, ..., n-1.

$$T(n) \leq T(n_1) + T(n_2) + n$$

$$\leq n_1 \lceil \lg n_1 \rceil + n_2 \lceil \lg n_2 \rceil + n$$

$$\leq n_1 \lceil \lg n_2 \rceil + n_2 \lceil \lg n_2 \rceil + n$$

$$= n \lceil \lg n_2 \rceil + n$$

$$\leq n(\lceil \lg n \rceil - 1) + n$$

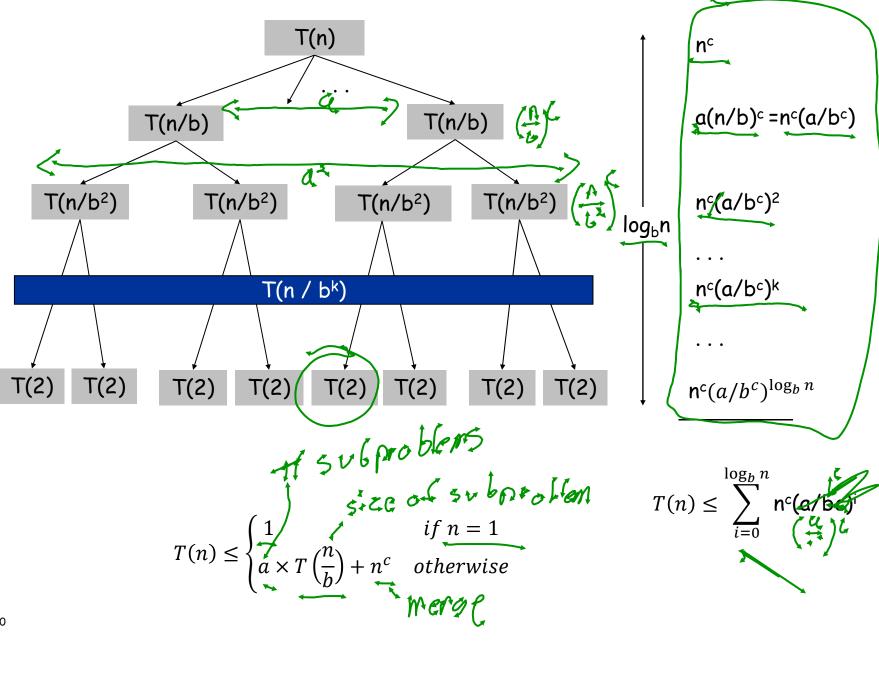
$$= n \lceil \lg n \rceil$$

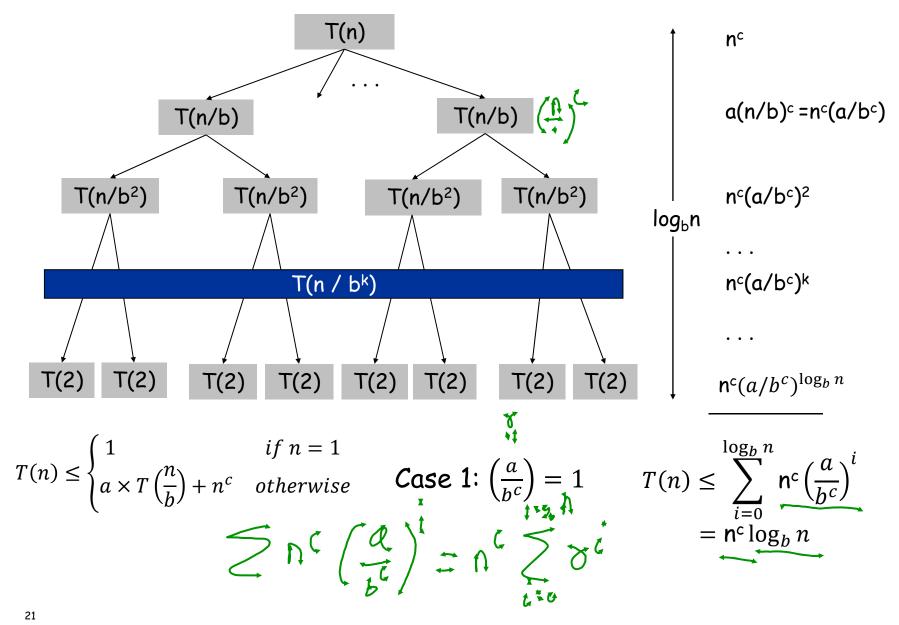
$$n_{2} = |n/2|$$

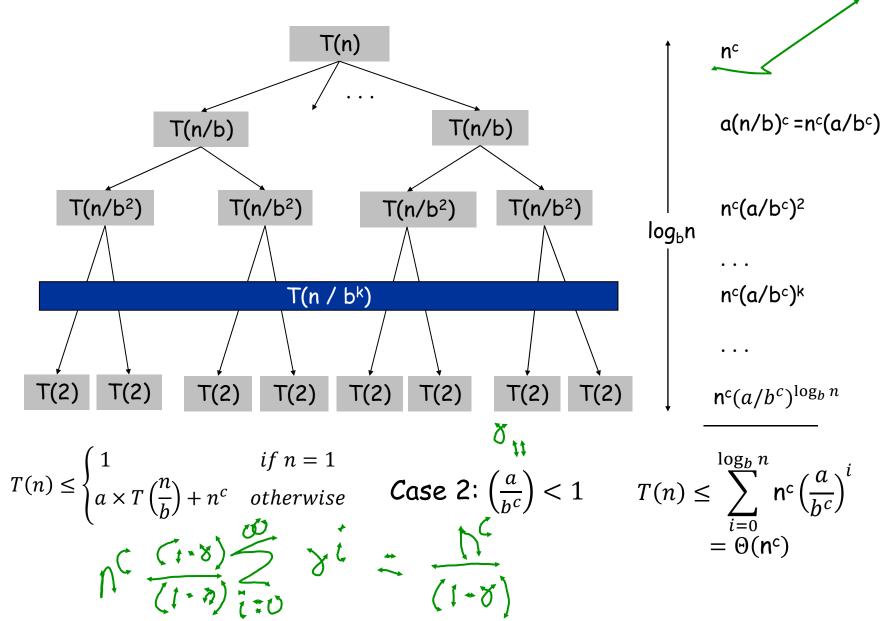
$$\leq \lceil 2^{\lceil \lg n \rceil} / 2 \rceil$$

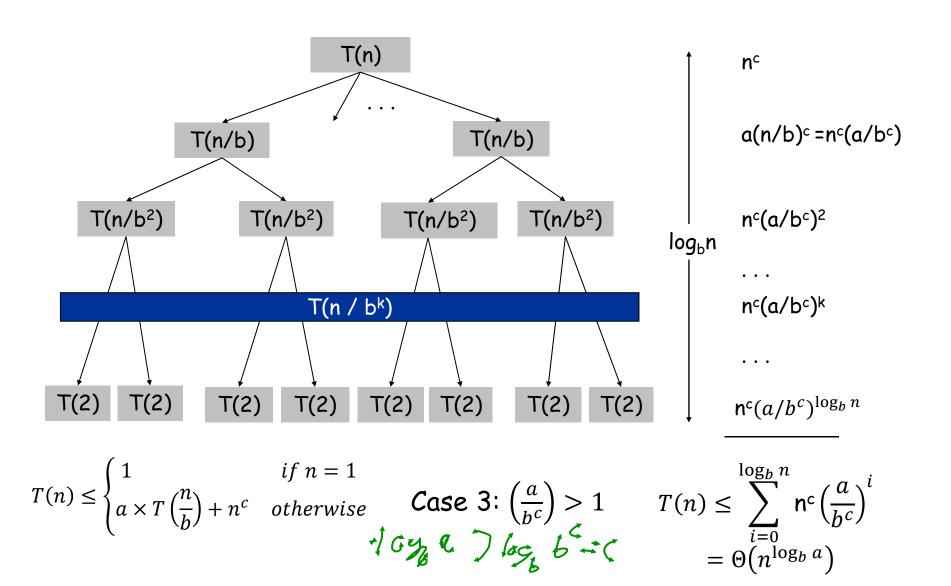
$$= 2^{\lceil \lg n \rceil} / 2$$

$$\Rightarrow \lg n_{2} \leq \lceil \lg n \rceil - 1$$









5.3 Counting Inversions

Counting Inversions

Music site tries to match your song preferences with others.

- You rank n songs.
- Music site consults database to find people with similar tastes.

Similarity metric: number of inversions between two rankings.

- My rank: 1, 2, ..., n.
- Your rank: $a_1, a_2, ..., a_n$.
- Songs i and j inverted if i < j, but $a_i > a_j$.

	Α	В	С	D	Ε
Me	1	2	3	4	5
You	1	3	4	2	5

Inversions 3-2, 4-2

Brute force: check all $\Theta(n^2)$ pairs i and j.

Applications

Applications.

- Voting theory.
- Collaborative filtering.
- Measuring the "sortedness" of an array.
- Sensitivity analysis of Google's ranking function.
- Rank aggregation for meta-searching on the Web.
- Nonparametric statistics (e.g., Kendall's Tau distance).

Divide-and-conquer.

1 5 4 8 10 2 6 9 12 11 3 7

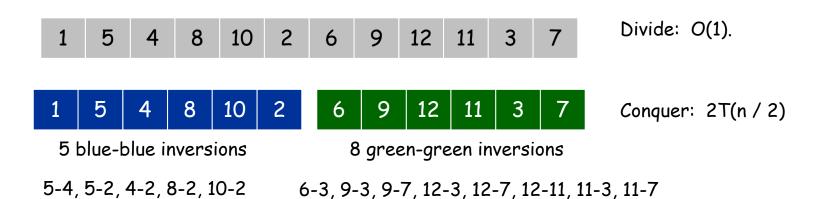
Divide-and-conquer.

Divide: separate list into two pieces.



Divide-and-conquer.

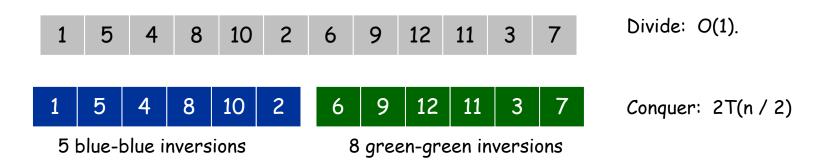
- Divide: separate list into two pieces.
- Conquer: recursively count inversions in each half.



Divide-and-conquer.

- Divide: separate list into two pieces.
- Conquer: recursively count inversions in each half.
- Combine: count inversions where a_i and a_j are in different halves, and return sum of three quantities.

Combine: ???



9 blue-green inversions 5-3, 4-3, 8-6, 8-3, 8-7, 10-6, 10-9, 10-3, 10-7

Total = 5 + 8 + 9 = 22.

Counting Inversions: Combine

Combine: count blue-green inversions



Assume each half is sorted.

play

- ${\color{blue} {\bf L}}$ Count inversions where a_i and a_j are in different halves.
- Merge two sorted halves into sorted whole.

to maintain sorted invariant



13 blue-green inversions: 6 + 3 + 2 + 2 + 0 + 0 Count: O(n)

Merge: O(n)

$$T(n) \le T(\lfloor n/2 \rfloor) + T(\lfloor n/2 \rfloor) + O(n) \Rightarrow T(n) = O(n \log n)$$

Counting Inversions: Implementation

Pre-condition. [Merge-and-Count] A and B are sorted. Post-condition. [Sort-and-Count] L is sorted.

```
Sort-and-Count(L) {
   if list L has one element
      return 0 and the list L

   Divide the list into two halves A and B
   (r<sub>A</sub>, A) ← Sort-and-Count(A)
   (r<sub>B</sub>, B) ← Sort-and-Count(B)
   (r , L) ← Merge-and-Count(A, B)

return r = r<sub>A</sub> + r<sub>B</sub> + r and the sorted list L
}
```

5.4 Closest Pair of Points

Closest Pair of Points

Closest pair. Given n points in the plane, find a pair with smallest Euclidean distance between them.

Fundamental geometric primitive.

- Graphics, computer vision, geographic information systems, molecular modeling, air traffic control.
- Special case of nearest neighbor, Euclidean MST, Voronoi.

fast closest pair inspired fast algorithms for these problems

Brute force. Check all pairs of points p and q with $\Theta(n^2)$ comparisons.

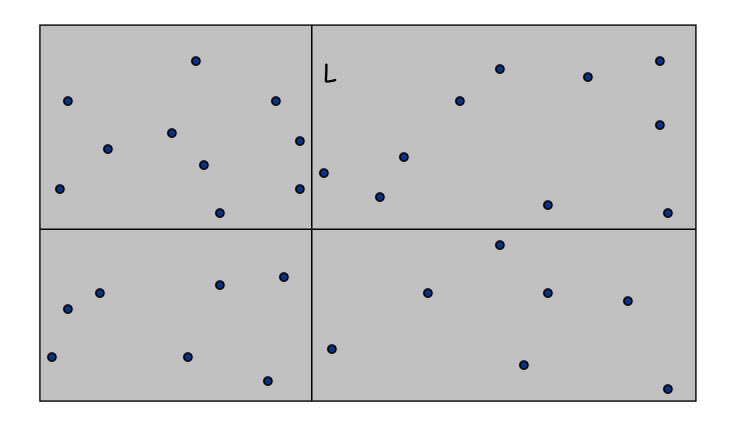
1-D version. O(n log n) easy if points are on a line.

Assumption. No two points have same x coordinate.

to make presentation cleaner

Closest Pair of Points: First Attempt

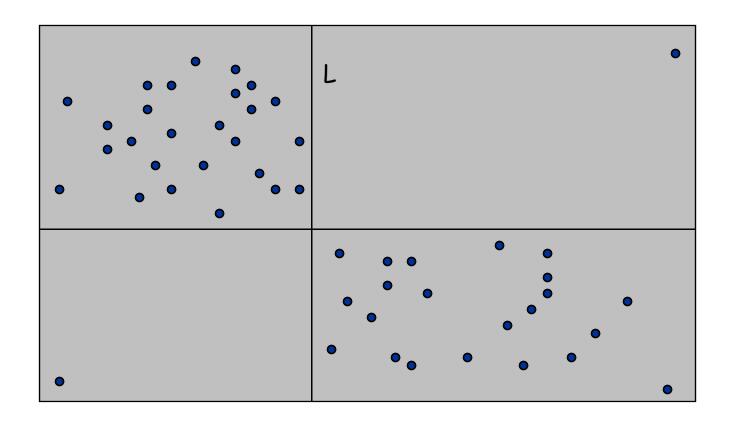
Divide. Sub-divide region into 4 quadrants.



Closest Pair of Points: First Attempt

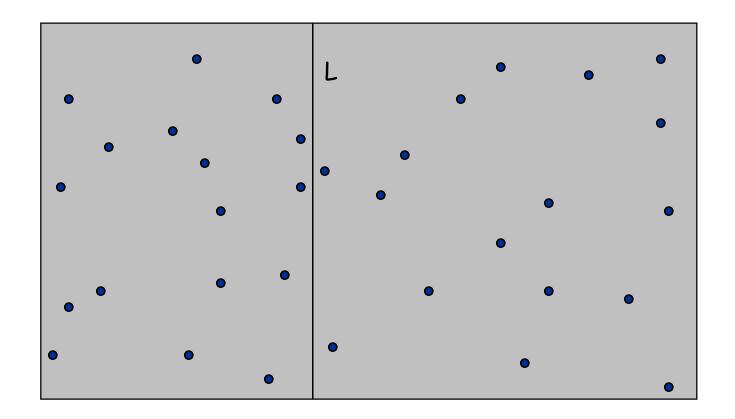
Divide. Sub-divide region into 4 quadrants.

Obstacle. Impossible to ensure n/4 points in each piece.



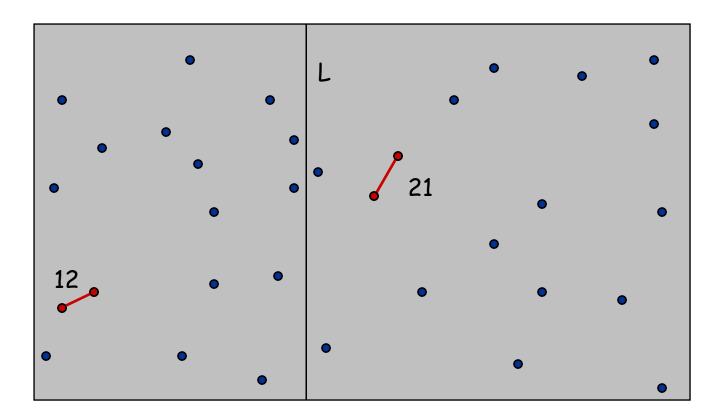
Algorithm.

• Divide: draw vertical line L so that roughly $\frac{1}{2}$ n points on each side.



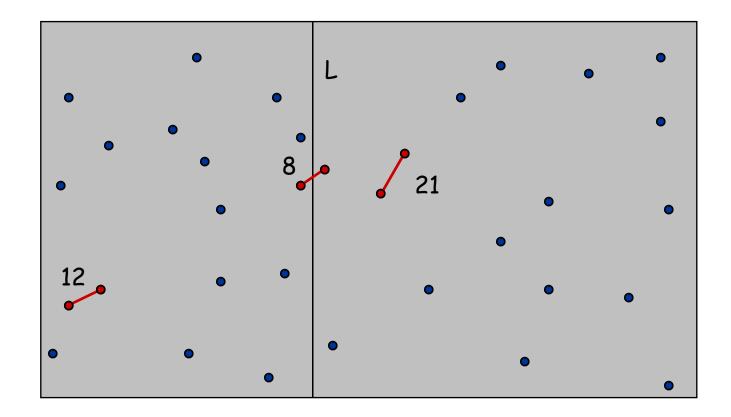
Algorithm.

- Divide: draw vertical line L so that roughly $\frac{1}{2}$ n points on each side.
- Conquer: find closest pair in each side recursively.

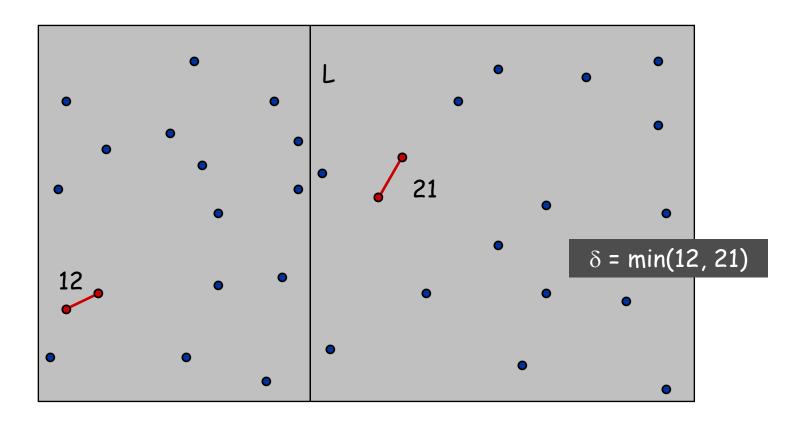


Algorithm.

- Divide: draw vertical line L so that roughly $\frac{1}{2}$ n points on each side.
- Conquer: find closest pair in each side recursively.
- Combine: find closest pair with one point in each side. \leftarrow seems like $\Theta(n^2)$
- Return best of 3 solutions.

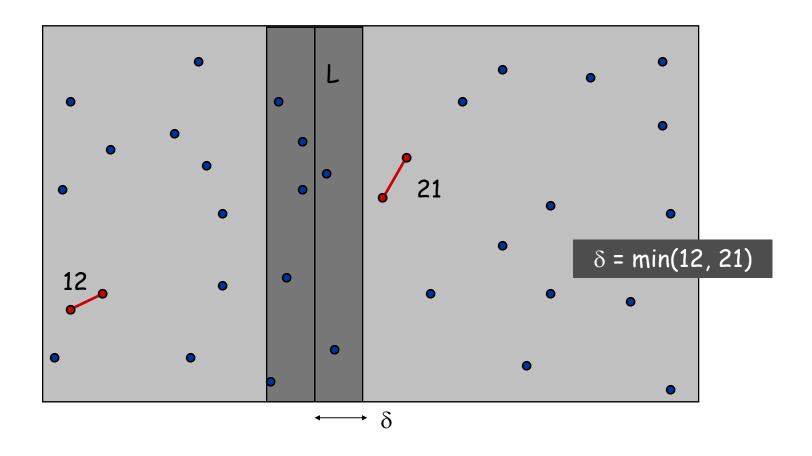


Find closest pair with one point in each side, assuming that distance $< \delta$.



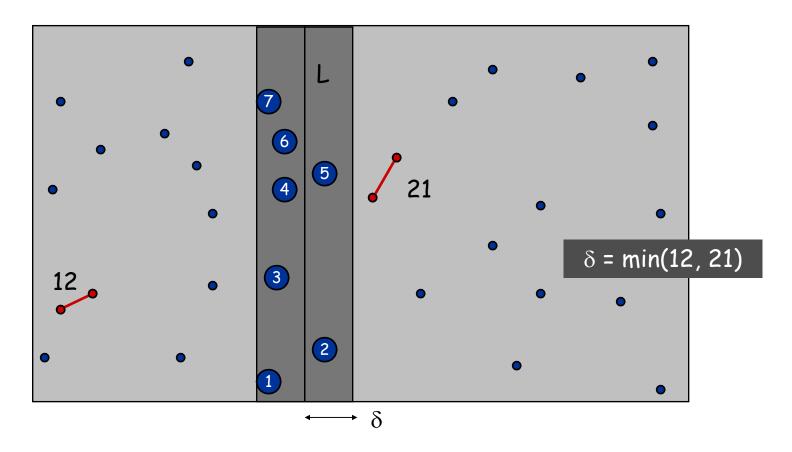
Find closest pair with one point in each side, assuming that distance $< \delta$.

 \blacksquare Observation: only need to consider points within δ of line L.



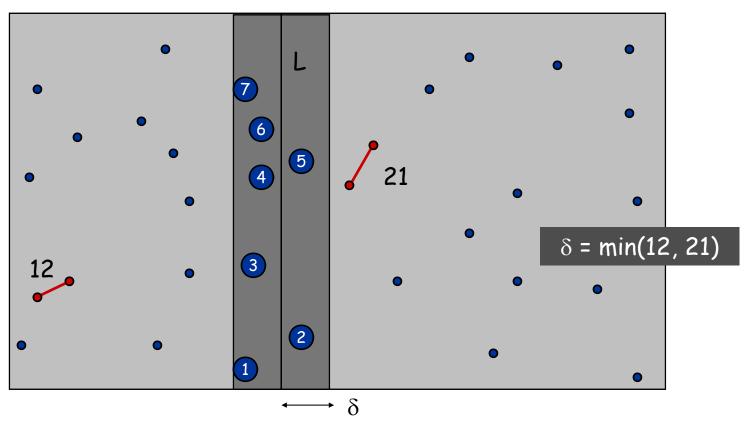
Find closest pair with one point in each side, assuming that distance $< \delta$.

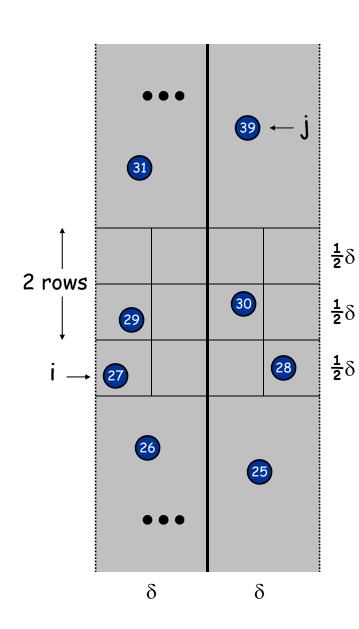
- Observation: only need to consider points within δ of line L.
- Sort points in 2δ -strip by their y coordinate.



Find closest pair with one point in each side, assuming that distance $< \delta$.

- \blacksquare Observation: only need to consider points within δ of line L.
- Sort points in 2δ -strip by their y coordinate.
- Only check distances of those within 11 positions in sorted list!





Def. Let s_i be the point in the 2δ -strip, with the i^{th} smallest y-coordinate.

Claim. If $|i-j| \ge 12$, then the distance between s_i and s_j is at least δ . Pf.

- No two points lie in same $\frac{1}{2}\delta$ -by- $\frac{1}{2}\delta$ box.
- Two points at least 2 rows apart have distance $\geq 2(\frac{1}{2}\delta)$. •

Fact. Still true if we replace 12 with 7.

Closest Pair Algorithm

```
Closest-Pair(p_1, ..., p_n) {
   Compute separation line L such that half the points
                                                                       O(n \log n)
   are on one side and half on the other side.
   \delta_1 = Closest-Pair(left half)
                                                                       2T(n / 2)
   \delta_2 = Closest-Pair(right half)
   \delta = \min(\delta_1, \delta_2)
   Delete all points further than \delta from separation line L
                                                                       O(n)
                                                                       O(n \log n)
   Sort remaining points by y-coordinate.
   Scan points in y-order and compare distance between
                                                                       O(n)
   each point and next 11 neighbors. If any of these
   distances is less than \delta, update \delta.
   return \delta.
```

Closest Pair of Points: Analysis

Running time.

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

Q. Can we achieve O(n log n)?

- A. Yes. Don't sort points in strip from scratch each time.
 - Each recursive returns two lists: all points sorted by y coordinate,
 and all points sorted by x coordinate.
 - Sort by merging two pre-sorted lists.

$$T(n) \le 2T(n/2) + O(n) \implies T(n) = O(n \log n)$$

MST Algorithms: Theory

Deterministic comparison based algorithms.

O(m log n) [Jarník, Prim, Dijkstra, Kruskal, Boruvka]

O(m log log n).[Cheriton-Tarjan 1976, Yao 1975]

• $O(m \beta(m, n))$. [Fredman-Tarjan 1987]

• $O(m \log \beta(m, n))$. [Gabow-Galil-Spencer-Tarjan 1986]

• $O(m \alpha (m, n))$. [Chazelle 2000]

Holy grail. O(m).

Notable.

O(m) randomized. [Karger-Klein-Tarjan 1995]

O(m) verification. [Dixon-Rauch-Tarjan 1992]

Euclidean.

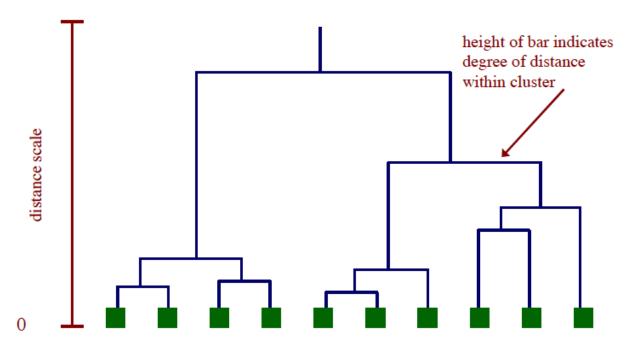
2-d: O(n log n). compute MST of edges in Delaunay

• k-d: $O(k n^2)$. dense Prim

Dendrogram

Dendrogram. Scientific visualization of hypothetical sequence of evolutionary events.

- Leaves = genes.
- Internal nodes = hypothetical ancestors.

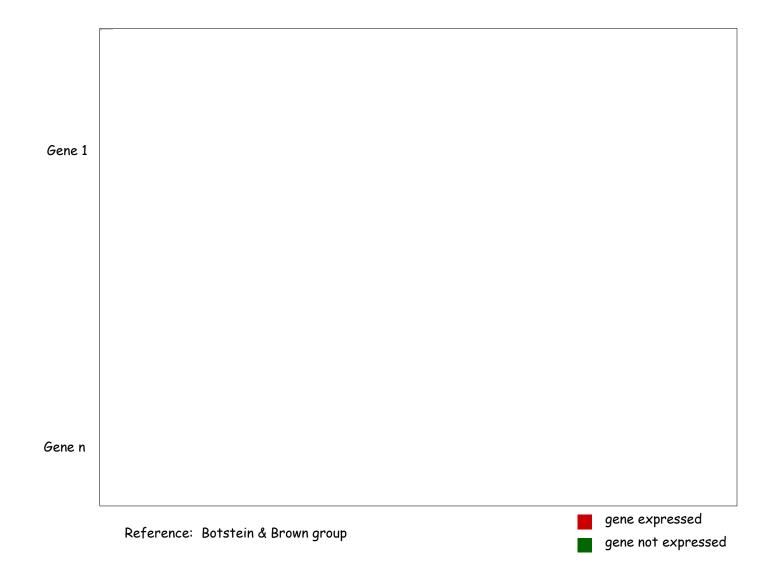


leaves represent instances (e.g. genes)

Reference: http://www.biostat.wisc.edu/bmi576/fall-2003/lecture13.pdf

Dendrogram of Cancers in Human

Tumors in similar tissues cluster together.



Extra Slides