CSC2420: Algorithm Design, Analysis & Theory

Lecture 9 (Sub-linear time / space (streaming) algorithms)

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1 Sub-linear time algorithms

In last lecture, we looked at some of the problems that can be solved (or approximated) using sub-linear time algorithms:

- Diameter of a metric space
- Searching in sorted linked-list
- Estimating the average degree of a graph (incomplete)

1.1 Estimating the average degree of a graph

Problem: Given a graph G = (V, E) and |V| = n, we want to estimate the average degree d of all vertices of G.

The $O(\sqrt{n}/\epsilon^{2.5})$ time algorithm presented in last lecture computes an estimate within a factor ~ 2 with sufficiently high probability. As the case with most sub-linear time algorithms, presentation of this algorithm is also simple but the analysis is not trivial.

Algorithm [1]

for $i = 1 \dots 8/\epsilon$ do Pick a set S_i of s =Compute d_{S_i} = average degree of vertices in $|S_i| = s = \sqrt{n}/(\epsilon^{2.5})$ end for Output min_i d_{S_i}

To prove the correctness of this algorithm we will prove the following claims:

Let d be the true average degree and S be one of these S_i

Claim 1: $Prob\left[d_S > (1+\epsilon)d\right] \le 1 - \frac{\epsilon}{2}$ (proved in last lecture)

Claim 2: $Prob\left[d_S < \frac{1}{2}(1-\epsilon)d\right] \leq \frac{\epsilon}{64}$

Theorem (Chernoff's bound): Let Z_1, \dots, Z_s independent "trials" of Z. Let $Z_i \in \{0, 1\}$ and $Z = \sum_{i=1}^s Z_i$ and $\mu = E[Z] = E[\sum_{i=1}^s Z_i]$. Then

$$Prob\left[\sum_{i=1}^{s} Z_i < (1-\epsilon)\mu\right] \le e^{-\mu\epsilon^2/4} \tag{1}$$

Proof of Claim 2: Let *H* be $\sqrt{\epsilon n}$ vertices of highest degree in the graph. Assume that the random selection of samples is done from *L* where,

$$L = V - H \tag{2}$$

By removing high degree vertices from random samples the probability of obtaining an average degree $d_S < \frac{1}{2}(1-\epsilon)d$ goes up. Now, the expected value of d_S when sampling from L is,

$$E[d_S] \ge \frac{1}{2} \left(\frac{d \cdot |V| - \binom{H}{2}}{|L|} \right) = \frac{1}{2} (d - \epsilon)$$
(3)

Therefore,

$$Prob\left[d_S < \frac{1}{2}(1-\epsilon)d\right] = Prob\left[d_S < (1-\epsilon)E[d_S]\right]$$
(4)

Let x_i be the degree of vertex choosen,

$$Prob\left[d_{S} < (1-\epsilon)E[d_{S}]\right] = Prob\left[\frac{\sum x_{i}}{d_{H}} \le (1-\epsilon)E\left[\frac{\sum x_{i}}{d_{H}}\right]\right]$$
(5)

$$\leq e^{-\epsilon^2 s.E[x_i]/d_H}$$
 (Chernoff's bound) (6)

If $s \ge \epsilon^{-2} \frac{d_H}{E[x_i]}$ we will be done; but we want our bound without knowing d_H . There are two cases:

• Case 1: $d_H \ge \frac{|H|}{\epsilon}$

$$E[x_i] = \sum_{v \in L} \frac{d(v)}{|L|}$$
(7)

$$\geq \frac{|H|d_H - |H|^2}{|L|} \tag{8}$$

$$\geq \frac{|H|(1-\epsilon)d_H}{|L|} \tag{9}$$

$$\implies \frac{d_H}{E[x_i]} \leq \frac{|V|}{|L|} \qquad (|V| > |L|) \tag{10}$$

$$= \frac{n}{\sqrt{\epsilon n}} \tag{11}$$

Thus,

$$s \ge \epsilon^{-2} \epsilon^{-1/2} \sqrt{n} \tag{12}$$

• Case 2: $d_H < \frac{|H|}{\epsilon}$

$$\epsilon^{-2} \frac{d_H}{E[x_i]} \leq \frac{\epsilon^{-2}}{\epsilon} |H| \tag{13}$$

$$\leq \epsilon^{-3}\sqrt{\epsilon n}$$
 (14)

$$= \epsilon^{-2.5} \sqrt{n} \tag{15}$$

1.2 Property testing

Definition: "Given the ability to perform (local) queries concerning a particular object (e.g., a function, or a graph), the task is to determine whether the object has a predetermined (global) property (e.g., linearity or bipartiteness), or is far from having the property. The task should be performed by inspecting only a small (possibly randomly selected) part of the whole object, where a small probability of failure is allowed [2]."

Property testing grew out of program testing. In program testing the goal is to check whether the program computes a specified function. One can test whether a program satisfies a certain property before checking whether the program computes a specified function. This paradigm has been followed both in theory of program testing and in practice through debugging. Different types of problems are studied in the context of property testing: graph properties, algebraic properties of functions, string properties, clustering, properties of boolean functions and more [2].

1.2.1 Testing an array for monotonicity

Goal: Given an array of length n with distinct values, test whether it is monotone or ϵ -far away from monotone [3].

Algorithm

for $O(1/\epsilon)$ trials do
Randomly choose j where $1 \leq j \leq n$ and let $v_j = A[j]$
Perform a binary search to determine whether v_j is in A
if not found report A is not monotone
end for
report A is monotone

The complexity of algorithm is $O((1/\epsilon) \log n)$.

Let S be a set of successful searches.

Lemma: S is a monotone sub-sequence.

Proof: Given, i < j and $i, j \in S$, at some point the binary search for v_i must diverge from the binary search for v_j . Let k be that point then at k,

$$A(i) \leq A(k) \tag{16}$$

$$A(k) \leq A(j) \tag{17}$$

This implies that,

$$A(i) \le A(j) \tag{18}$$

Therefore, S is an increasing sub-sequence.

Claim: If A is monotone the algorithm reports it with sufficiently high probability and if A is ϵ -far from monotone the algorithm rejects with sufficiently high probability.

Proof: If A is monotone then all the binary searches will succeed and the algorithm always reports that A is monotone. Suppose A is ϵ -far away from monotone. This implies $|S| < (1 - \epsilon)n$ since S is a monotone sub-sequence and if $|S| \ge (1 - \epsilon)n$, then changing at most $n\epsilon$ coordinates $j \notin S$ would make the input monotone. That would make A ϵ -close to monotone. Hence the probability with which the algorithm reports A as monotone is,

=

$$Prob[ALG \text{ accepts}] < (1-\epsilon)^{1/\epsilon}$$
 (19)

$$= (1 - \frac{1}{\delta})^{\delta} , \delta = \frac{1}{\epsilon}$$
 (20)

$$\Rightarrow e^{-1}$$
 (21)

Thus if A is ϵ -far from monotone, the algorithm rejects with probability $1 - e^{-1}$.

1.2.2 Testing for element distinctness

Goal: Given unsorted array A of length n, test if all A(i) are distinct.

Algorithm

Randomly choose set X with \sqrt{n}/ϵ elements if X has a repeated element **report** failure **else** report success

The complexity of algorithm is $O((\sqrt{n}/\epsilon) \log n)$. If we use hashing the we can get rid of the $\log n$ factor. Proof of correctness is based on "birthday paradox".

1.2.3 Graph property testing

There are several models for testing properties of graphs. Let G = (V, E), n = |V|, and m = |E|,

- 1. Dense model: These graphs are represented by its $n \times n$ adjacency matrix. We say that a graph is ϵ -far from having a property in this model if more than an ϵ -fraction (ϵn^2) of its adjacency matrix need to be modified in order to obtain the property.
- 2. Sparse/bounded degree model: In this model there is an upper bound d (some constant) on the degree of vertices. The graph is represented by an $n \times d$ matrix. We say that a graph is ϵ -far from having a property in this model if more than an ϵ -fraction (ϵdn) of its adjacency matrix should be modified in order to obtain the property.

Testing K-colorability

Given a dense graph G = (V, E) test,

- G is k-colorable.
- G is ϵ -far from k-colorable, i.e. need to remove at least ϵn^2 edges to make it k-colorable.

For k = 2, the problem reduces to testing the bipartiteness of graph. Given a dense graph G = (V, E), determine with high probability if it is bipartite or ϵ -far from it.

Algorithm

Randomly selects $\Theta\left(\frac{\log(1/\epsilon)}{\epsilon^2}\right)$ vertices Accept **if** the sub-graph induced on them is bipartite

In dense model \exists constant time algorithm (with the constant $C_{k,\epsilon}$ depending on k and ϵ) such that the algorithm tests for k-colorability (i.e. whether the graph is bipartite or ϵ far from being bipartite).

In sparse model, for constant degree d and ϵ , testing bipartiteness requires $\Omega(\sqrt{n})$ queries of the "incidence vector".

Algorithm

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for \Theta\left(\frac{1}{\epsilon}\right) times
Select a vertex v \in V
if \exists odd length cycle of v, report graph is not bipartite
end for
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In bipartite graph all cycles are of even length.

2 Sub-linear space (streaming) algorithms

In streaming model input is a sequence of data $A(1), A(2), \ldots, A(m), \ldots$ which is too large to be stored in memory. The space available is less than linear space << m. Common types of problems analyzed by streaming algorithms are:

1. Computing frequency (moments) statistics [4]: Let $A = (a_1, a_2, ..., a_n)$ be a sequence of elements, where each a_i is a member of $N = \{1, 2, 3, ..., n\}$. Let m_i denote the number of occurrences of a_i in the sequence A, then,

$$F_k = \sum_{i=1}^n m_i^k \tag{22}$$

 F_k are called the frequency moments of A and provide useful statistics on the sequence. F_0 is the number of distinct elements appearing in the sequence, F_1 is the length of the sequence, and F_2 is the repeat rate or Gini's index of homogeneity needed in order to compute the surprise index of the sequence. Surprise index for event (i),

$$S_i = \frac{\sum_j P_j^2}{P_j} \tag{23}$$

where, $P_j = \frac{m_j}{m}$. Alon, Matias, and Szegedy [4] showed that for every k > 0, F_k can be approximated randomly using at most $O(n^{1-1/k} \log n)$ memory bits.

- 2. Finding k "heavy hitters": Heavy hitters are the items occurring with frequency above a given threshold. E.g. those $a_i : a_i$ occurs at least m/k times in the stream.
- 3. Finding rare or unique values.

References

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