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A. Pavan¹, Kanat Tangwongsan², Srikanta Tirthapura¹, Kun-Lung Wu²

¹Iowa State University

²IBM Research Division Thomas J. Watson Research Center P.O. Box 208 Yorktown Heights, NY 10598 USA



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Counting and Sampling Triangles from a Graph Stream^{*}

A. Pavan^{*} Kanat Tangwongsan[†] Srikanta Tirthapura^{*} Kun-Lung Wu[†]

Iowa State University* and IBM Research †

Abstract

This paper presents a new space-efficient algorithm for counting and sampling triangles, and more generally, constant-sized cliques, in a massive graph whose edges arrive as a stream. When compared with prior work, our algorithm yields significant improvements in the space and time complexity for these fundamental problems. Our algorithm is simple to implement and has very good practical performance on large graphs.

Keywords: graph streams, data streams, triangle counting, cliques, social networks, sampling

1 Introduction

Triangle counting has emerged as an important building block in the study of social networks [WF94, New03], identifying thematic structures of networks [EM02], spam and fraud detection [BBCG08], link classification and recommendation [TDM⁺11], and more. In these applications, streaming algorithms provide an attractive option for real-time processing of live data; they also benefit the analysis of large disk-resident graph data, allowing computations in one or a small number of passes over the data.

In this paper, we address the question of counting and sampling triangles, as well as complete subgraphs, in the adjacency stream model. Specifically, we study the following closely related problems:

- (1) *Triangle Counting:* maintain an (accurate) estimate of the number of triangles in a graph;
- (2) Triangle Sampling: maintain a random triangle from the set of all triangles in a graph; and
- (3) counting and sampling cliques of 4 or more vertices (K_4, K_5, \ldots) .

In the *adjacency stream model* [BYKS02, JG05, BFL⁺06], a graph G = (V, E) is presented as a stream of edges $S = \langle e_1, e_2, e_3, \dots, e_{|E|} \rangle$. In this notation, e_i denotes the *i*-th edge in the stream order, which is arbitrary and potentially chosen by an adversary. For this graph, let n = |V|, m = |E|, $\mathcal{T}(G)$ denote set of all triangles, and $\tau(G)$ denote the number of triangles (i.e., $\tau(G) = |\mathcal{T}(G)|$). We will assume that the input graph is simple (no parallel edges, no self-loops, and no multiple copies of the same edge).

Our algorithms are randomized and provide the following notion of probabilistic guarantees: for parameters $\varepsilon, \delta \in [0, 1]$, an (ε, δ) -approximation for a quantity X is a random variable \hat{X} such that $|\hat{X} - X| \leq \varepsilon X$ with probability at least $1 - \delta$. We write $s(\varepsilon, \delta)$ as a shorthand for $1/\varepsilon^2 \cdot \log(1/\delta)$.

1.1 Our Contributions

— **Neighborhood Sampling:** We present *neighborhood sampling*, a new technique for counting and sampling cliques from a graph stream. Neighborhood sampling is a multi-level inductive random sampling procedure, where a random edge in the stream is first sampled, and then in subsequent steps, an edge with an endpoint in common to the sampled edge(s) is sampled. We show that this simple technique leads to significant improvements in the space complexity of triangle counting and related problems.

— Counting and Sampling Triangles: Using neighborhood sampling, we present a one-pass streaming algorithm for triangle counting and triangle sampling. The space complexity of triangle counting is $O(s(\varepsilon, \delta)m\Delta/\tau(G))$ and triangle sampling is $O(m\Delta/\tau(G))$, where Δ is the maximum degree of any vertex. This significantly improves upon

^{*}E-mail addresses: pavan@cs.iastate.edu, ktangwo@us.ibm.com, snt@iastate.edu, klwu@us.ibm.com

	Space Complexity	Time Per Element
Buriol <i>et al.</i> [BFL ⁺ 06]	$O\left(s(\varepsilon,\delta)\frac{mn}{\tau(G)} ight)$	$O\left(1+s(\varepsilon,\delta)\tfrac{n\log m}{\tau(G)}\right)$
Jowhari and Ghodsi [JG05]	$s_2 = O\left(s(\varepsilon, \delta) \frac{m\Delta^2}{\tau(G)}\right)$	$\Theta(s_2)$
This Paper (amortized with bulk processing)	$O\left(s(\varepsilon,\delta)\frac{m\Delta}{\tau(G)} ight)$	$\Theta(1)$

Table 1: Performance of streaming algorithms for counting triangles in a graph. The space complexity is expressed in terms of the number of words used, and assumes that vertex identifiers, and counters, including the number of edges and vertices, can be stored in a constant number of words.

prior algorithms for the same problem, as shown in Table 1. We provide a sharper bound for the space complexity of our algorithm in terms of the "tangle coefficient" of the graph, which we define in our analysis. While in this worst case, this results in the space bound we have stated above, in typical cases, it can be smaller.

We also present a method for quickly processing edges in bulks, which leads to an amortized constant processing time per edge. This allows the possibility of processing massive graphs quickly even on a modest machine.

— Counting and Sampling Cliques: We extend neighborhood sampling to the problem of sampling and counting the set of all cliques of size ℓ in the graph, $\ell \ge 4$. For $\ell = 4$, the space complexity of the counting algorithm is $O(s(\varepsilon, \delta) \cdot \eta / \tau_4(G))$, and the space complexity of the sampling algorithm is $O(\eta / \tau_4(G))$, where $\eta = \max\{m\Delta^2, m^2\}$ and $\tau_4(G)$ is the number of 4-cliques in G. General bounds for ℓ -cliques are presented in Section 3. To our knowledge, this is the best space complexity for counting the number of ℓ -cliques in a graph in the streaming model and improves on prior work due to Kane *et al.* [KMSS12].

— Experiments: Our experiments with real-world large graphs show that our streaming algorithm for counting triangles is fast and accurate in practice. For instance, the Orkut network (for a description, see Section 4) with 117 million edges and 633 million triangles can be processed in 103 seconds, with a (mean) relative error of 3.55 percent, using 1 million instances of estimators (and hence a few MB of memory). This experiment was run on a laptop, with an implementation that did not use parallelism. Our experiments show that in order to get good estimates, far fewer than $\Theta(s(\varepsilon, \delta)m\Delta/\tau(G))$ estimators may be necessary.

Prior Work: For triangle counting in adjacency streams, Bar-Yossef *et al.* [BYKS02] present the first algorithms based on reductions to the problem of computing the zeroth and second frequency moments of appropriately defined streams, derived from the edge stream of the graph. Their algorithm on the adjacency stream model takes space $s = O(\frac{s(\varepsilon,\delta)}{\varepsilon} \cdot (mn/\tau(G))^3)$ and poly(s) time per item. They also present a lower bound showing that in general, the worst case streaming complexity of approximating $\tau(G)$ is $\Omega(n^2)$.

The space and time bounds were subsequently improved. Jowhari and Ghodsi [JG05] present a one-pass streaming algorithm that uses space and per-edge processing time of $O(s(\varepsilon, \delta)m\Delta^2/\tau(G))$. Our algorithm significantly improves upon this algorithm in both space and time complexity. Note that Δ for large graphs can be significant in practice; for instance, the Orkut graph has a maximum degree of greater than 66,000. They also present a three pass streaming algorithm with space and per-edge processing time of $O(s(\varepsilon, \delta) \cdot (1 + T_2(G)/\tau(G)))$, where $T_2(G)$ is the number of vertex triples in the graph with exactly two edges connecting them. Later, Buriol *et al.* [BFL⁺06] present algorithms for counting the number of triangles with space complexity $O(s(\varepsilon, \delta)mn/\tau(G))$. If the maximum degree Δ is small compared with *n*, our algorithm substantially improves upon theirs in terms of space. Another difference is that the algorithm of [BFL⁺06] needs to know the set of vertices in the graph stream in advance, but ours does not. This can be a significant advantage in practice when vertices are being dynamically added to graph, or being discovered by the stream processor.

On counting cliques, Kane *et al.* [KMSS12] present estimators for the number of occurrences of an arbitrary subgraph H in the stream. When applied to counting cliques on ℓ vertices in a graph, their space complexity is $O\left(s(\varepsilon, \delta) \cdot m^{\binom{\ell}{2}} / \tau_{\ell}^2(G)\right)$ which is much higher than the space complexity that we obtain. We note that their

algorithm works in the model where edges can be inserted or deleted (turnstile model), while ours is insert-only.

Related Work: Manjunath *et al.* [MMPS11] present an algorithm for counting the number of cycles of length k in a graph; their algorithm works under dynamic inserts and deletes of edges. Since a triangle is also a cycle, this algorithm applies to counting the number of triangles in a graph, but uses space and per item processing time $\Theta(s(\varepsilon, \delta)m^3/\tau^2(G))$. When compared with our algorithm, their space and time bound can be much larger, especially for graphs with a small number of triangles. Recent work on graph sketches by Ahn, Guha, and McGregor [AGM12] also yield algorithms for counting triangles in a graph, with space complexity, whose dependence on m and n is the same as in [BFL⁺06].

Buriol *et al.* [BFLS07] present algorithms to estimate the clustering index of a graph in the *incidence stream* model, which assumes that all edges incident at a vertex arrive together, and that each edge appears twice, once for each endpoint. We note that in the incidence streams model, counting triangles is an easier problem, and there are streaming algorithms [BFL+06] that use space $O(s(\varepsilon, \delta)(1 + T_2(G)/\tau(G)))$. In this work, we focus on the adjacency streams model which is a more realistic model for real-time analytics on an evolving graph.

Becchetti *et al.* [BBCG08] present algorithms for counting the number of triangles in a graph in a model where the processor is allowed $O(\log n)$ passes through the data and O(n) memory. Their algorithm also returns for each vertex, the number of triangles that the vertex is a part of. There is a significant body of work on counting the number of triangles in a graph in the non-streaming setting, for example [SV11, TKMF09]. We do not attempt to survey this literature. An experimental study of algorithms for counting and listing triangles in a graph is presented in [SW05].

Roadmap: In Section 2, we present our technique and its use in counting and sampling triangles, followed by extensions to counting and sampling cliques in Section 3, experimental results in Section 4 and extensions to sliding windows in Section 5.

Preliminaries: For an edge e, let V(e) denotes the two end vertices of e. We say that two edges are adjacent to each other if they share a vertex. Given an edge e_i , the *neighborhood of* e_i , denoted by $N(e_i)$, is the set of all edges in the stream that arrive after e_i and are adjacent to e_i . Let $c(e_i)$ denote the size of $N(e_i)$. For $\ell \ge 4$, let $\mathcal{T}_{\ell}(G)$ denote the set of all ℓ -cliques of graph G, and $\tau_{\ell}(G)$ the number of ℓ -cliques. Further, for a triangle $t^* \in \mathcal{T}(G)$, define $C(t^*)$ to be c(f), where f is its first edge in the stream. Our algorithms use a procedure coin(p) which returns heads with probability p. We assume this procedure takes constant time.

2 Sampling and Counting Triangles

In this section, we present algorithms to sample and count triangles. We begin by describing *neighborhood sampling*, a basic method upon which we build an algorithm for counting triangles (Section 2.2), an efficient implementation for bulk processing (Section 2.3), and an algorithm for sampling triangles (Section 2.4).

2.1 Neighborhood Sampling Algorithm for Triangles



Figure 1: An example graph, where the edges arrive in order $e_1, e_2, ...$ in the stream, forming triangles $t_1 = \{e_1, e_2, e_3\}$ and $t_2 = \{e_4, e_5, e_6\}$.

Overview. The algorithm first samples a random edge, say r_1 , from the edge stream; this can be maintained using reservoir sampling. It then samples a random edge from the set of edges in the stream that appear after r_1 and are adjacent to r_1 . That is, the second edge is sampled from the neighborhood of the first edge. This sample can also be maintained using reservoir sampling on the appropriate substream. Once such an edge, say r_2 , has been found, a potential triangle t made up of the two edges r_1 and r_2 is implicitly defined, and the algorithm waits for the third edge of t to appear in the stream and complete the potential triangle.

The triangle found by this procedure, however, is *not* necessarily uniformly chosen from $\mathcal{T}(G)$. As an example, in the graph in Figure 1, the probability that the neighborhood sampling procedure chooses triangle t_1 is the probability that e_1 is chosen into r_1 (which is $\frac{1}{10}$), and then from among the edges adjacent to e_1 (i.e., e_2 and e_3), e_2 is chosen into r_2 , for a total probability of $\frac{1}{2} \cdot \frac{1}{10} = \frac{1}{20}$. But the probability of choosing t_2 is the probability of choosing e_4 into

 r_1 (which is $\frac{1}{10}$), and then from among those edges adjacent to e_4 and arrive after e_4 (i.e., $\{e_5, e_6, e_7, e_8, e_9, e_{10}, e_{11}\}$), e_5 is chosen into r_2 (which is $\frac{1}{7}$), for a total probability of $\frac{1}{7} \cdot \frac{1}{10} = \frac{1}{70}$. This bias poses a challenge in our algorithms but can be normalized away by keeping track of how much bias is incurred on the potential triangle.

We briefly contrast our algorithm with an algorithm for adjacency streams due to Buriol *et al.* [BFL⁺06], which also employs random sampling. Like ours, their algorithm first samples a random edge from the stream, say r_1 , but then unlike ours, it picks a random vertex that is not necessarily incident on an endpoint of r_1 . The edge and the vertex together form a potential triangle, and the algorithm then waits for the triangle to be completed by the remaining two edges. In our algorithm, instead of selecting a random third vertex, we select a vertex that is already connected to r_1 . This leads to a greater chance that the triangle is completed, and hence better space bounds.

We now describe the neighborhood sampling algorithm in detail. The algorithm maintains the following state:

- Edge r_1 , sampled uniformly from among all edges so far. We call this the "level 1 edge".
- Edge r_2 , sampled uniformly from among all edges in $N(r_1)$, i.e., those edges in the graph stream that are adjacent to r_1 and come after r_1 . We call this the "level 2 edge".
- Counter c, equal to $c(r_1) = |N(r_1)|$, i.e, the number of edges that are adjacent to r_1 and have appeared after r_1 .
- A triangle t that is potentially a sample

Presented in Algorithm 1 is the neighborhood sampling algorithm, which is used to maintain the state. Algorithms 2 and 3 build on this algorithm to solve triangle counting and triangle sampling, respectively.

Algorithm 1: Algorithm NSAMP-TRIANGLE

```
Initialization: Set r_1, r_2, t to \phi, and c to 0.
Upon getting edge e_i, i \ge 1;
begin
     if coin(1/i) = "head" then
          /\!/ e_i is the new sampled edge at level 1.
          r_1 \leftarrow e_i;
          r_2 \leftarrow \phi; c \leftarrow 0; t \leftarrow \phi;
     else
          if e_i is adjacent to r_1 then
                c \leftarrow c + 1;
                if coin(1/c) = "head" then
                     // e_i is the new sampled edge at level 2.
                     r_2 \leftarrow e_i;
                     t \leftarrow \phi;
                else
                     if e_i forms a triangle with r_1 and r_2 then
                        t \leftarrow \{r_1, r_2, e_i\}
```

Lemma 2.1 Let t^* be a triangle in the graph. The probability that $t = t^*$ in the state maintained by Algorithm 1 after observing all edges (note t may be empty) is

$$\mathbf{Pr}[t=t^*] = \frac{1}{m \cdot C(t^*)}$$

where we recall that $C(t^*) = c(f)$ if f is the t^* 's first edge in the stream.

Proof: Let $t^* = \{f_1, f_2, f_3\}$ be a triangle in the graph, whose edges arrived in the order f_1, f_2, f_3 in the stream, so $C(t^*) = c(f_1)$ by definition. Let \mathcal{E}_1 be the event that f_1 is stored in r_1 , and \mathcal{E}_2 be the event that f_2 is stored in r_2 . We can easily check that the neighborhood sampling algorithm produces t^* at the end if and only if both \mathcal{E}_1 and \mathcal{E}_2 hold.

Now we know from reservoir sampling that $\mathbf{Pr}[\mathcal{E}_1] = \frac{1}{m}$. Furthermore, we claim that $\mathbf{Pr}[\mathcal{E}_2|\mathcal{E}_1] = \frac{1}{c(f_1)}$. This holds because given the event \mathcal{E}_1 , the edge r_2 is randomly chosen from $N(f_1)$, so the probability that $r_2 = f_2$ is exactly $1/|N(f_1)|$, which is $1/c(f_1)$, since c tracks the size of $N(r_1)$. Hence, we have

$$\mathbf{Pr}[t=t^*] = \mathbf{Pr}[\mathcal{E}_1 \cap \mathcal{E}_2] = \mathbf{Pr}[\mathcal{E}_1] \cdot \mathbf{Pr}[\mathcal{E}_2 \mid \mathcal{E}_1] = \frac{1}{m} \cdot \frac{1}{c(f_1)} = \frac{1}{m \cdot C(t^*)}$$

2.2 Counting Triangles in a Graph

The neighborhood sampling algorithm produces a triangle t with probability $\frac{1}{m \cdot C(t)}$. We show in Algorithm 2 how to build an unbiased estimator from this sampling algorithm. The basic idea is to output a value which counterbiases the probability so that in expectation, the contribution of a triangle is exactly 1. This is easy to achieve because we know both C(t) and the number of edges m. The following lemma formally shows that Algorithm 2 gives an unbiased estimator for the number of triangles.

Algorithm 2: COUNT-TRIANGLE	
Run Algorithm 1 and let t and c be the variables it maintains.	_
Return $c \cdot m$ if t is defined or 0 otherwise.	

Lemma 2.2 Let X denote the return value of Algorithm 2 after graph G has been observed. Then $\mathbf{E}[X] = \tau(G)$.

Proof: By Lemma 2.1, the probability that Algorithm 2 samples a particular triangle t^* is precisely $\mathbf{Pr}[t = t^*] = 1/mC(t^*)$. Further, the counting algorithm returns $mC(t^*)$ if $t = t^*$, and 0 if t is empty. Therefore,

$$\mathbf{E}[X] = \sum_{t^* \in \mathcal{T}(G)} mC(t^*) \cdot \mathbf{Pr}[t = t^*] = \tau(G).$$

To obtain an accurate estimate, we apply a standard technique which runs multiple copies of an unbiased estimator in parallel and outputs the average. In the following lemma, we give a bound on the number of copies needed to achieve an (ε, δ) -approximation. In Appendix A, we give a sharper space bound in terms of a measure we term "tangle index," which helps to explain why we usually need less space in practice.

Theorem 2.3 For any graph G, for parameters $0 \le \delta, \varepsilon \le 1$, there is a streaming algorithm that observes the edges of G in an arbitrary order, and returns an (ε, δ) -approximation for the number of triangles in G using space $O\left(\frac{1}{\varepsilon^2}\frac{m\Delta}{\tau(G)}\log\left(\frac{1}{\delta}\right)\right)$.

The proof follows from standard concentration bounds and Lemma 2.2; we defer it to Appendix C

2.3 A Nearly-Linear Time Algorithm for Triangle Counting

Our discussion thus far directly leads to a simple O(mr)-time implementation, where r is the number of copies of estimators we decide to maintain, but this leaves much to be desired in terms of performance. For large graphs, we would ideally like the algorithm to take time linear in the number of edges and the number of estimators.

The result in this section is motivated by the observation that in many real-world applications, the algorithm receives edges in bulk (e.g., block reads from disk, HTTP PUT requests). We show that it is possible to achieve significantly better performance through bulk processing. The basic intuition is that with bulk processing, we update the estimators much less often. As one example, by processing in bulks of $\Theta(r)$ edges, the total running time becomes $\Theta(m+r)$ using O(r) space—at the expense of a constant factor more space, we are able to achieve O(m+r) time bound as opposed to O(mr). More precisely, we obtain the following bounds:

Theorem 2.4 Let w be a block size and r be the number of estimators. There is an algorithm for triangle counting that on input a graph with m edges, runs in total time O(m(1 + r/w)) using space O(r + w).

We obtain this bound by developing a routine to process a block $B = \langle b_1, \ldots, b_{|B|} \rangle$ of edges in roughly O(|B|+r) time. Our goal is to advance the states of all r estimators to the point after including the block B, simulating the effects of playing these edges one by one faithfully. In the interest of space, we can only highlight a few key ingredients here and give detailed descriptions in Appendix B.

The challenge in fastforwarding over the block B lies in maintaining level-2 edges, finding the completed triangles, and keeping track of the counters; maintaining the level-1 edges is relatively straightforward. There are two important ingredients in our solution: The first ingredient is a data structure that maintains the degrees of about O(r) nodes; these degrees only count the edges in B. More specifically, let R be the set of the endpoints of all level-1 edges. As we make a pass through $b_1, b_2, \ldots, b_{|B|}$, we maintain for each $v \in R$, a counter λ_v which contains the number of edges in this block so far that is incident on v. This is useful for updating the counter c and for determining which level-2 edge to pick. As an example, for each estimator, its c value is increased by the difference between the λ values of the endpoints when its level-1 edge is encountered and the λ values at the end of the pass. This whole computation requires one pass and O(|B| + |R|) = O(|B| + r) time and space using a (hash) table which maps level-1 edges to their corresponding estimator states and a (hash) table storing the degree values.

The second ingredient is the observation that the event "the edge that increments the λ_x to a certain value z" uniquely identifies an edge in B. Given the degree information obtained previously, we can describe our random selection of a level-2 edge in this form—and by keeping a (hash) table mapping (vertex, desired-restricted-degree) to the corresponding estimator, a second pass through the data allows us to find level-2 replacement edges. Since the table contains at most O(r) entries, the space and time usage for this pass is O(|B| + r).

Finally, in the second pass, we also keep another (hash) table for queries of the form: "*if an edge e is seen, we have a complete triangle for estimator X*." This table is updated as we discover a new level-2 edge. The size of this table is, again, at most O(r). Hence, each block can be processed in *two* passes in O(|B| + r) space and time.

2.4 Sampling a Triangle

We now present an algorithm that picks a triangle of the graph stream uniformly at random. As discussed previously, the neighborhood sampling algorithm does not necessarily return a uniform sample. To fix it, we resort to a simple normalization procedure.

Algorithm 3: Algorithm SAMPLE-TRIANGLE: a query arrives for a random triangle
Run Algorithm 1 and let t and c be the variables it maintains.
Return the triangle t with probability $c/2\Delta$; else return "Failed".

Lemma 2.5 If a query for a random triangle is posed after observing a graph G, each triangle in $\mathcal{T}(G)$ is equally likely to be returned by Algorithm 3. Further, the probability that a triangle is returned by Algorithm 3 is $\frac{\tau(G)}{2m\Delta}$, where Δ is the maximum degree of any vertex in G.

Proof: The proof is similar to the proof of Lemma 2.2. By 2.1, the probability that Algorithm 2 samples a particular triangle t^* is precisely $\Pr[t = t^*] = 1/mC(t^*)$. Further, note that $C(t^*) \leq 2\Delta$. Therefore, if the triangle t^* was returned by Algorithm 1, the probability that t^* is passed on as our output is $\frac{1}{mC(t^*)} \cdot \frac{C(t^*)}{2\Delta} = \frac{1}{2m\Delta}$, where we note that $c = C(t^*)$ and the normalization factor $\frac{c}{2\Delta} \leq 1$. Finally, since the events of different triangles being returned are all disjoint from each other, the probability that some triangle is returned by the algorithm is $\frac{\tau(G)}{2m\Delta}$.

The following establishes the main theorem on triangle sampling.

Theorem 2.6 Assuming that $\tau(G) > 0$, for a parameter δ , $0 < \delta < 1$, there is a streaming algorithm with space complexity $\Theta\left(\frac{m\Delta \log(1/\delta)}{\tau(G)}\right)$ that observes graph G as an adjacency stream and returns a random triangle in G with probability at least $1 - \delta$.

Proof: Let $\alpha = \left(\frac{2m\Delta \ln(1/\delta)}{\tau(G)}\right)$. The streaming algorithm runs α copies of Algorithm 3. This algorithm fails to return a triangle only if all these α copies fail to produce a triangle; otherwise, it can return any of them. Therefore, the overall failure probability is at most $(1 - \beta)^{\alpha} \le e^{-\beta\alpha} \le \delta$, where $\beta = \frac{\tau(G)}{2m\Delta}$ is the success probability in Lemma 2.5. The space complexity follows since the space taken by each copy of the algorithm is a constant number of words.

A Lower Bound: It is natural to ask whether better space bounds are possible for triangle sampling and counting. In particular, can we meet the space complexity of $O(1+T_2(G)/\tau(G))$, which is the space-complexity of an algorithm for triangle counting in the incidence stream model [BFL⁺06]? Note that $T_2(G)$ is the number of vertex triples with exactly two edges in them. We show that this space bound is not possible in the adjacency stream model.

Lemma 2.7 There exists a graph G^* and an order of arrival of edges, such that any randomized streaming algorithm that can estimate the number of triangles in G^* with a relative error of better than 1/2 must have space complexity $\omega(1 + T_2(G)/\tau(G))$.

The proof uses a reduction from the index problem from communication complexity and is deferred to Appendix C in the interest of space. We note that the same proof technique also applies to triangle sampling.

3 Counting and Sampling Cliques

We extend the neighborhood sampling technique to counting and sampling from the set of cliques of size greater than 3. For ease of presentation, we focus on 4-cliques. Let $\mathcal{T}_4(G)$ denote the set of all 4-cliques in G, and $\tau_4(G)$ the number of 4-cliques in G. We partition $\mathcal{T}_4(G)$ into two classes, Type I cliques and Type II cliques, defined based on the order in which the edges of the clique appear in the stream. Let κ^* be a 4-clique and let f_1, f_2, \dots, f_6 be the order in which the edges of κ^* appear in the stream. We say that κ^* is a Type I clique if f_2 and f_1 share a common vertex, otherwise κ^* is a Type II clique. Clearly, every clique in $\mathcal{T}_4(G)$ is either a Type I clique or a Type II clique. Let $\mathcal{T}_4^1(G)$ denote the set of Type I cliques in G and $\mathcal{T}_4^2(G)$ denote the set of Type II cliques; let $\tau_4^1(g)$ and $\tau_4^2(G)$ denote the corresponding cardinalities. We present two estimators, one for $\tau_4^1(G)$ and the other for $\tau_4^2(G)$. The sum of the two estimators will be an estimator for $\tau_4(G)$. Due to space constraints all proofs of this section appear in Appendix D.

3.1 Neighborhood Sampling for 4-Cliques

Our first algorithm, described in Algorithm 4, is concerned with sampling Type I cliques. This algorithm maintains two sets, a sample of up to three edges, r_1, r_2, r_3 , and a potential clique, κ_1 . Each of the r_i s is also included in κ_1 , which could also contain additional edges required for completing the clique.

Lemma 3.1 Consider a Type I clique κ^* and let f_1, \ldots, f_6 be its edges in the order they appeared in the stream. After Algorithm 4 has processed the entire graph, the probability that κ_1 equals κ^* is $\frac{1}{m \cdot c(f_1) \cdot c(f_1, f_2)}$.

Next we describe a neighborhood sampling algorithm, Algorithm 5 that processes Type II cliques. This algorithm is simpler that Algorithm 4, and maintains a sample consisting of two edges r_1 and r_2 , and a potential clique κ_2 .

Lemma 3.2 Consider a Type II clique κ^* . After Algorithm 5 has processed the entire graph, the probability that κ_2 is equal to κ^* is $\frac{1}{m^2}$

Counting 4-cliques. We obtain following results for counting 4-cliques. Let $\eta = \max\{m\Delta^2, m^2\}$.

Algorithm 4: NSAMP-Type I

```
Initialization: Set r_1, r_2, r_3 to \phi, \kappa_1 to \emptyset, and c_1, c_2 to 0.
Edge e_i arrives;
if coin(1/i) = "head" then
 | r_1 = e_i; c_1 = 0; \kappa_1 = \{r_1\};
else
    if e_i is adjacent to r_1 then
         Increment c_1
         if coin(1/c_1) = "head" then
             r_2 = e_i; c_2 = 0; \kappa_1 = \{r_1, r_2\};
         else
              if e_i, r_1, and r_2 form a triangle then
                 \kappa_1 = \kappa_1 \cup \{e_i\};
              else
                  if e_i is adjacent to r_1 or r_2 then
                       c_2 = c_2 + 1;
                       if coin(1/c_2) = "head" then
                        | r_3 = e_i; \kappa_1 = \kappa_1 \cup \{r_3\}
                       else
                        | If e_i is an edge connecting two end points of r_1, r_2 or r_3, then add e_i to \kappa_1.
```

```
Algorithm 5: NSAMP-Type IIInitialization: r_1 = \phi, r_2 = \phi; \kappa_2 = \emptyset.Edge e_i arrives;If (coin(1/i) = "head") set r_1 \leftarrow e_i;If (coin(1/i) = "head") set r_2 \leftarrow e_i;If (coin(1/i) = "head") set r_2 \leftarrow e_i;If (coin(1/i) = "head") set r_2 \leftarrow e_i;If (Neither \ coin \ returned \ "head") then\downarrow \ \epsilon_i \ is \ adjacent \ to \ both \ r_1 \ and \ r_2 \ then\lfloor \ \kappa_2 = \kappa_2 \cup \{e_i\};else\downarrow \ \kappa_2 = \{r_1, r_2\}; else \kappa_2 = \{r_1\};
```

Theorem 3.3 There is a $O\left(s(\varepsilon, \delta)\frac{\eta}{\tau_4(G)}\right)$ -space bounded streaming algorithm that observes a graph G and returns $a(\varepsilon, \delta)$ approximation of the number of 4-cliques in G.

Theorem 3.4 There is a $O\left(s(\varepsilon, \delta)\frac{\eta_{\ell}}{\tau_{\ell}(G)}\right)$ -space bounded algorithm that returns an (ε, δ) approximation of the number of ℓ -cliques in a stream, where $\eta_{\ell} = \max \cup_{\alpha=1}^{\lfloor \ell/2 \rfloor} \{m^{\alpha} \Delta^{\ell-2\alpha}\}.$

For instance, if ℓ was even, and $\Delta = O(\sqrt{m})$, then the above algorithm for ℓ -cliques takes space $O\left(s(\varepsilon, \delta)\frac{m^{\ell/2}}{\tau_{\ell}(G)}\right)$ **Sampling 4-Cliques.** We show the following analogous results for sampling cliques.

Theorem 3.5 Assume that $\tau_4(G) > 0$, and let $\eta = \max\{m\Delta^2, m^2\}$. For every δ , $0 < \delta < 1$, there is a streaming algorithm with space complexity $O\left(\frac{\eta \log(1/\delta)}{\tau_4(G)}\right)$ that observes a graph G with m edges and returns a random 4-clique in G with probability at least $1 - \delta$.

Theorem 3.6 Assume that ℓ is a constant and $\tau_{\ell}(G) > 0$. There is an $O\left(\frac{\eta_{\ell}}{\tau_{\ell}}\log(1/\delta)\right)$ -space bounded algorithm that observes a graph G and returns an ℓ -clique from the graph, chosen uniformly at random from the set of all ℓ -cliques in the graph, where $\eta_{\ell} = \max \bigcup_{\alpha=1}^{\lfloor \ell/2 \rfloor} \{m^{\alpha} \Delta^{\ell-2\alpha}\}.$

4 **Experiments**

We empirically study the proposed triangle counting algorithm. We implemented the version of the algorithm which processes edges in batches, and also the other state-of-the-art algorithms due to [BFL⁺06, JG05].

4.1 Experimental Setup

Datasets and Environment. Our study uses a collection of popular social media graphs, obtained from the publicly available data provided by the SNAP project at Stanford [Les]. We present a summary of these datasets in Table 2. We remark that while these datasets stem from social media, our algorithm does not assume any special property about them. Our experiments were performed on a 2.2 Ghz Intel Core i7 laptop machine with 8GB of memory, but our experiments used no more than a few MB of RAM. The machine is running Mac OS X 10.7.5. All programs were compiled with GNU g++ version 4.2.1 (Darwin) using the flag -O3. We measure and report wall-clock time using gettimeofday, which has enough resolution for our experiments. Our prototype implementation uses GNU's STL implementation of collections, including unordered_map for hash maps; we did not attempt to optimize our code.

Our algorithm is randomized and may behave differently on different runs. Thus, for robustness, we perform *five* trials and gather the following statistics: (1) the minimum, mean, and maximum deviation (relative error) values from the true answer across the trials, (2) the median wall-clock overall runtime, and (3) the median I/O time. Mean deviation is a well accepted measure of error, which we believe to give an accurate picture of how well the algorithm performs. For completeness, we also report the min/max deviation values, but we note that as we perform more trials, the minimum becomes smaller and the maximum becomes larger, so they are not robust.

Dataset	n	m	Δ	au	$m\Delta/ au$
Amazon	334,863	925,872	1,098	667,129	1,523.85
DBLP	317,080	1,049,866	686	2,224,385	323.78
Youtube	1,134,890	2,987,624	57,508	3,056,386	56,214.20
LiveJournal	3,997,962	34,681,189	29,630	177,820,130	5,778.89
Orkut	3,072,441	117,185,083	66,626	633,319,568	12,328.02

Table 2: A summary of the datasets used in our experiments, showing for every dataset the number of nodes (n), the number of edges (m), the maximum degree (Δ), the number of triangles in the graph (τ), and the ratio $m\Delta/\tau$.

Baseline. We implemented the state-of-the-art algorithms for adjacency streams due to Buriol et al. [BFL⁺06], and Jowhari and Ghodsi [JG05]. Our implementation of Buriol et al.'s algorithm follows the description of the optimized version in their paper, which achieves O(m + r) running time for m edges and r estimators, through certain approximations. However, we were unable to obtain meaningful results from these algorithms on the datasets we consider: The estimators due to Buriol et al.fail to find a triangle most of the time, resulting in low-quality estimates, or producing no estimates at all—even when using millions of estimators (see Section 2.1 for a related discussion); this behavior is consistent with Buriol et al.'s findings for their adjacency-stream algorithm. In the case of Jowhari-Ghodsi's algorithm, its O(mr) running time is too slow even on a modest dataset. Instead, we directly compare our results with the true count and focus our study on the scalability of the approach as the graph size and the number of estimators increase. For most datasets, the exact triangle count is provided by the source; in other cases, we compute the exact count using an algorithm developed as part of the Problem-Based Benchmark Suite [SBF⁺12].

4.2 Accuracy

The first set of experiments aims to study the accuracy of our estimates on different datasets. Our theoretical results predict that as the number of estimators r increases, so does the accuracy. We are interested in verifying this prediction, as well as studying the dependence of the accuracy on parameters such as the number of edges m, maximum degree Δ and the number of triangles τ .

Dataset	$r = 1 \mathrm{K}$		$r = 128 \mathrm{K}$		r = 1M		
	min/mean/max dev.	Time	min/mean/max dev.	Time	min/mean/max dev.	Time	
Amazon	1.60 / 6.28 / 12.45	0.41	0.11 / 0.84 / 1.52	1.06	0.08 / 0.25 / 0.40	3.72	0.26
DBLP	8.04 / 18.28 / 36.53	0.45	0.08 / 0.50 / 0.97	1.08	0.07 / 0.19 / 0.42	3.90	0.28
Youtube	12.56 / 59.45 / 79.76	1.25	9.37 / 21.46 / 38.49	2.39	1.75 / 4.42 / 10.18	5.26	0.79
LiveJournal	0.24 / 11.53 / 29.76	15.00	1.41 / 2.35 / 4.02	23.10	0.19 / 0.60 / 1.45	33.40	10.00
Orkut	4.61 / 31.93 / 58.93	52.40	2.13 / 4.69 / 12.69	75.20	1.48 / 3.55 / 5.93	103.00	33.40

Table 3: The accuracy (min/mean/max deviation in percentage), median total running time (in seconds), and I/O time (in seconds) of our bulk algorithm across five runs as the number of estimators r is varied.

Table 3 shows the median total running time, accuracy (showing min., mean, and max. relative errors in percentage), and median I/O time of our algorithm across five runs as we vary the number of estimators r (1024, 131072, 1047576). We show the I/O time for each dataset as it makes up of a non-negligible fraction of the total running time. Several things are clear from this experiment. First, *our algorithm is accurate with only a modest number of estimators*. In all but the Youtube dataset, the algorithm achieves less than 5% mean deviation using only about 100 thousand estimators. Furthermore, the accuracy significantly improves as we increase the number of estimators r to 1M. Second, *a high degree graph with few triangles needs more estimators*. Consistent with the theoretical findings, Youtube and Orkut, which have the largest $m\Delta/\tau(G)$ values, need more estimators than others to reach the same accuracy. Third, but perhaps most importantly, *in practice, far fewer estimators than suggested by the pessimistic theoretical bound is necessary to reach a desired accuracy*. For example, on Orkut, using $\varepsilon = 0.0355$, the expression $s(\varepsilon, \delta)m\Delta/\tau$ is at least 9.78 million, but we already get this accuracy using 1 million estimators.

4.3 Performance

In the second set of experiments, we selected two graphs—Youtube and LiveJournal—to study runtime performance and accuracy in more detail. The goal of these experiments is to understand how the number of estimators affects the running time, as well as the accuracy.

First, we consider running time. It is clear from Figure 2 that the running time increases with the number of estimators r, as expected. The theory predicts that the running time is O(m + r); that is, it scales linearly with r. This is hard to confirm visually since we do not know how much of it is due to the O(m) term; however, in both cases, we are able to compute a value t_0 such that the running times minus t_0 scale roughly linearly with r.



Figure 2: Running time (in seconds) as the number of estimators r is varied (r = 1K, 2K, ..., 8M). The x-axis is in log scale.



Figure 3: Mean deviation (**in percentage**) of our estimates as the number of estimators r is varied ($r = 1K, 2K, \dots, 8M$). The *x*-axis is in log scale.

As for accuracy, (ε, δ) -type approximation algorithms are difficult to experiment with, in general: the observed error could stem from the controlled error term ε or it could stem from the unlikely event that happens with probability δ where we have no control over the error. The general trend from Figure 3 is that the mean deviation decreases as we increase the number of estimators.

5 Extensions

Our basic algorithms for sampling and counting can be extended to the case the case of sliding windows. For simplicity, we consider sequence-based sliding window, where the scope of relevant data is restricted to the w most recent edges. Thus, we would like to estimate the number of triangles in the graph induced by the most recent w edges. There are several algorithms to sample an element from a sliding window [BOZ09, BDM02, Haa, ZLY⁺05, GL08] that can be adapted for use here. For simplicity, we use the algorithm from [BDM02]. Recall that the neighborhood sampling algorithm maintains two edges r_1 —a randomly chosen edge, and r_2 —a randomly chosen edge from $N(r_1)$. At time instance t, let $\langle e_{t-w+1}, e_{t-w+2}, \cdots, e_{t-1}, e_t \rangle$ denote the last w edges seen. For each edge e_i , we pick a random number $\rho(i)$ chosen uniformly between 0 and 1. We maintain a chain of samples $S = \{e_{\ell_1}, e_{\ell_2}, \cdots, e_{\ell_k}\}$ from the current window. The first edge e_{ℓ_1} is the edge such that $\rho(\ell_1) = \min\{\rho(t-w+1), \cdots, \rho(t)\}$. For $2 \le i \le k$, e_{ℓ_i} is the edge such that $\rho(\ell_i) = \min\{\rho(\ell_{i-1}+1), \cdots, \rho(t)\}$. For each $e_{\ell_i} \in S$, we also maintain a random adjacent neighbor r_2^i from $N(e_{\ell_i})$. Note that the second edge can be chosen using standard reservoir sampling, because, if e_{ℓ_i} lies in the current window, any neighbor that arrives after it will also be in the current window. Thus all of r_2^i s also belong to the current window. We chose r_1 to be e_{ℓ_1} and r_2 to r_2^1 . When r_1 falls out of window, we remove it from S, and update r_1 to e_{ℓ_2} and r_2 to r_2^2 , and so on. This will ensure that r_1 is always a random edge in the current window and r_2 is a random neighbor of r_1 from the current window, and the rest of the analysis follows. It is easy to see that that the expected size of the set S is $\Theta(\log w)$ [BDM02]. Thus, the total expected space used by the algorithm increases by a factor of $\log w$.

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A Appendix: Improved Space Bound For Triangle Counting

In this section, we prove a sharper bound on the space needed to obtain an accurate estimate. We will define a measure that captures in the amount of interaction between triangles and non-triangle edges in the graph. The *tangle coefficient* of a graph G, denoted by γ is given by

$$\gamma := \frac{1}{\tau(G)} \sum_{t' \in \mathcal{T}(G)} C(t').$$

This is equivalent to $\gamma = \frac{1}{\tau(G)} \sum_{e \in E} c(e)s(e)$, where s(e) counts the number of triangles $t' \in \mathcal{T}(G)$ such that the first edge of t' is e. Using the tangle coefficient, we prove the following theorem (we provide some discussions about the tangle coefficient after that):

Theorem A.1 (Improved Triangle Counting) Let γ be the tangle coefficient of G. There is an (ε, δ) -approximation to the triangle counting problem that requires $O(1/\varepsilon^2 \cdot m\gamma/\tau(G) \cdot \log(\frac{1}{\delta}))$.

Proof: (Sketch) Each estimator has variance $m \sum_{t \in \mathcal{T}(G)} C(t) = m\tau(G)\gamma$. We will run $\alpha = 4/\varepsilon^2 \cdot \gamma m/\tau(G)$ independent parallel copies of such an estimator. Let the average of these estimates be Y. By Chebyshev's inequality, we have

$$\Pr\Big[|Y - \mathbf{E}[Y]| > \varepsilon \cdot \tau(G)\Big] \le \frac{1}{4}.$$

To boost the success probability up to $1 - \delta$, we will run $\beta = 12 \ln(1/\delta)$ independent copies of Y estimators and take the median. Note that our median estimator fails to produce an ε -approximation only if more than $\beta/2$ fails to produce an ε -approximation. In expectation, the number of "failed" estimators is at most $\beta/4$. Therefore, by a standard Chernoff bound, we fail with probability at most

$$\mathbf{Pr}[\mathsf{FAILED}] \le e^{-\frac{1^2(\beta/4)}{3}} = \delta,$$

proving our final estimate is an (ε, δ) -approximation using space at most $\alpha\beta$.

Clearly, γ is at most 2Δ , recovering the original bound we proved. We can gain more understanding of the quantity γ by considering the following random process: Fix a stream and pick a random triangle from this graph. If *e* is the first edge in the stream of this triangle, then the value of γ is the number of edges that incident on *e* that comes after it. In this view, γ can be seen as a measure of how entangled the triangles in this stream are—as our intuition suggests, if the triangles "interact" with many non-triangle triples, we will need more space to obtain an accurate answer.

B Bulk-Processing Edges

Our algorithms take advantage of a procedure unif for generating a uniform random number from the interval [0, 1] as well as a procedure randInt(n) for drawing a number uniformly at random from $\{0, \ldots, n-1\}$. We assume these take constant time.

We now give detailed descriptions; the bookkeeping, though conceptually simple, turns out to be quite messy. For a block B, computing the level-1 edge after incorporating B is straightforward. Let i_B be the total number of edges so far (including the edges in B). An edge in B will replace an existing level-1 edge with probability b/i_B —and when this happens, each edge in B has an equal probability of being a replacement edge. Thus, with one call to unif, we can compute the level-1 edge after incorporating B, so in O(r) time, we can determine the level-1 edges for all r estimators we maintain. This can be further optimized by noticing that in later stages, the number of estimators that end up updating their level-1 edges are progressively smaller. In this case, our task boils down to generating a binary vector where the probability p that a position is 1 is small. We can generate this vector by generating a few geometric random variables representing the gap between 1's in the vector. Since we expect only a p-th fraction of the estimators to be updated, this is more efficient than going over all the estimators. Using this technique, the cost of updating level-1 across the whole stream is at most $O(r \log(m/w))$.

The challenge in fastforwarding over the block B lies in maintaining level-2 edges, finding the completed triangles, and keeping track of the counters. But these, too, can be maintained efficiently if we maintain some additional information and perform an extra pass through the block.

We first address the problem of maintaining the count c; the information we gather here will also be useful for updating level-2 edges. At this point, we have determined the level-1 edges for all r estimators. Let R be the set of the endpoints of these edges. We will maintain an integer counter for each node in R, requiring O(r) space. Further, we augment the state of each estimator est with the following fields: est. d_1 , est. d_2 .

Pass 1. We will make a pass through *B*. As we go through the edges $b_1, b_2, \ldots, b_{|B|}$, we keep a count λ_w of the edges incident on *w* so far. For an estimator est with level-1 edge $e = \{x, y\}$, we record est $d_1 = \lambda_x$ and est $d_2 = \lambda_y$ when seeing *e* in *B* and when we finish the pass, the count est *c* is incremented by $\lambda_x - \text{est.} d_1 + \lambda_y - \text{est.} d_2$, where λ_x, λ_y are the values at the end of the pass. So far, we require one pass and O(|B| + |R|) = O(|B| + r) time and space using a (hash) table which maps level-1 edges to their corresponding estimator states and a (hash) table storing the degree values.

Conveniently, the λ values are also useful for updating level-2 edges. Consider an estimator est with level-1 edge $e = \{x, y\}$. The information we gather in the first pass provides us with clues about which edge to pick as a level-2 edge. We know that the number of edges in *B* incident to *x* that come after *e* is exactly $\alpha = \lambda_x - \text{est.} d_1$. Similarly, the number of edges in *B* incident to *y* that come after *e* is exactly $\beta = \lambda_y - \text{est.} d_2$. Therefore, the probability that an edge from *B* will replace the current level-2 edge is exactly $(\alpha + \beta)/\text{est.} c$. At this point, we can number the edges incident on *x* with numbers $\{0, 1, \ldots, \alpha - 1\}$ and the edges incident on *y* with numbers $\{0, 1, \ldots, \beta - 1\}$. Picking a random level-2 edge boils down to flipping a coin to decide whether to replace the orignal edge and if so, pick a random number between 0 and $\alpha + \beta - 1$ corresponding the replacement edge.

Our task now is simply to recognize the selected edge when we perform the second pass. This is where the second ingredient comes in: the event "the edge that increments the λ_x to a certain value z" uniquely identifies an edge in B. Given the degree information obtained previously, we can describe our random selection of a level-2 edge in this form. Once these decisions are made, we store them a (hash) table mapping (vertex, desired-restricted-degree) to the corresponding estimator

Pass 2. A second pass through the data allows us to recognize level-2 replacement edges as well as their corresponding estimators that need to be updated. Since the table contains at most O(r) entries, the space and time usage for this pass is O(|B| + r). The second pass has another important role: we can recognize the edges that will complete the triangles partially formed by level-1 and 2 edges. For this, we keep a (hash) table for queries of the form: "*if an edge e is seen, we have a complete triangle for estimator* X." This table is updated as we discover a new level-2 edge, and it is important that we perform our pass sequentially in the same order as we did in pass 1. Note that the size of this table is, again, at most O(r). Hence, each block requires two passes, which can be completed in O(|B| + r) space and time.

We can be more precise about our runtime bounds. Let C_1 and C_2 be constants independent of r. Using the optimized level-1 processing, the running time of our bulk-processing algorithm is at most

$$C_1 r(2 + H_{m/w}) + C_2 (1 + \frac{m}{w})(w + r).$$

C Omitted Proof from Section 2

We first state a measure concentration bound that will be used in the proofs.

Theorem C.1 (A Chernoff Bound) Let $\lambda > 0$ and $X = X_1, \ldots, X_n$, where each X_i , $i = 1, \ldots, n$, is independently distributed in [0, 1]. Then,

$$\mathbf{Pr}\Big[X \ge (1+\lambda)\,\mathbf{E}[X]\Big] \le e^{-\frac{\lambda^2}{2+\lambda}\cdot\mathbf{E}[X]} \quad and \quad \mathbf{Pr}\Big[X \ge (1-\lambda)\,\mathbf{E}[X]\Big] \le e^{-\frac{\lambda^2}{2}\cdot\mathbf{E}[X]}.$$

Proof of Theorem 2.3: Let $\alpha = \frac{6}{\varepsilon^2} \frac{m\Delta}{\tau(G)} \log\left(\frac{2}{\delta}\right)$. We show that the average of α independent unbiased estimators from Algorithm 2 is an (ε, δ) -approximation. For $i = 1, ..., \alpha$, let X_i be the value returned by the *i*-th estimator. Let $\bar{X} = \frac{1}{\alpha} \sum_{i=1}^{\alpha} X_i$ denote the average of these estimators. Then, as a direct consequence of Lemma 2.2, we have $\mathbf{E}[X_i] = \tau(G)$ and $\mathbf{E}[\bar{X}] = \tau(G)$. Further, for $e \in E$, we have $c(e) \leq 2\Delta$, so know $X_i \leq 2m\Delta$. Let $Y_i = X_i/(2m\Delta)$ so that $Y_i \in [0, 1]$. By letting $Y = \sum_{i=1}^{\alpha} Y_i$, we have $\mathbf{E}[Y] = \alpha \tau(G)/(2m\Delta)$; thus, by Chernoff bound (Theorem C.1),

$$\mathbf{Pr}\big[\bar{X} > (1+\varepsilon)\,\mathbf{E}[X]\big] = \mathbf{Pr}\big[\sum_{i} Y_i > (1+\varepsilon)\,\mathbf{E}[Y]\big] \le e^{-\frac{\varepsilon^2}{3}\,\mathbf{E}[Y]} = e^{-\frac{\varepsilon^2}{3}\frac{\mathbf{\alpha}\cdot\boldsymbol{\tau}(G)}{2m\Delta}} = \delta/2.$$

Similarly, we can show that $\Pr[X < (1 - \varepsilon) \mathbb{E}[X]] \le \delta/2$. Hence, with probability at least $1 - \delta$, the average \overline{X} approximates the true count within $1 \pm \varepsilon$. Since each estimator only takes O(1) space, the total space is $O(\alpha)$.

Proof of Lemma 2.7: We use a reduction from the index problem from communication complexity: Alice is given a bit vector $x \in \{0, 1\}^n$ and Bob is given an index $k \in \{1, 2, ..., n\}$, and wants to compute x_k , the bit in the kth position in x. It is known that in the model where Alice can send exactly one message to Bob, the communication cost of a randomized protocol is $\Omega(n)$ bits (see Chapter 4.2 in [KN97]).

Suppose there is a streaming algorithm \mathcal{A} that estimates the number of triangles. We can use this algorithm to solve the index problem as follows. Given a bit vector $x \in \{0,1\}^n$, Alice constructs a graph G^* on 3(n + 1) vertices with the vertex set $\{a_0, a_1, \dots, a_n\} \cup \{b_0, \dots, b_n\} \cup \{c_0, \dots, c_n\}$. Alice places three edges among a_0, b_0, c_0 to form a triangle. For each $i \in \{1, 2, \dots, n\}$, Alice places the edge (a_i, b_i) if and only if x_i equals 1. Alice processes this graph using \mathcal{A} and sends the state of the algorithm to Bob, who continues the algorithm using the state sent by Alice, and adds the two edges (b_k, c_k) and (c_k, a_k) . By querying the number of triangles in this graph with relative error of smaller that 0.5, Bob can distinguish between the following cases: (1) G^* has two triangles, and (2) G^* has one triangle. In case (1), $x_k = 1$ and in Case (2), $x_k = 0$, and hence Bob has solved the index problem.

It follows that the memory used by the streaming algorithm at Alice must be $\Omega(n)$ bits. Note that the graph G^* sent by Alice has no triples with two edges between them, and hence $O(1 + T_2(G^*)/\tau(G^*)) = O(1)$. We note that a similar proof applies to triangle sampling also.

D Omitted Proofs from Section 3

Before we present proofs, we extend the notion of neighborhood to a set of edges. Let f_1 and f_2 be two edges of S so that f_2 arrives after f_1 . Let f be an edge that is adjacent to both f_1 and f_2 . Note that such an edge may not exist. The neighborhood of f_1 and f_2 , denote $N(f_1, f_2)$, is the following set:

$$N(f_1, f_2) = \{ e \in N(f_1) \mid e \text{ arrives after } f_2 \} \cup N(f_2) - \{ f \}.$$

Let $c(f_1, f_2)$ denote the cardinality of $N(f_1, f_2)$.

Proof of Lemma 3.1.: Since κ^* is a Type I clique, f_1 and f_2 are adjacent to each other, and they together fix 3 vertices of the clique. The edge f_3 is adjacent to either one or both of f_1 and f_2 . We consider the case when f_3 is adjacent to only one of f_1 or f_2 , but not both. The other case can be handled similarly. Note that κ equals κ^* when the following events are all true: (1) \mathcal{E}_1 : f_1 equals r_1 , (2) \mathcal{E}_2 : f_2 equals r_2 , (3) \mathcal{E}_3 : f_3 equals r_3 .

Since r_1 is chosen uniformly at random among all possible m edges, the probability of \mathcal{E}_1 is 1/m. Since r_2 is an edge that is chosen uniformly at random from $N(r_1)$, $\Pr[\mathcal{E}_2 | \mathcal{E}_1] = \frac{1}{c(f_1)}$. Finally note that r_3 is chosen uniformly at random from $N(r_1, r_2)$. Thus $\Pr[\mathcal{E}_3 | \mathcal{E}_1, \mathcal{E}_2] = \frac{1}{c(f_1, f_2)}$. Thus the probability that κ equals κ^* is $\frac{1}{m} \cdot \frac{1}{c(f_1)} \cdot \frac{1}{c(f_1, f_2)}$, as desired.

Proof of Lemma 3.2: Suppose that the edges of κ^* in the stream order were f_1, f_2, \ldots, f_6 respectively. We note that $\kappa_2 = \kappa^*$ if and only if at the end of observation, $r_1 = f_1$ and $r_2 = f_2$ in Algorithm 5. To see these, note that if at

the end of observation, $r_1 = f_1$ and $r_2 = f_2$, then $\kappa_2 = \kappa^*$, since the edges f_3, \ldots, f_6 will certainly be added to κ_2 by the algorithm. The converse can also be easily seen to be true. The proof follows due to the face that the events $r_1 = f_1$ and $r_2 = f_2$ are independent, and each has a probability of 1/m of being true.

Before we present proof Theorem 3.3, we will first describe algorithms that estimate the number of Type I and Type II cliques.

Algorithm 6: COUNT-Type I
Run Algorithm 4 and let κ_1 and c_1, c_2 be the variables it maintains.
If κ_1 is a 4-Clique, then return $c_1 \cdot c_2 \cdot m$, else return 0.

The following states that Algorithm 6 returns an unbiased estimator form $\tau_4^1(G)$.

Lemma D.1 Let X denote the random variable returned by Algorithm 6 after the graph G has been observed. Then $\mathbf{E}[X] = \tau_4^1(G)$.

Proof: Note that Algorithm 6 returns a non-zero value only when κ_1 is a Type I clique. For a Type I clique κ^* let $f_1, f_2, f_3, f_4, f_5, f_6$ be the edges in the order of arrival. First, we note that if $\kappa = \kappa^*$, it must be true that $c_1 = c(f_1)$ and $c_2 = c(f_1, f_2)$.

$$\begin{split} \mathbf{E}[X] &= \sum_{\kappa^* \in S_1} \mathbf{E}[X \mid \kappa = \kappa^*] \cdot \Pr[\kappa = \kappa^*] = \sum_{\kappa^* \in \mathcal{T}_4^1(G)} \Pr[\kappa = \kappa^*] \cdot m \cdot c_1 \cdot c_2 \\ &= \sum_{\kappa^* \in \mathcal{T}_4^1(G)} \Pr[\kappa = \kappa^*] \cdot m \cdot c(f_1) \cdot c(f_1, f_2) \\ &= \sum_{\kappa^* \in \mathcal{T}_4^1(G)} \frac{1}{m} \cdot \frac{1}{c(f_1)} \cdot \frac{1}{c(f_1, f_2)} \cdot m \cdot c(f_1) \cdot c(f_1, f_2) (\text{ By Lemma 3.1}) \\ &= \tau_4^1(G) \end{split}$$

The next algorithm estimates the number of Type II cliques.

Algorithm 7: COUNT-Type II	
Run Algorithm 5 and let κ_2 be the variable it maintains.	ntains.
If κ_2 is a 4-Clique, then return m^2 , else return 0.	

Lemma D.2 Let X denote the random variable returned by Algorithm 7. Then $\mathbf{E}[X] = \tau_4^2(G)$.

The proof of the above lemma is similar to the proof of Lemma D.1, we omit the proof. Now we are ready to describe the algorithm to estimate the number of 4-cliques and prove Theorem 3.3

Proof of Theorem 3.3: Algorithm 8 is our estimator of number of 4-cliques. The correctness of the algorithm, space and accuracy bounds follow from Lemma D.1 and Lemma D.2 and applying a Chernoff bound on the resulting estimates.

By running $O(Dm \log(1/\delta))$ repetitions of Algorithm 9, we obtain Theorem 3.5, where $D = \max{\{\Delta^2, m\}}$.

Algorithm 8: COUNT-CLIQUES

Result: An (ε, δ) estimate for $\tau_4(G)$ $N \leftarrow \max\{6m\Delta^2, m^2\}\frac{\log(1/\delta)}{\tau_4(G)\varepsilon^2}$; $X_1 \leftarrow 0, X_2 \leftarrow 0$; Run N independent copies of Algorithm 6 and of Algorithm 7; Set X_1 to the mean of all values returned by Algorithm 6; Set X_2 to the mean of all values returned by Algorithm 7; Return $X_1 + X_2$.

Algorithm 9: Sample from $\mathcal{T}_4(G)$

Run in parallel Algorithms 4, and 5. Let κ_1, c_1, c_2 be the state maintained by Algorithm 4 and κ_2 be the state maintained by Algorithm 5. **Query arrives for a random Clique** Uniformly at random pick $b \in \{1, 2\}$; If b = 1, then if κ_1 is a 4-clique, return κ_1 with probability $\frac{c_1c_2}{D}$. If b = 2, then if κ_2 is a 4-clique, return κ_2 with probability m/D. Return "Fail"

Lemma D.3 For any 4-clique in G, say κ^* , the probability that Algorithm 9 returns κ^* is $\frac{1}{2mD}$. The probability that the algorithm returns a 4-clique is $\frac{\tau_4(G)}{2mD}$.

Proof: Suppose that κ^* was a Type I clique, and suppose its first two edges in the stream order were f_1 , f_2 respectively. Algorithm 9 returns κ^* if: (1) \mathcal{E}_1 :*b* is chosen to be 1 by the algorithm, and (2) \mathcal{E}_2 : κ^* is chosen by κ_1 , and (3) \mathcal{E}_3 : κ^* is finally returned by the algorithm in the final step, with prob. $c(f_1)c(f_1, f_2)/m\Delta$.

We know from Lemma 3.1 that $\Pr[\mathcal{E}_2] = \frac{1}{mc(f_1)c(f_1,f_2)}$. Thus,

$$\Pr[\mathcal{E}_1 \cap \mathcal{E}_2 \cap \mathcal{E}_3] = \frac{1}{2} \cdot \frac{1}{mc(f_1)c(f_1, f_2)} \cdot \frac{c(f_1)c(f_1, f_2)}{D} = \frac{1}{2mD}$$

It is possible to similarly show that the probability of returning each Type II clique is also $\frac{1}{2mD}$. Since the events of returning different cliques are all disjoint, the probability that Algorithm 9 returns some 4-clique is $\frac{\tau_4(G)}{2mD}$.