

CSC2420: Algorithm Design, Analysis and Theory

Spring 2019

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Week 11

Announcements:

- Geoff Hinton receives Turing Award.
- I think everyone now has a topic for the project. But please ask if there are further questions.
- I posted the remaining questions for the final assignment problems on Sunday, March 24. The due date is April 4.
- I posted the Data Stream Algorithm Lecture Notes by Amit Chakrabarti. He has more recent 2011 and 2015 versions of his data stream analysis course on his Dartmouth College web site.

Today's agenda:

- Continue sublinear time algorithms after a review of last week brief introduction.
- Streaming algorithms

Review of some results from last week

Last week we quickly presented three sublinear time algorithms. Before considering some further sublinear algorithms, here is a quick review.

- A deterministic algorithm for approximating the diameter of points in a finite metric space. **Note:** we said that this was an exception as almost all sublinear time algorithms require randomization and, moreover, for explicitly defined metric spaces (e.g., such as Euclidean spaces given by the point locations) this is no longer a sublinear time algorithm.
- Searching for an element in an anchored list. The algorithm we presented was just an adaption of the Chazelle, Liu and Magen algorithm for searching in a doubly linked list. There is also a matching lower bound that we will sketch today.
- Estimating the average degree in a graph. Today we will also fill in some of the analysis. There are some substantial technical details so best to read the paper (SICOMP 2006). We will assume connected graphs so that $1 \leq \text{mindegree} \leq \text{averagedegree}$.

What is the difference between the average degree problem and estimating the average of n numbers?

To estimate the average of a set of numbers $\{x_i\}$ (with $1 \leq x_i \leq n - 1$) requires $\Omega(n)$ time. This is a needle in a haystack problem if all but one of the $x_i = 1$ and the other one is n .

The average degree in a graph seems pretty much like the average of a set of numbers. **What is different?** Not much if the graph consists of a single edge.

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To estimate the average of a set of numbers $\{x_i\}$ (with $1 \leq x_i \leq n-1$) requires $\Omega(n)$ time. This is a needle in a haystack problem if all but one of the $x_i = 1$ and the other one is n .

The average degree in a graph seems pretty much like the average of a set of numbers. **What is different?** Not much if the graph consists of a single edge.

For a connected graph, the difference is intuitively that while we may not sample a high degree vertex, we are likely to find their neighbours and this will wind up accounting for the high degree edges.

A more precise argument would use the following theorem:

Erdos-Gallai

The sequence $d_1 \geq d_2 \geq \dots \geq d_n \geq 1$ is a graph degree sequence if and only if $\sum_{i=1}^n d_i$ is even and $\sum_{i=1}^k d_i \leq k(k-1) + \sum_{i=k+1}^n d_i$.

Estimating average degree in a graph

Here is where we ended last lecture.

- Given a graph $G = (V, E)$ with $|V| = n$, we want to estimate the average degree d of the vertices.
- We want to construct an algorithm that approximates the average degree within a factor less than $(2 + \epsilon)$ with probability at least $3/4$ in time $O(\frac{\sqrt{n}}{\text{poly}(\epsilon)})$. More We will assume that we can access the degree d_i of any vertex v_i in one step.
- Like a number of results in this area, the algorithm is simple but the analysis requires some care.

The Feige algorithm

Sample $8/\epsilon$ random subsets S_i of V each of size (say) $\frac{\sqrt{n}}{\epsilon^{2.5}}$

Compute the average degree a_i of nodes in each S_i .

The output is the minimum of these $\{a_i\}$.

The precise statement of Feige's approximation

We will just sketch a slightly weaker result but here is precisely the statement in the Feige 2006 SICOMP paper.

For any d_0 (the minimum degree in the graph) and for $\rho = 2 + \epsilon$, the Feige algorithm computes an estimation within a factor of ρ with high probability (e.g., any constant) and uses $O(\frac{1}{\epsilon} \sqrt{n/d_0})$ degree queries.

The analysis of the approximation

Since we are sampling subsets to estimate the average degree, we might have estimates that are too low or too high. But we will show that with high probability these estimates will not be too bad.

The proof will follow from the following two lemmas concerning the average degree a_i of a subset S_i .

- ❶ Lemma 1: $Prob[a_i < \frac{1}{2}(1 - \epsilon)\bar{d}] \leq \frac{\epsilon}{64}$
- ❷ Lemma 2: $Prob[a_i > (1 + \epsilon)\bar{d}] \leq 1 - \frac{\epsilon}{2}$

How the proof follows from the lemmas

The probability bound in Lemma 2 is amplified to any constant (say $1/8$) by repeated trials; i.e., the $8/\epsilon$ independent trials of random S_i . That is,

$$\begin{aligned} \text{Prob}[\text{Alg} > (1 + \epsilon)\bar{d}] &= \text{Prob}[a_i > (1 + \epsilon)\bar{d}] \quad \forall i \\ &< \left(1 - \frac{\epsilon}{2}\right)^{8/\epsilon} \leq e^{-4} \leq 1/8. \end{aligned}$$

For Lemma 1, we fall outside the desired bound if any of the repeated trials gives a very small estimate of the average degree but by the union bound this is no worse than the sum of the probabilities for each trial. That is, $\text{Prob}[\text{ALG} < \frac{1}{2}(1 - \epsilon)\bar{d}] \leq \sum_{i=1}^{\epsilon/8} (\epsilon/64) = 1/8$.

Sketch of the lemmas in Feige's average degree algorithm

Lemma 2 is relatively easy. Consider any S_i . Letting X_j be the degree of vertex v_j we have $\mathbb{E}[X_j] = \bar{d}$, and

$$\mathbb{E}[a_i] = \mathbb{E}\left[\frac{1}{|S_i|} \left(\sum_{j:j \in S_i} X_j\right)\right] = \frac{1}{|S_i|} \sum_{j:j \in S_i} \mathbb{E}[X_j] = \bar{d}$$

We can then use Markov's inequality to obtain Lemma 2.

The proof of Lemma 1 is more involved. We cannot simply amplify an error bound for a random subset since each S_i trial gives another chance of finding a low estimate. Instead we will have to use a Chernoff bound for the probability that a sum of independent random variables deviates from its mean. The following is sufficient:

A Chernoff bound

Let Z_1, \dots, Z_s be a sequence of independent r.v.s with $Z_j \in [0, 1]$ and let $\mu = \mathbb{E}[\sum_j Z_j]$. Then $\text{Prob}[\sum_j Z_j \leq (1 - \epsilon)s\mu] \leq e^{-\epsilon^2 s \frac{\mu}{4}}$

Continuing the proof sketch for Lemma 1

Let H be the $\sqrt{\epsilon' n}$ vertices of the the highest degree. Here ϵ' will be chosen to satisfy $(1 - \epsilon') \cdot (1/2 - \epsilon') \leq (1/2 - \epsilon)$

Assume that the random selection of nodes in the algorithm was restricted to just $L = V \setminus H$.

Of course, by removing high degree vertices from the random sampling, the probability of concluding that $\bar{d} \leq \frac{1}{2}(1 - \epsilon)d$ increases.

Claim $\sum_{i \in L} d_i \geq (\frac{1}{2} - \epsilon') \sum_{i \in V} d_i - \epsilon' n$

The Claim is a counting algorithm noting that the $\sum_{i \in V}$ counts every edge twice while $\sum_{i \in L}$ might omit $\epsilon' n$ edges within H and only count edges between H and L once.

Continuation of proof of Lemma 1

Thus the average degree in L is at least $\frac{1}{2}(\bar{d} - \epsilon')$. So it remain to obtain a lower bound on the average degree of vertices in L . We use the following observations:

- Let d_H = minimum degree of any vertex in H
- Let X_j = degree of a vertex $v_j \in S_i$ which implies $X_j \in [1, d_H]$
- Let $Z_j = X_j/d_H$
- The Chernoff bound and the bound we have for the average degree of vertices in L , then gives us : $Prob[a_i < (1/2)(1 - \epsilon)\bar{d}] \leq e^{-\frac{\epsilon^2 s \mathbb{E}[X_j]}{4d_H}}$ using the Chernoff bound.

If $s = |S_i|$ were sufficiently large (i.e. $s \geq \epsilon^2 \frac{d_H}{\mathbb{E}[X_j]}$), we are done. But we would like the bound to not depend on d_H and $\mathbb{E}[X_j]$.

This is handled by considering two cases; namely for when $d_H < |H|$ and when $d_H \geq |H|$.

Understanding the input query model

- As we initially noted, sublinear time algorithms almost invariably sample (i.e. query) the input in some way. The nature of these queries will clearly influence what kinds of results can be obtained.
- Feige's [2006] algorithm for estimating the average degree uses only "degree queries"; that is, "what is the degree of a vertex v ".
- Feige shows that in this degree query model, any algorithm that achieves a $(2 - \epsilon)$ approximation (for any $\epsilon > 0$) requires time $\Omega(n)$.
- In contrast, Goldreich and Ron [2008] consider the same average degree problem in the "neighbour query" model; that is, upon a query (v, j) , the query oracle returns the j^{th} neighbour of v or a special symbol indicating that v has degree less than j . A degree query can be simulated by $\log n$ neighbour queries.
- Goldreich and Ron show that in the neighbour query model, that the average degree \bar{d} can be $(1 + \epsilon)$ approximated (with one sided error probability $2/3$) in time $O(\sqrt{n} \text{poly}(\log n, \frac{1}{\epsilon}))$
- They show that $\Omega(\sqrt{n/\epsilon})$ queries is necessary to achieve a $(1 + \epsilon)$ approximation.

Approximating the size of a maximum matching in a bounded degree graph

- We recall that the size of any *maximal* matching is within a factor of 2 of the size of a maximum matching. Let m be smallest possible maximal matching.
- Our goal is to compute with high probability a *maximal* matching in time depending only on the maximum degree D .

Nguyen and Onak Algorithm

Choose a random permutation p of the edges $\{e_j\}$

% Note: this will be done “on the fly” as needed

The permutation determines a maximal matching M as given by the greedy algorithm that adds an edge whenever possible.

Choose $r = O(D/\epsilon^2)$ nodes $\{v_i\}$ at random

Using an “oracle” let X_i be the indicator random variable for whether or not vertex v_i is in the maximal matching.

Output $\tilde{m} = \sum_{i=1 \dots r} X_i$

Performance and time for the maximal matching

Claims

- 1 $m \leq \tilde{m} \leq m + \epsilon n$ where $m = |M|$.
 - 2 The algorithm runs in time $2^{O(D)}/\epsilon^2$
- This immediately gives an approximation of the *maximum* matching m^* such that $m^* \leq \tilde{m} \leq 2m^* + \epsilon n$
 - A more involved algorithm by Nguyen and Onak yields the following result:

Nguyen and Onak maximum matching result

Let $\delta, \epsilon > 0$ and let $k = \lceil 1/\delta \rceil$. There is a randomized one sided algorithm (with probability $2/3$) running in time $\frac{2^{O(D^k)}}{\epsilon^{2k+1}}$ that outputs a maximum matching estimate \tilde{m} such that $m^* \leq \tilde{m} \leq (1 + \delta)m^* + \epsilon n$.

Property Testing

- Perhaps the most prevalent and useful aspect of sublinear time algorithms is for the concept of property testing. This is its own area of research with many results.
- Here is the concept: Given an object G (e.g. a function, a graph), test whether or not G has some property P (e.g. G is bipartite) or is in some sense far away from that property.
- The tester determines with sufficiently high probability (say $2/3$) if G has the property or is “ ϵ -far” from having the property. The tester can answer either way if G does not have the property but is “ ϵ -close” to having the property.
- We will usually have a 1-sided error in that we will always answer YES if G has the property.
- We will see what it means to be “ ϵ -far” (or close) from a property by some examples.

Tester for linearity of a function

- Let $f : Z_n \rightarrow Z_n$; f is linear if $\forall x, y \ f(x + y) = f(x) + f(y)$.
- Note: this will really be a test for group homomorphism
- f is said to be ϵ -close to linear if its values can be changed in at most a fraction ϵ of the function domain arguments (i.e. changing the values of at most ϵn elements of Z_n) so as to make it a linear function. Otherwise f is said to be ϵ -far from linear.

The tester

Repeat $4/\epsilon$ times

Choose $x, y \in Z_n$ at random

If $f(x) + f(y) \neq f(x + y)$

then Output f is not linear

End Repeat If all these $4/\epsilon$ tests succeed then Output linear

- Clearly if f is linear, the tester says linear.
- For $\epsilon < 2/9$, if f is ϵ -far from being linear then the probability of detecting this is at least $2/3$.

Testing a list for monotonicity

- Given a list $A[i] = x_i, i = 1 \dots n$ of distinct elements, determine if A is a monotone list (i.e. $i < j \Rightarrow A[i] < A[j]$) or is ϵ -far from being monotone in the sense that more than $\epsilon * n$ list values need to be changed in order for A to be monotone.
- The algorithm randomly chooses $2/\epsilon$ random indices i and performs binary search on x_i to determine if x_i is in the list. The algorithm reports that the list is monotone if and only if all binary searches succeed.
- Clearly the time bound is $O(\log n/\epsilon)$ and clearly if A is monotone then the tester reports monotone.
- If A is ϵ -far from monotone, then the probability that a random binary search will succeed is at most $(1 - \epsilon)$ and hence the probability of the algorithm failing to detect non-monotonicity is at most $(1 - \epsilon)^{\frac{2}{\epsilon}} \leq \frac{1}{e^2}$

Graph Property testing

- Graph property testing is an area by itself. There are several models for testing graph properties.
- Let $G = (V, E)$ with $n = |V|$ and $m = |E|$.
- Dense model: Graphs represented by adjacency matrix. Say that graph is ϵ -far from having a property P if more than ϵn^2 matrix entries have to be changed so that graph has property P .
- Sparse model, bounded degree model: Graphs represented by vertex adjacency lists. Graph is ϵ -far from property P is at least ϵm edges have to be changed.
- In general there are substantially different results for these two graph models.

The property of being bipartite

- In the dense model, there is a constant time one-sided error tester. The tester is (once again) conceptually what one might expect but the analysis is not at all immediate.

Goldreich, Goldwasser, Ron bipartite tester

Pick a random subset S of vertices of size $r = \Theta\left(\frac{\log(\frac{1}{\epsilon})}{\epsilon^2}\right)$

Output bipartite iff the induced subgraph is bipartite

- Clearly if G is bipartite then the algorithm will always say that it is bipartite.
- The claim is that if G is ϵ -far from being bipartite then the algorithm will say that it is not bipartite with probability at least $2/3$.
- The algorithm runs in time quadratic in the size of the induced subgraph (i.e. the time needed to create the induced subgraph).

Testing bipartiteness in the bounded degree model

- Even for degree 3 graphs, $\Omega(\sqrt{n})$ queries are required to test for being bipartite or ϵ -far from being bipartite. Goldreich and Ron [1997]
- There is a nearly matching algorithm that uses $O(\sqrt{n} \text{poly}(\log n/\epsilon))$ queries. The algorithm is based on random walks in a graph and utilizes the fact that a graph is bipartite iff it has no odd length cycles.

Goldreich and Ron [1999] bounded degree algorithm

Repeat $O(1/\epsilon)$ times

Randomly select a vertex $s \in V$

If algorithm $OddCycle(s)$ returns cycle found then REJECT

End Repeat

If case the algorithm did not already reject, then ACCEPT

- $OddCycle$ performs $\text{poly}(\log n/\epsilon)$ random walks from s each of length $\text{poly}(\log n/\epsilon)$. If some vertex v is reached by both an even length and an odd length prefix of a walk then report cycle found; else report odd cycle not found

Sublinear space: A slight detour into complexity theory

- Sublinear space has been an important topic in complexity theory since the start of complexity theory. While not as important as the $P = NP$ or $NP = co - NP$ question, there are two fundamental space questions that remain unresolved:
 - 1 Is $NSPACE(S) = DSPACE(S)$ for $S \geq \log n$?
 - 2 Is P contained in $DSPACE(\log n)$ or $\cup_k SPACE(\log^k n)$? Equivalently, is P contained in polylogarithmic parallel time.
- Savitch [1969] showed a non deterministic S space bounded TM can be simulated by a deterministic S^2 space bounded machine (for space constructible bounds S).
- Further in what was considered a very surprising result, Immerman [1987] and independently Szelepcsényi [1987] $NSPACE(S) = co - NSPACE(S)$. (Savitch's result was also considered surprising by some researchers when it was announced.)

USTCON vs STCON

We let *USTCON* (resp. *STCON*) denote the problem of deciding if there is a path from some specified source node s to some specified target node t in an undirected (resp. directed) graph G .

- As previously mentioned the Aleliunas' et al [1979] random walk result showed that *USTCON* is in $RSPACE(\log n)$ and after a sequence of partial results about *USTCON*, Reingold [2008] was eventually able to show that *USTCON* is in $DSPACE(\log n)$
- It remains open if
 - ① *STCON* (and hence $NSPACE(\log n)$) is in $RSPACE(\log n)$ or even $DSPACE(\log n)$.
 - ② $STCON \in RSPACE(S)$ or even $DSAPCE(S)$ for any $S = o(\log^2 n)$
 - ③ $RSPACE(S) = DSPACE(S)$.

The streaming model

- In the data stream model, the input is a sequence A of input items (or input elements) a_1, \dots, a_n which is assumed to be too large to store in memory. Let $a_i \in [1, D]$

Small notational dilemma: The seminal paper in the streaming area is the Alon, Matias and Szegedy (AMS) paper for computing the frequency moment problem. Their notation is that a stream is a sequence a_1, \dots, a_m with $a_i \in \{1, 2, \dots, n\}$. To be more consistent with the online literature, I will use n for the length of the sequence and D for the domain of the a_i with the exception of the discussion of the AMS paper where I will use their notation.

- We usually assume that n is not known and hence one can think of this model as a type of online or dynamic algorithm.
- The space available $S(n, D)$ is some sublinear function. The input items stream by and one can only store information in space S .

The streaming model continued

- In some papers, space is measured in bits (which is what we will do) and sometimes in words, each word being $O(\log n)$ bits.
- It is also desirable that each input item is processed efficiently, say $\log(n) + \log(m)$ time, and perhaps even in time $O(1)$ (assuming we are counting operations on words as $O(1)$).
- The initial (and primary) work in streaming algorithms is to approximately compute some function (say a statistic) of the data or identify some particular item(s) in the data stream.
- Lately, the model has been extended to consider “semi-streaming” algorithms for optimization problems. For example, for a graph problem such as matching for a graph $G = (V, E)$, the goal is to obtain a good approximation using space $\tilde{O}(|V|)$ rather than $O(|E|)$.
- Most results concern the space required for a one pass algorithm. But there are results concerning multi-pass algorithms and also results concerning the tradeoff between the space and number of passes.

An example of a deterministic streaming algorithms

As in sublinear time, it will turn out that almost all of the results in this area are for randomized algorithms. Here is one exception.

The missing item problem

Suppose we are given a stream $A = a_1, \dots, a_{n-1}$ and we are promised that the stream A is a permutation of $\{1, \dots, m\} - \{x\}$ for some integer x in $[1, n]$. (Here $D = n$.) The goal is to compute the missing x .

- Space n is obvious using a bit vector $c_j = 1$ iff j has occurred.
- Instead we know that $\sum_{j \in A} j = n(n+1)/2 - x$.
So if $s = \sum_{i \in A} a_i$, then $x = n(n+1)/2 - s$.
This uses only $2 \log m$ space and constant time/item.

Generalizing to k missing elements

Now suppose we are promised a stream A of length $n - k$ whose input elements consist of a permutation of $n - k$ distinct elements in $\{1, \dots, n\}$. We want to find the missing k elements.

- Generalizing the one missing element solution, to the case that there are k missing elements we can (for example) maintain the sum of j^{th} powers ($1 \leq j \leq k$) $s_j = \sum_{i \in A} (a_i)^j = c_j(n) - \sum_{i \notin A} x_i^j$. Here $c_j(m)$ is the closed form expression for $\sum_{i=1}^n i^j$. This results in k equations in k unknowns using space $k^2 \log n$ but without an efficient way to compute the solution.
- As far as I know there may not be an efficient small space *deterministic* streaming algorithm for this problem.
- Using randomization, much more efficient methods are known; namely, there is a streaming alg with space and time/item $O(k \log k \log n)$; it can be shown that $\Omega(k \log(n/k))$ space is necessary.

Some well-studied streaming problems

- Computing **frequency moments**. Let $A = a_1 \dots a_m$ be a data stream with $a_i \in [n] = \{1, 2, \dots, n\}$. Let m_i denote the number of occurrences of the value i in the stream A . For $k \geq 0$, the k^{th} frequency moment is $F_k = \sum_{i \in [n]} (m_i)^k$. The frequency moments are most often studied for integral k .
 - ① $F_1 = m$, the length of the sequence which can be simply computed.
 - ② F_0 is the number of distinct elements in the stream
 - ③ F_2 is a special case of interest called the repeat index (also known as Gini's homogeneity index).
- Finding **k -heavy hitters**; i.e. those elements appearing more than m/k times in stream A .
- Finding **rare or unique elements** in A .

The majority element problem

While most streaming algorithms concern one pass over the input stream, there are results that use two or more passes.

One relatively easy (but still very interesting) result is the Misra-Gries algorithm for computing k heavy hitters. As a special case, we have the majority problem (i.e. the k -hitter problem for $k = 2$).

There is a temptation to solve this problem by divide and conquer; divide the sequence in half, find the heavy hitters in each half and then check.

The streaming model facilitates thinking about a much better solution. In the case of majority, let's just try to maintain one possible candidate in the first pass and then check to see if the candidate is a true more than majority item in the second pass. See the Chakrabarti Lecture notes.

The Misra-Gries algorithm

Maintain a candidate for the majority element and a counter for that candidate.

When the counter is empty, the next element in the stream becomes the candidate.

Every time the next element in the stream is the candidate increase the counter by 1. If the next element is not the candidate decrease the counter by 1.

Claim: If there is a majority element then it has to be the current candidate.

We can use a second pass over the elements to check if the candidate occurs more than $n/2$ times.

The space used is $O(\log n + \log D)$ and the time is $(\log n)$ (or $O(1)$ if counting element comparisons) per input element.

Some comments on the Misra-Gries algorithm

It can be shown that no (even randomized) one pass streaming algorithm can return a truly majority element. So the second pass is needed to verify a candidate.

As suggested the majority algorithm can be extended to obtain solve the k heavy hitters problem for any small k . This is related to your assignment so I will leave this comment for you to think about or read in Chakrabarti.

Consider the following extension to the ϵ -approximate ϕ -hitters problem where the algorithm is required to return a set S of elements such that

- 1) Every element in S appears at least ϕn times.
- 2) S does not contain any elements less than $(\phi - \epsilon)n$ times.

The Misra-Gries algorithm can be extended to solve this problem by setting $k = \frac{1}{\epsilon}$ and then outputting all elements a that occur at least $(\phi - \epsilon)n$ times.

Another way to solve an upper bound version of the ϵ -approximate ϕ -heavy hitters.

Suppose we want The following ideas are another very general approach to solving various streaming problems. We introduce it here as it is an easy case to expose the ideas. And now we are going to use randomization.

Suppose we just want the counts for the ϕ -heavy hitters. We want to get all the counts and have the guarantee that $\text{Prob}[\text{count} \geq \text{truecount} + \epsilon m]$ is small.

The idea is to hash values and then just keep counts of the hashed values. Let $k = 2/\epsilon$. We use a 2-universal hash function h so that $\text{Prob}[h(v) = h(v')] = 1/k$ so any two distinct values v and v' .

Count Min Sketch uses a number of different hash functions (in parallel) and then takes the min of these counters.

Frequency Moments: What is known about computing F_k ?

Given an error bound ϵ and confidence bound δ , the goal in the frequency moment problem is to compute an estimate F'_k such that $\text{Prob}[|F_k - F'_k| > \epsilon F_k] \leq \delta$.

- The seminal paper in this regard is by Alon, Matias and Szegedy (AMS) [1999]. AMS establish a number of results:
 - 1 For $k \geq 3$, there is an $\tilde{O}(m^{1-1/k})$ space algorithm. The \tilde{O} notation hides factors that are polynomial in $\frac{1}{\epsilon}$ and polylogarithmic in $m, n, \frac{1}{\delta}$.
 - 2 For $k = 0$ and every $c > 2$, there is an $O(\log n)$ space algorithm computing F'_0 such that $\text{Prob}[(1/c)F_0 \leq F'_0 \leq cF_0 \text{ does not hold}] \leq 2/c$.
 - 3 For $k = 1$, $\log n$ is obvious to exactly compute the length but an estimate can be obtained with space $O(\log \log n + 1/\epsilon)$
 - 4 For $k = 2$, they obtain space $\tilde{O}(1) = O(\frac{\log(1/\delta)}{\epsilon^2})(\log n + \log m)$
 - 5 They also show that for all $k > 5$, there is a (space) lower bound of $\Omega(m^{1-5/k})$.

Results following AMS

- A considerable line of research followed this seminal paper. Notably settling conjectures in AMS:
- The following results apply to real as well as integral k .
 - ① An $\tilde{\Omega}(m^{1-2/k})$ space lower bound for all $k > 2$ (Bar Yossef et al [2002]).
 - ② Indyk and Woodruff [2005] settle the space bound for $k > 2$ with a matching upper bound of $\tilde{O}(m^{1-2/k})$
- The basic idea behind these randomized approximation algorithms is to define a random variable Y whose expected value is close to F_k and variance is sufficiently small such that this r.v. can be calculated under the space constraint.
- We will just sketch the (non optimal) AMS results for F_k for $k > 2$ and the result for F_2 .

The AMS F_k algorithm

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

AMS algorithm for F_k

The output Y of the algorithm is the median of s_2 random variables Y_1, Y_2, \dots, Y_{s_2} where Y_i is the mean of s_1 random variables $X_{ij}, 1 \leq j \leq s_1$. All X_{ij} are independent identically distributed random variables. Each $X = X_{ij}$ is calculated in the same way as follows: Choose random $p \in [1, \dots, m]$, and then see the value of a_p . Maintain $r = |\{q | q \geq p \text{ and } a_q = a_p\}|$. Define $X = m(r^k - (r-1)^k)$.

- Note that in order to calculate X , we only require storing a_p (i.e. $\log n$ bits) and r (i.e. at most $\log m$ bits). Hence the Each $X = X_{ij}$ is calculated in the same way using only $O(\log n + \log n)$ bits.
- For simplicity we assume the input stream length m is known but it can be estimated and updated as the stream unfolds.
- We need to show that $\mathbf{E}[X] = F_k$ and that the variance $\text{Var}[X]$ is small enough so as to use the Chebyshev inequality to show that $\text{Prob}[|Y_i - F_k| > \epsilon F_k]$ is small.

AMS analysis sketch

- Showing $E[X] = F_k$.

$$\frac{m}{m}[(1^k + (2^k - 1^k) + \dots + (m_1^k - (m_1 - 1)^k)) +$$

$$(1^k + (2^k - 1^k) + \dots + (m_2^k - (m_2 - 1)^k)) + \dots + \\ (1^k + (2^k - 1^k) + \dots + (m_n^k - (m_n - 1)^k))]$$

(by telescoping)

$$= \sum_i^n m_i^k$$

$$= F_k$$

AMS analysis continued

- Y is the median of the Y_i . It is a standard probabilistic idea that the median Y of identical r.v.s Y_i (each having constant probability of small deviation from their mean F_k) implies that Y has a high probability of having a small deviation from this mean.
- $E[Y_i] = E[X]$ and $\text{Var}[Y_i] \leq \text{Var}[X]/s_1 \leq E[X^2]/s_1$.
- The result needed is that $\text{Prob}[|Y_i - F_k| > \epsilon F_k] \leq \frac{1}{8}$
- The Y_i values are an average of independent $X = X_{ij}$ variables but they can take on large values so that instead of Chernoff bounds, AMS use the Chebyshev inequality:

$$\text{Prob}[|Y - E[Y]| > \epsilon E[Y]] \leq \frac{\text{Var}[Y]}{\epsilon^2 E[Y]}$$

- It remains to show that $E[X^2] \leq kF_1F_{2k-1}$ and that $F_1F_{2k-1} \leq n^{1-1/k}F_k^2$

Sketch of F_2 improvement

- They again take the median of $s_2 = 2 \log(\frac{1}{\delta})$ random variables Y_i but now each Y_i will be the sum of only a constant number $s_1 = \frac{16}{\epsilon^2}$ of identically distributed $X = X_{ij}$.
- The key additional idea is that X will not maintain a count for each particular value separately but rather will count an appropriate sum $Z = \sum_{t=1}^n b_t m_t$ and set $X = Z^2$.
- Here is how the vector $\langle b_1, \dots, b_n \rangle \in \{-1, 1\}^n$ is randomly chosen.
- Let $V = \{v_1, \dots, v_h\}$ be a set of $O(n^2)$ vectors over $\{-1, 1\}$ where each vector $v_p = \langle v_{p,1}, \dots, v_{p,n} \rangle \in V$ is a 4-wise independent vector of length n .
- Then p is selected uniformly in $\{1, \dots, h\}$ and $\langle b_1, \dots, b_n \rangle$ is set to v_p .