## Lecture 10 Sublinear Time Algorithms (contd)

## Recap

- Sublinear time algorithms
  - > Deterministic + exact: binary search
  - Deterministic + inexact: estimating diameter in a metric space
  - Randomized + exact: searching in a sorted list
     Lower bound (thus optimality) using Yao's principle
  - Randomized + inexact:
    - $\,\circ\,$  Estimating average degree in a graph
    - $\odot$  Estimating size of maximal matching in a graph
    - Property testing
      - Testing linearity of a Boolean function

# Today

- Continue sublinear time property testing
  - > Testing if an array is sorted
  - > Testing if a graph is bipartite
- Some comments about sublinear space algorithms
- Begin streaming algorithms
  - > Find the missing element(s)
  - > Finding very frequent or very rare elements
  - > Counting the number of distinct elements

- Input: Array A of length n with O(1) access to A[i]
- Check: A[i] < A[i + 1] for every  $i \in \{1, ..., n 1\}$
- Definition of "at least  $\epsilon$ -far": You need to change at least  $\epsilon n$  entries to make it monotonic
  - Equivalently, there are at least entries that are not between their adjacent values.
- Goal: 1-sided algorithm with  $O\left(\frac{\log n}{\epsilon}\right)$  queries

#### • Proposal:

> Pick t random indices i, and return "no" even if  $x_i > x_{i+1}$  for even one of them.

• No!

> For 1 1 1 ... 1 0 0 0 ... 0 (n/2 each), we'll need  $t = \Omega(n)$ 

#### • Proposal:

> Pick t random pairs (i, j) with i < j, and return "no" if  $x_i > x_j$  for even one of them.

#### • No!

- 102132435465 ... (two interleaved sorted lists)
- >  $\frac{1}{2}$ -far (WHY?), but need  $t \ge \Omega(n)$  (by Birthday Paradox, we also must access  $\Omega(\sqrt{n})$  elements) (WHY?)

#### • Algorithm:

- > Choose  $2/\epsilon$  random indices *i*.
- > For each *i*, do a binary search for A[i].
- > Return "yes" if all binary searches succeed.
- Assume all elements are distinct w.l.o.g.
  - Can replace A[i] by (A[i], i) and use lexicographic comparison
- Important observation:
  - > "searchable" elements form an increasing subsequence! (WHY?)

#### • Algorithm:

- > Choose  $2/\epsilon$  random indices *i*.
- > For each *i*, do a binary search for A[i].
- > Return "yes" if all binary searches succeed.

#### • Thus:

- > If  $\alpha \cdot n$  elements searchable  $\Rightarrow$  array is at most  $(1 \alpha)$ -far from monotonic
- > If array is at least  $\epsilon$ -far from monotonic  $\Rightarrow$  at least  $\epsilon \cdot n$  elements must not be searchable

 $\,\circ\,$  Each iteration fails to detect violation w.p. at most  $1-\epsilon$ 

• All  $2/\epsilon$  iterations fail to detect w.p. at most  $(1-\epsilon)^{\frac{2}{\epsilon}} \leq 1/3$ 

# **Graph Property Testing**

- It's an active area of research by itself.
- Let G = (V, E) with n = |V| and m = |E|
- Input models:
  - Dense: Represented by adjacency matrix
    - Query if  $(i, j) \in E$  in O(1) time
    - $\circ \epsilon$ -far from satisfying P if  $\epsilon n^2$  matrix entries must be changed to satisfy P
    - $\circ$  Change required =  $\epsilon$ -fraction of the input

# **Graph Property Testing**

- It's an active area of research by itself.
- Let G = (V, E) with n = |V| and m = |E|
- Input models:
  - Sparse: Represented by adjacency lists
    - $\circ$  Query if (v, i) to get the  $i^{th}$  neighbor of v in O(1) time
    - $\circ$  We only use it for graphs with degrees bounded by d
    - $\epsilon$ -far from satisfying P if  $\epsilon(dn)$  matrix entries must be changed to satisfy P
    - $\circ$  Change required =  $\epsilon$ -fraction of the input
  - Generally, dense is *easier* than sparse

- Dense model:
  - > Upper bound:  $O(1/\epsilon^2)$  (independent of n) > Lower bound:  $\Omega(1/\epsilon^{1.5})$
- Sparse model (for constant d):
   > Upper bound: O (√n · poly (<sup>log n</sup>/<sub>ϵ</sub>))
   > Lower bound: Ω(√n)

- In the dense model:
- Algorithm [Goldreich, Goldwasser, Ron]
  - > Pick a random subset of vertices S,  $|S| = \Theta\left(\frac{\log\frac{1}{\epsilon}}{\epsilon^2}\right)$

> Output "bipartite" iff the induced subgraph is bipartite

- Analysis:
  - > Easy: If the graph is bipartite, algorithm always accepts.
  - > Claim: If the graph is  $\epsilon$ -far, it rejects w.p. at least 2/3
  - > Running time: trivially constant (i.e., independent of n)

- Q: Why doesn't this work for the sparse model?
  - > Take a line graph of n nodes. Throw  $\epsilon n$  additional edges.
  - > In the dense model, we don't care about this instance because it's not  $\epsilon$ -far (only  $\epsilon/n$ -far).
  - In the sparse model, we care about it, and the previous algorithm will not work.

- In the sparse model:
- Algorithm [Goldreich, Ron]
- Repeat  $O(1/\epsilon)$  times:
  - $\succ$  Pick a random vertex v
  - > Run OddCycle(v), and if it finds an odd cycle, REJECT.
- If no trial rejected, then ACCEPT.
- OddCycle:
  - > Performs  $poly(\log n/\epsilon)$  random walks from v, each of length  $poly(\log n/\epsilon)$ .
  - If a vertex is reachable by an even-length path and an odd-length prefix, an odd cycle is detected.

## Limitations of Sublinear Time

- The problems we saw are rather exceptions.
- For most problems, there is not much you can do in sublinear time.
- For instance, these problems require  $\Omega(n^2)$  time:
  - > Estimating  $\min_{i,j} d_{i,j}$  in a metric space d.
    - $\circ$  Contrast this with the sublinear algorithm we saw for estimating  $\max_{i,j} d_{i,j}$  (diameter)
  - > Estimating the cost of the minimum-cost matching
  - > Estimating the cost of k-median for  $k = \Omega(n)$

≻ ...

## Sublinear Space Algorithms

- An important topic in complexity theory
- Fundamental unsolved questions:
  - > Is NSPACE(S) = DSPACE(S) for  $S \ge \log n$ ?
  - > Is P = L? ( $L = DSPACE(\log n)$ , and we know  $L \subseteq P$ )
  - > What's the relation between P and polyL =  $DSPACE((\log n)^{O(1)})$ ?
    - We know  $P \neq polyL$ , but don't know if  $P \subset polyL$ ,  $polyL \subset P$ , or if neither is contained in the other.
- Savitch's theorem:
  - $\succ DSPACE(S) \subseteq NSPACE(S) \subseteq DSPACE(S^2)$

## USTCON vs STCON

- USTCON (resp. STCON) is the problem of checking if a given source node has a path to a given target node in an undirected (resp. directed) graph.
  - > USTCON  $\in$  RSPACE(log n) was shown in 1979 through a random-walk based algorithm
  - ➤ After much effort, Reingold [2008] finally showed that USTCON ∈ DSPACE(log n)
- Open questions:
  - > Is STCON in RSPACE( $\log n$ ), or maybe even in RSPACE( $\log n$ )?
  - > What about  $o(\log^2 n)$  instead of  $\log n$  space?
  - > Is RSPACE(S) = DSPACE(S)?

# Streaming Algorithms

- Input data comes as a stream  $a_1, \dots, a_m$ , where, say, each  $a_i \in \{1, \dots, n\}$ .
  - > The stream is typically too large to fit in the memory.
  - > We want to use only S(m, n) memory for sublinear S.
    - $\circ$  We can measure this in terms of the number of integers stored, or the number of actual bits stored (might be  $\log n$  times).
  - It is also desired that we do not take too much processing time per element of the strem.

 $\circ O(1)$  is idea, but  $O(\log(m + n))$  might be okay!

If we don't know m in advance, this can often act as an online algorithm.

# Streaming Algorithms

- Input data comes as a stream  $a_1, \dots, a_m$ , where, say, each  $a_i \in \{1, \dots, n\}$ .
  - > Most questions are about some statistic of the stream.
  - E.g., "how many distinct elements does it have?", or "count the #times the most frequent element appears"
  - > Once again, we will often approximate the answer.
  - Most algorithms process the stream in one pass, but sometimes you can achieve more if you can do two or more passes.

## Missing Element Problem

- Problem: Given a stream {a<sub>1</sub>, ..., a<sub>n-1</sub>}, where each element is a distinct integer from {1, ..., n}, find the unique missing element.
- An *n*-bit algorithm is obvious
  - > Keep a bit for each integer.
  - > At the end, spend O(n) time to search for the 0 bit.
- We can do O(log n) bits by maintaining the sum.
   ➤ Missing element = <sup>n(n+1)</sup>/<sub>2</sub> SUM
- Deterministic + exact.

#### Missing Elements Problem

- Problem: Given a stream {a<sub>1</sub>, ..., a<sub>n-k</sub>}, where each element is a distinct integer from {1, ..., n}, find all k missing elements.
- The previous algorithm can be generalized:
   > Instead of just computing the sum, compute power-sums.
   > {S<sub>j</sub>}<sub>1≤j≤k</sub> where S<sub>j</sub> = ∑<sup>n-k</sup><sub>i=1</sub>(a<sub>i</sub>)<sup>j</sup>
  - > At the end, we have k equations, and k unknowns.
  - > This uses  $O(k^2 \log n)$  space.
  - Computationally expensive to solve the equations
    - Using Newton's identities followed by finding roots of a polynomial

## Missing Elements Problem

- We can design much more efficient algorithms if we use randomization.
  - There is a streaming algorithm with space and time/item that is O(k log k log n).
  - > It can also be shown that  $\Omega\left(k\log\left(\frac{n}{k}\right)\right)$  space is necessary.

- Another classic problem is that of computing frequency moments.
  - > Let  $A = a_1, \dots, a_m$  be a data stream with  $a_i \in \{1, \dots, n\}$ .
  - > Let  $m_i$  denote the number of occurrences of value i.
  - > Then for  $k \ge 0$ , the  $k^{th}$  frequency moment is defined as  $F_k = \sum_{i \in [n]} (m_i)^k$
  - $> F_0 = #$  distinct elements
  - $> F_1 = m$
  - >  $F_2$  = Gini's homogeneity index

 $\circ$  The greater the value of  $F_2$ , the greater the homogeneity in A

• Goal: Given  $\epsilon, \delta$ , find  $F'_k$  s.t.

$$\Pr[|F_k - F'_k| > \epsilon F_k] \le \delta$$

• Seminal paper by Alon, Matias, Szegedy [AMS'99] > k = 0: For every c > 2,  $O(\log n)$  space algorithm s.t.  $\Pr[(1/c)F_0 \le F'_0 \le cF_0] \ge 1 - 2/c$ > k = 2:  $O((\log n + \log m)\log(1/\delta)/\epsilon) = \tilde{O}(1)$  space >  $k \ge 3$ :  $\tilde{O}(m^{1-1/k} \operatorname{poly}(1/\epsilon) \operatorname{polylog}(m, n, 1/\delta))$  space > k > 5: Lower bound of  $\Omega(m^{1-5/k})$ 

• Goal: Given  $\epsilon$ ,  $\delta$ , find  $F'_k$  s.t.

$$\Pr[|F_k - F'_k| > \epsilon F_k] \le \delta$$

• Seminal paper by Alon, Matias, Szegedy [AMS'99]

> k = 0: For every c > 2,  $O(\log n)$  space algorithm s.t.  $\Pr[(1/c)F_0 \le F'_0 \le cF_0] \ge 1 - 2/c$ 

- > Exactly counting  $F_0$  requires  $\Omega(n)$  space:
  - Once the stream is processed, the algorithm acts as a membership tester. On new element x, the count increases by 1 iff x was not part of the stream.
  - $\circ$  Algorithm must have enough memory to distinguish between all possible  $2^n$  states

• Goal: Given  $\epsilon$ ,  $\delta$ , find  $F'_k$  s.t.

$$\Pr[|F_k - F'_k| > \epsilon F_k] \le \delta$$

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> k = 0: For every c > 2,  $O(\log n)$  space algorithm s.t.  $\Pr[(1/c)F_0 \le F'_0 \le cF_0] \ge 1 - 2/c$ 

> State-of-the-art is "HyperLogLog Algorithm"

Uses hash functions

- Widely used, theoretically near-optimal, practically quite fast
- $\circ$  Uses  $O(\epsilon^{-2} \log \log n + \log n)$  space
- $\circ$  It can estimate > 10<sup>9</sup> distinct elements with 98% accuracy using only 1.5kB memory!

• Goal: Given  $\epsilon$ ,  $\delta$ , find  $F'_k$  s.t.

$$\Pr[|F_k - F'_k| > \epsilon F_k] \le \delta$$

- Seminal paper by Alon, Matias, Szegedy [AMS'99]
  - > k > 2: The  $\Omega(m^{1-5/k})$  bound was improved to  $\Omega(m^{1-2/k})$  by Bar Yossef et al.

 $\circ$  Their bound also works for real-valued k.

> Indyk and Woodruff [2005] gave an algorithm that works for real-valued k > 2 with a matching upper bound of  $\tilde{O}(m^{1-2/k})$ .

- The basic idea is to define a random variable Y whose expected value is close to  $F_k$  and variance is sufficiently small such that it can be calculated under the space constraint.
- We will present the AMS algorithm for computing  $F_k$ , and sketch the proof for  $k \ge 3$  as well as the improved proof for k = 2.

#### Algorithm:

> Let 
$$s_1 = 8\epsilon^{-2}k m^{1-1/k}$$
 and  $s_2 = 2\log^{1/\delta}$ .

> Let 
$$Y = median(Y_1, \dots, Y_{s_2})$$
, where

> 
$$Y_i = mean(X_{i,1}, ..., X_{i,s_1})$$
, where

- $\circ X_{i,j}$  are i.i.d. random variables that are calculated as follows:
- For each  $X_{i,j}$ , choose a random  $p \in [1, ..., m]$  in advance.
- $\circ$  When  $a_p$  arrives, note down this value.

○ In the remaining stream, maintain  $r = |\{q | q \ge p \text{ and } a_q = a_p\}|$ .

o 
$$X_{i,j} = m(r^k - (r-1)^k).$$

#### • Space:

- > For  $s_1 \cdot s_2$  variables X,  $\log n$  space to store  $a_p$ ,  $\log m$  space to store r.
- Note: This assumes we know *m*. But it can be estimated as the stream unfolds.

• We want to show:  $E[X] = F_k$ , and Var[X] is small.

• 
$$E[X] = E\left[m\left(r^k - (r-1)^k\right)\right]$$

- > The *m* different choices of  $p \in [m]$  have probability 1/m.
- ➤ Thus, E[X] is just the sum of r<sup>k</sup> (r 1)<sup>k</sup> across all choices of p.
- > For each distinct value *i*, there will be  $m_i$  terms:  $((m_i)^k - (m_i - 1)^k) + ((m_i - 1)^k - (m_i - 2)^k) + \dots + (1^k - 0^k) = (m_i)^k$

> Thus, the overall sum is  $F_k = \sum_i (m_i)^k$ .

• Thus, 
$$E[Y] = E[X] = F_k$$

- To show: Pr[|Y<sub>i</sub> − F<sub>k</sub>| > εF<sub>k</sub>] ≤ <sup>1</sup>/<sub>8</sub>
  ≻ Median over 2 log 1/δ many Y<sub>i</sub> will do the rest.
- Chebyshev's inequality:

$$\begin{split} & \operatorname{Pr} \left[ |Y_i - E[Y_i]| > \epsilon E[Y_i] \right] \leq \frac{\operatorname{Var}[Y]}{\epsilon^2 (E[Y])^2} \\ & \operatorname{Var}[Y_i] \leq \frac{\operatorname{Var}[X]}{s_1} \leq \frac{E[X^2]}{s_1}, \text{ and } E[Y] = E[X] = F_k. \\ & \text{Thus, probability bound is:} \\ & \frac{E[X^2]}{s_1 \epsilon^2 (F_k)^2} = \frac{E[X^2]}{8 \epsilon^{-2} \, k m^{1-1/k} \epsilon^2 (F_k)^2} \\ & \text{To show that this is at most } 1/8, \text{ we want to show:} \\ & E[X^2] \leq k m^{1-1/k} (F_k)^2 \\ & \text{Show that:} E[X^2] \leq k F_1 F_{2k-1}, \text{ and } F_1 F_{2k-1} \leq m^{1-\frac{1}{k}} (F_k)^2 \\ \end{split}$$

# Sketch of *F*<sup>2</sup> improvement

- They retain  $s_2 = 2 \log 1/\delta$ , but decrease  $s_1$  to just a constant  $16/\epsilon^2$ .
  - The idea is that X will not maintain a count for each value separately, but rather an aggregate.

$$> Z = \sum_{t=1}^{n} b_t m_t$$
, then  $X = Z^2$ 

- > The vector  $(b_1, ..., b_n) \in \{-1, 1\}^n$  is chosen at random as follows:
  - Let  $V = \{v_1, ..., v_h\}$  be  $O(n^2)$  "four-wise independent" vectors ○ Each  $v_p = (v_{p,1}, ..., v_{p,n}) \in \{-1,1\}^n$
  - Choose  $p \in \{1, ..., h\}$  at random, and set  $(b_1, ..., b_n) = v_p$ .

- Input: Stream  $A = a_1, \dots, a_m$ , where  $a_i \in [n]$
- Q: Is there a value *i* that appears more than *m*/2 times?

#### • Algorithm:

> Store candidate  $a^*$ , and a counter c (initially c = 0).

> For 
$$i = 1 ... m$$

$$\circ$$
 If  $c = 0$ : Set  $a^* = a_i$ , and  $c = 1$ .

#### $\circ$ Else:

• If 
$$a^* = a_i$$
,  $c \leftarrow c + 1$ 

• If 
$$a^* \neq a_i$$
,  $c \leftarrow c - 1$ 

- Space: Clearly  $O(\log m + \log n)$  bits
- Claim: If there exists a value v that appears more than m/2 times, then  $a^* = v$  at the end.
- Proof:
  - > Take an occurrence of v (say  $a_i$ ), and let's pair it up:
    - If it decreases the counter, pair up with the unique element  $a_j$  (j < i) that contributed the 1 we just decreased.
    - $\circ\,$  If it increases the counter:
      - If the added 1 is never taken back, QED!
      - If it is decreased by  $a_j$  (j > i), pair up with that.
  - > Because at least occurrence of v is not paired, the "never taken back" case happens at least once.

- Space: Clearly  $O(\log m + \log n)$  bits
- Claim: If there exists a value v that appears more than m/2 times, then  $a^* = v$  at the end.
- A simpler proof:
  - > At any step, let c' = c if  $a^* = v$ , and c' = -c otherwise.
  - > Every occurrence of v must increase c' by 1.
  - > Every occurrence of a value other than v either increases or decreases c' by 1.
  - > Majority  $\Rightarrow$  more increments than decrements in c'.
  - > Thus, a positive value at the end!

- Note 1: When a majority element does not exist, the algorithm doesn't necessarily find the mode.
- Note 2: If a majority element exists, it correctly finds that element. However, if there is no majority element, the algorithm does not detect that and still returns a value.
  - It can be trivially checked if the returned value is indeed a majority element if a second pass over the stream is allowed.
  - Surprisingly, we can prove that this cannot be done in 1pass. (Next lecture!)