A Linear-time Algorithm for Sparsification of Unweighted Graphs

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Abstract

Given an undirected graph $G$ and an error parameter $\varepsilon > 0$, the graph sparsification problem requires sampling edges in $G$ and giving the sampled edges appropriate weights to obtain a sparse graph $G_\varepsilon$ with the following property: the weight of every cut in $G_\varepsilon$ is within a factor of $(1 \pm \varepsilon)$ of the weight of the corresponding cut in $G$. If $G$ is unweighted, an $O(m \log n)$-time algorithm for constructing $G_\varepsilon$ with $O(n \log n / \varepsilon^2)$ edges in expectation, and an $O(m)$-time algorithm for constructing $G_\varepsilon$ with $O(n \log^2 n / \varepsilon^2)$ edges in expectation have recently been developed [9]. In this paper, we improve these results by giving an $O(m)$-time algorithm for constructing $G_\varepsilon$ with $O(n \log n / \varepsilon^2)$ edges in expectation, for unweighted graphs. Our algorithm is optimal in terms of its time complexity; further, no efficient algorithm is known for constructing a sparser $G_\varepsilon$. Our algorithm is Monte-Carlo, i.e. it produces the correct output with high probability, as are all efficient graph sparsification algorithms.
1 Introduction

A cut of an undirected graph is a partition of its vertices into two disjoint sets. The weight of a cut is the sum of weights of the edges crossing the cut, i.e., edges having one endpoint in each of the two vertex subsets of the partition. For unweighted graphs, each edge is assumed to have unit weight. Cuts play an important role in many problems in graphs; e.g., the maximum flow between a pair of vertices is equal to the minimum weight cut separating them.

A skeleton \( G' \) of an undirected graph \( G \) is a subgraph of \( G \) on the same set of vertices where each edge in \( G' \) can have an arbitrary weight. The problem of finding an appropriately weighted sparse skeleton for an undirected graph \( G \) that approximately preserves the weights of all cuts in \( G \) was introduced and studied by Karger et al. in a series of results [11, 12, 3] culminating in the following theorem. Throughout this paper, for any undirected graph \( G \) and any \( \varepsilon \in (0, 1] \), \( (1 \pm \varepsilon)G \) will denote the set of all appropriately weighted subgraphs of \( G \) where the weight of every cut in the subgraph is within a factor of \( (1 \pm \varepsilon) \) of the weight of the corresponding cut in \( G \).

**Theorem 1** (Benczúr-Karger [3]). For any undirected graph \( G \) with \( m \) edges and \( n \) vertices, and for any error parameter \( \varepsilon \in (0, 1] \), a skeleton \( G_\varepsilon \) containing \( O(n \log^2 n/\varepsilon^2) \) edges in expectation such that \( G_\varepsilon \subseteq (1 \pm \varepsilon)G \) with high probability [9] can be found in \( O(m \log^2 n) \) time if \( G \) is unweighted and \( O(m \log^3 n) \) time otherwise.

Besides its combinatorial ramifications, the importance of this result stems from its use as a pre-processing step in several graph algorithms, e.g., to obtain an \( \tilde{O}(n^{3/2} + m) \)-time algorithm for approximate maximum flow using the \( \tilde{O}(m^{3/2}) \)-time algorithm for exact maxflow due to Goldberg and Rao [7]; and more recently, \( \tilde{O}(n^{3/2} + m) \)-time algorithms for approximate sparsest cut [15, 19].

Subsequent to Benczúr and Karger’s work, Spielman and Teng [21, 22] extended their results to preserving all quadratic forms, of which cuts are a special case; however, the size of the skeleton constructed was \( O(n \log^c n) \) for some large constant \( c \). Spielman and Srivastava [20] improved this result by constructing skeletons of size \( O(n \log n/\varepsilon^2) \) in \( O(m \log O(1) n) \) time, while continuing to preserve all quadratic forms. Recently, this result was further improved by Batson et al. [2] who gave a deterministic algorithm for constructing skeletons of size \( O(n/\varepsilon^2) \) that preserve the weights of all cuts whp. While their result is optimal in terms of the size of the skeleton constructed, the time complexity of their algorithm is \( O(m \log n/\varepsilon^2) \), rendering it somewhat useless in terms of applications.

Recently, further progress has been made on efficiently constructing a skeleton graph in the form of the following theorem due to Hariharan and Panigrahi [9].

**Theorem 2** (Hariharan-Panigrahi [9]). For an undirected graph \( G \) with \( m \) edges and \( n \) vertices, and for any error parameter \( \varepsilon \in (0, 1] \), the following algorithmic results can be obtained for constructing a skeleton graph \( G_\varepsilon \) that is in \((1 \pm \varepsilon)G\) with high probability:

- If the expected number of edges in \( G_\varepsilon \) is \( O(n \log^2 n/\varepsilon^2) \), then \( G_\varepsilon \) can be constructed in \( O(m) \) time if \( G \) has polynomial edge weights and \( O(m \log^2 n) \) time if \( G \) has arbitrary edge weights.

- If the expected number of edges in \( G_\varepsilon \) is \( O(n \log n/\varepsilon^2) \), then \( G_\varepsilon \) can be constructed in \( O(m \log n) \) time if \( G \) is unweighted, and \( O(m \log^2 n) \) time if \( G \) has polynomial edge weights.

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\(^1\) We say that a property holds with high probability (or whp) for a graph on \( n \) vertices if its failure probability can be bounded by the inverse of a fixed polynomial in \( n \).
Combining the above two results, one can obtain an algorithm to construct a skeleton graph $G_ε$ that preserves the weights of all cuts whp and has $O(n \log n/ε^2)$ edges in expectation in $O(m + n \log^4 n/ε^2)$ time if $G$ has polynomial edge weights.

A natural conclusion for this line of work would be to obtain an $O(m)$-time algorithm for constructing a skeleton graph $G_ε$ containing $O(n \log n/ε^2)$ edges in expectation. In this correspondence, we obtain this result in the form of the following theorem if $G$ is unweighted. It may be noted that even if $G$ is unweighted, the best sparsification result known previously was Theorem[2].

**Theorem 3.** For an undirected unweighted graph $G$ with $m$ edges and $n$ vertices, and for any error parameter $ε \in (0, 1]$, a skeleton graph $G_ε$ that is in $(1 \pm ε)G$ with high probability and has $O(n \log^2 n/ε^2)$ edges in expectation, can be constructed in $O(m)$ time.

Note that the above algorithm is optimal in terms of its running time, and there is no efficient algorithm known for constructing a sparser skeleton, even for unweighted graphs. (As mentioned previously, the only algorithm known that constructs a sparser skeleton has a time complexity of $O(n^3 m/ε^2)$.)

Before describing our algorithm in more detail, it is worth mentioning some of the related research in graph sparsification. In a recent result, Fung and Harvey[5] show that sampling uniformly random spanning trees of a graph produces good sparsifiers. This approach was used previously to obtain coarser sparsifiers by Goyal et al[8]. Fung and Harvey also show that sampling edges according to their standard connectivities also produces good sparsifiers, a result obtained independently by Hariharan and Panigrahi[9]. The problem of graph sparsification in the semi-streaming model was first considered by Ahn and Guha[1] who gave a one-pass algorithm for constructing a skeleton $G_ε$ containing $O(n \log n \log(m/n)/ε^2)$ edges. Recently, Goel et al have given the following algorithms for this problem[6]:

- An $O(m \log \log n)$-time one-pass algorithm for constructing a skeleton graph $G_ε$ containing $O(n \log^2 n/ε^2)$ edges in expectation. The size of the skeleton can be improved to $O(n \log n/ε^2)$ edges; however, the time complexity of the algorithm then becomes $O(m \log \log n + n \log^5 n/ε^2)$.

- An $O(m)$-time two-pass algorithm for constructing a skeleton graph $G_ε$ containing $O(n \log n/ε^2)$ edges in expectation, if $m = Ω(n^{1+δ})$ for some constant $δ > 0$.

Observe that both results, if applied to a non-streaming model, are weaker than Theorem[2]. Another area of recent interest, though not directly related to our problem, is that of vertex sparsification[15][14]. Given a graph $G = (V, E)$ and a subset of vertices $S \subset V$, the goal here is to create a graph $G_S = (S, E_S)$ that approximately preserves some desired connectivity property of $G$ (e.g. minimum steiner cut[15], maximum multi-commodity flow[14]).

### 1.1 Our Techniques

All previous algorithms for graph sparsification have two phases: in the first phase, a suitable probability $p_e$ for sampling each edge $e$ is determined; and then, in the second phase, every edge is independently sampled with probability $p_e$ and given weight $1/p_e$ in the skeleton graph if selected in the sample. Our main technical novelty is in interleaving the sampling process with that of estimating sampling probabilities. Such interleaving leads to several technical hurdles:

- It introduces dependence between the sampling of different edges. Such dependence has appeared previously in sparsification algorithms for the semi-streaming model, but the nature of the dependence in our algorithm is somewhat different from that in the streaming algorithms.
An edge may now be sampled multiple times, and errors are accrued in each such sampling. This requires us to choose the interleaved sampling probabilities very carefully so that the errors do not add up.

At a high level, our algorithm has the same iterative structure as algorithms in [3] and [9]. In each iteration, the algorithm identifies suitable sampling probabilities of a subset of edges and removes them from the graph. It is in what the algorithm does with the remaining edges that our algorithm differs from previous work. While all remaining edges are retained for the next iteration in previous algorithms, we sample all the edges with probability 1/2 and retain only half of them in expectation for the next iteration. The intuition behind this sampling comes from the observation that the sampling probabilities decrease (approximately by a factor of 2) with every iteration; therefore, a natural approach is to sample the remaining edges with probability 1/2 and retain only the selected edges thereby reducing the time complexity of the next iteration.

Suppose $X_i$ be the set of remaining edges at the beginning of iteration $i$, $F_i$ be the set of edges whose sampling probabilities are determined in iteration $i$ and $Y_i = X_i \setminus F_i$ be the set of remaining edges after iteration $i$. (Note that $X_{i+1}$ is therefore constructed by sampling each edge in $Y_i$ with probability 1/2.) Our proof technique consists of two parts. First, we show that the graph $S$ containing appropriately weighted edges in $\cup_i F_i$ is in $(1 \pm \varepsilon/3)G$ whp, i.e. even though edges in $Y_i \setminus X_{i+1}$ are sampled out between iterations $i$ and $i+1$ for each $i$, the retained edges (when weighted appropriately) are sufficient to preserve all cuts. In the second step of the proof, we show that the skeleton graph $G_e$ constructed by sampling edges in $\cup_i F_i$ and giving them appropriate weights is in $(1 \pm \varepsilon/3)S$ whp. For this proof, we use the generic sparsification framework developed recently by Hariharan and Panigrahi [9]. Combining these two steps, we conclude that $G_e \in (1 \pm \varepsilon)G$ whp.

Roadmap. In section 2 we give an outline of the generic sampling framework from [9] that we use later in our proof. In section 3 we describe our sparsification algorithm, prove its correctness and derive its time complexity. Finally, we conclude with some open questions in section 4.

2 Preliminaries

We first need to introduce the notion of $k$-heavy edges, for any $k > 0$.

**Definition 1.** An edge $e = (u,v)$ of an undirected graph $G = (V,E)$ is said to be $k$-heavy if the maximum flow between vertices $u$ and $v$ in $G$ is at least $k$.

By Menger’s theorem (see e.g., [4]), it follows that if $e = (u,v)$ is $k$-heavy, then the weight of every cut in $G$ having $u$ and $v$ on different sides is at least $k$.

2.1 Outline of Sparsification Framework from [9]

Suppose $G = (V,E)$ is an undirected graph where edge $e \in E$ has a positive integer weight $w_e$. Let $G_M = (V, E_M)$ denote the multi-graph constructed by replacing each edge $e$ by $w_e$ unweighted parallel edges $e_1, e_2, \ldots, e_{w_e}$. Consider any $\varepsilon \in (0,1]$. Suppose we construct a skeleton $G_e$ where each edge $e \in E_M$ is present in graph $G_e$ independently with probability $p_e$, and if present, it is given a weight of $1/p_e$. Let $p_e = \min\left(\frac{\varepsilon}{32\alpha_n}, 1\right)$, where $\alpha$ is independent of $e$ and $\lambda_e$ is some parameter of $e$ satisfying $\lambda_e \leq 2^{n-1} - 1$. The authors describe a sufficient condition that characterizes a good choice of $\alpha$ and $\lambda_e$’s.

To describe this sufficient condition, partition the edges in $G_M$ according to the value of $\lambda_e$ into sets $R_0, R_1, \ldots, R_K$ where $K = \lceil \log_{\lambda_e} (\lambda_e) \rceil \leq n - 1$ and $e_i \in R_j$ iff $2^j \leq \lambda_e \leq 2^{j+1} - 1$. Now, let $Q = $
\( (Q_0, Q_1, Q_2, \ldots, Q_i) \) be a sequence of subgraphs of \( G_M \) (edges of \( G_M \) are allowed to be replicated multiple times in the \( Q_i \)'s) such that \( R_i \subseteq W_i \) for every \( i \). \( Q \) is said to be a \((\pi, \alpha)\)-certificate corresponding to the above choice of \( \alpha \) and \( \lambda_e \)'s if the following properties are satisfied:

**\( \pi \)-connectivity** For \( i \geq 0 \), any edge \( e_l \in R_i \) is \( \pi \)-heavy in \( Q_i \).

**\( \alpha \)-overlap** For any cut \( C \) containing \( c \) edges in \( G_M \), let \( w^{(C)}_i \) be the number of edges that cross \( C \) in \( Q_i \).

Then, for all cuts \( C, \sum_{i=0}^{k} \frac{w^{(C)}_i}{2^{i-1}} \leq \alpha c. \)

Then, the following theorem holds.

**Theorem 4** (Hariharan-Panigrahi [9] (Theorem 8)). If there exists a \((\pi, \alpha)\)-certificate for a particular choice of \( \alpha \) and \( \lambda_e \)'s, then the skeleton \( G_e \in (1 \pm \varepsilon)G \) with probability at least \( 1 - 4/n \). Further \( G_e \) has \( O\left(\frac{\alpha \log n}{\varepsilon^2} \sum_{e \in E} \frac{w_e}{\lambda_e} \right) \) edges in expectation.

We also need the following lemma, which is a slight variation of Lemma 5 from [9]. (For completeness, we give a proof in the appendix.) For an undirected unweighted graph \( G = (V, E) \), let \( R \subseteq E \) and \( Q \supseteq R \) be subsets of edges such that \( R \) is \( \pi \)-heavy in \( (V, Q) \). Suppose each edge \( e \in R \) is sampled with probability \( p \), and if selected, given a weight of \( 1/p \) to form a set of weighted edges \( \hat{R} \). Now, for any cut \( C \) in \( G \), let \( R^{(C)}, Q^{(C)} \) and \( \hat{R}^{(C)} \) be the sets of edges crossing cut \( C \) in \( R, Q \) and \( \hat{R} \) respectively; also let the total weight of edges in \( R^{(C)}, Q^{(C)} \) and \( \hat{R}^{(C)} \) be \( r^{(C)}, q^{(C)} \) and \( \hat{r}^{(C)} \) respectively. Then the following lemma holds.

**Lemma 1.** For any \( \delta \in (0, 1] \) satisfying \( \delta^2 \cdot p \cdot \pi \geq \frac{6\log n}{0.38} \),

\[ |r^{(C)} - \hat{r}^{(C)}| \leq \delta q^{(C)} \]

for all cuts \( C \) in \( G \) with probability at least \( 1 - 4/n^2 \).

### 2.2 Nagamochi-Ibaraki Forests

We first introduce the notion of spanning forests of a graph. As earlier, \( G \) denotes a graph with integer edge weights \( w_e \) for edge \( e \) and \( G_M \) is the unweighted multi-graph where \( e \) is replaced with \( w_e \) parallel unweighted edges.

**Definition 2.** A spanning forest \( T \) of \( G_M \) (or equivalently of \( G \)) is an (unweighted) acyclic subgraph of \( G \) satisfying the property that any two vertices are connected in \( T \) if and only if they are connected in \( G \).

We partition the set of edges in \( G_M \) into a set of forests \( T_1, T_2, \ldots \) using the following rule: \( T_i \) is a spanning forest of the graph formed by removing all edges in \( T_1, T_2, \ldots, T_{i-1} \) from \( G_M \) such that for any edge \( e \in G \), all its copies in \( G_M \) appear in a set of contiguous forests \( T_{i-k}, T_{i-k+1}, \ldots, T_{i-1} \). This partitioning technique was introduced by Nagamochi and Ibaraki in [18], and these forests are known as Nagamochi-Ibaraki forests (or NI forests). The following is a basic property of NI forests.

**Lemma 2** (Nagamochi-Ibaraki [18] [17]). For any pair of vertices \( u, v \), they are connected in NI forests \( T_1, T_2, \ldots, T_{k(u,v)} \) for some \( k(u,v) \) and not connected in any forest \( T_j \) for \( j > k(u,v) \).

Nagamochi and Ibaraki also gave an algorithm for constructing NI forests that runs in \( O(m + n) \) time if \( G_M \) is a simple graph (i.e. \( G \) is unweighted) and \( O(m + n \log n) \) time otherwise [18] [17].
3 The Algorithm

We describe our sparsification algorithm for an unweighted graph \( G = (V, E) \) with \( m \) edges and \( n \) vertices, and an error parameter \( \varepsilon \in (0, 1] \) as inputs. We prove that the skeleton graph \( G_\varepsilon \) produced by the algorithm is in \((1 \pm \varepsilon)G \) with high probability. We then show that the expected number of edges in \( G_\varepsilon \) is \( O(n \log n / \varepsilon^2) \). Finally, we prove that the expected time complexity of the algorithm is \( O(m) \).

Description of the Algorithm. The algorithm has three phases. The first phase has the following steps:

- If \( m \leq 2pn \), where \( p = \frac{1014 \ln n}{0.398^2} \), \( G \) is sparse enough itself. Therefore, we take \( G \) as our skeleton graph.
- Otherwise, we construct a set of NI forests of \( G \) and all edges in the first \( 2p \) NI forests are included in the skeleton graph \( G_\varepsilon \) with weight 1. We call these edges \( F_0 \). The edge set \( Y_0 \) is then defined as \( E \setminus F_0 \).

The second phase is iterative. The input to iteration \( i \) is a graph \((V, Y_{i-1})\), which is a subgraph of the input graph to iteration \( i - 1 \) (i.e. \( Y_{i-1} \subseteq Y_{i-2} \)). Iteration \( i \) comprises the following steps:

- If the number of edges in \( Y_{i-1} \) is at most \( 2pn \), we take all those edges in \( G_\varepsilon \) with weight \( 2^{i-1} \) each, and terminate the algorithm.
- Otherwise, all edges in \( Y_i \) are sampled with probability \( 1/2 \); call the sample \( X_i \) and let \( G_i = (V, X_i) \).

- We identify a set of edges in \( X_i \) (call this set \( F_i \)) that has the following properties:
  - The number of edges in \( F_i \) is at most \( 2k_i|V_\varepsilon| \), where \( k_i = p \cdot 2^{i+1} \), and \( V_\varepsilon \) is the set of components in \((V, Y_i)\), where \( Y_i = X_i \setminus F_i \).
  - Each edge in \( Y_i \) is \( k_i \)-heavy in \( G_i \).
- We give a sampling probability \( p_i = \min\left(\frac{3}{169 \cdot 2^{i-1} \cdot p}, 1\right) \) to all edges in \( F_i \).

The final phase consists of replacing each edge in \( F_i \) (for each \( i \)) with \( 2^i \) parallel edges, and then sampling each parallel edge independently with probability \( p_i \). If an edge is selected in the sample, it is added to the skeleton graph \( G_\varepsilon \) with weight \( 1/p_i \).

We now give a short description of the sub-routine that constructs the set \( F_i \) in iteration \( i \) of the second phase of the algorithm. This subroutine is iterative itself: we start with \( V_\varepsilon = V \) and \( E_\varepsilon = X_i \), and let \( G_\varepsilon = (V_\varepsilon, E_\varepsilon) \). We repeatedly construct \( k_i + 1 \) NI forests for \( G_\varepsilon \) where \( k_i = p \cdot 2^{i+1} \) and contract all edges in the \((k_i + 1)\)st forest to obtain a new \( G_i \), until \( |E_\varepsilon| \leq p \cdot 2k_i|V_\varepsilon| \). The set of edges \( E_\varepsilon \) that finally achieves this property forms \( F_i \).

The complete algorithm is given in Figure 1.

Cut Preservation. We first show that the skeleton graph \( G_\varepsilon \) produced by the above algorithm is in \((1 \pm \varepsilon)G \) with high probability. We use the following notation throughout: for any set of unweighted edges \( Z, cZ \) denotes these edges with a weight of \( c \) given to each edge.

Our goal is to prove the following theorem.

Theorem 5. \( G_\varepsilon \in (1 \pm \varepsilon)G \) with probability at least \( 1 - 8/n \).

As outlined in the introduction, our proof has two stages. Let \( K \) be the maximum value of \( i \) for which \( F_i \neq \emptyset \); let \( S = (\cup_{i=0}^{K} 2^i F_i) \cup 2^K Y_K \) and \( G_S = (V, S) \). Then, in the first stage, we prove the following theorem.
1. Set $\rho = \frac{\log n}{0.38e^2}$.

2. If $m \leq 2\rho n$, then $G_{e} = G$; else, go to step 3.

3. Construct NI forests $T_1, T_2, \ldots$ for $G$.

4. Set $i = 0$.

5. Set $X_i = E; F_i = \cup_{1 \leq j \leq 2\rho} T_j; Y_i = X_i \setminus F_i$.

6. Add each edge in $F_i$ to $G_e$ with weight 1.

7. If $|Y_i| \leq 2\rho n$, then add each edge in $Y_i$ to $G_e$ with weight $2^{i-1}$ and terminate; else, go to step 8.

8. Sample each edge in $Y_i$ with probability $1/2$ to construct $X_{i+1}$.

9. Increment $i$ by 1.

10. Set $E_c = X_i; V_c = V$.

11. Set $k_i = \rho \cdot 2^{i+1}$.

12. If $|E_c| \leq 2k_i|V_c|$, then

   (a) Set $F_i = E_c; Y_i = X_i \setminus E_c$.

   (b) For each edge $e \in F_i$, set $\lambda_e = \rho \cdot 4^i$.

   (c) Go to step 7.

   Else,

   (a) Construct NI forests $T_1, T_2, \ldots, T_{k_i+1}$ for graph $G_c = (V_c, E_c)$.

   (b) Update $G_c = (V_c, E_c)$ by contracting all edges in $T_{k_i+1}$.

   (c) Go to step 12.

13. For each edge $e \in \cup_i F_i$,

   (a) Set $p_e = \min(\frac{\log n}{0.38e^2}, 1) = \min(\frac{3}{109 \cdot 2e^{1.5}}, 1)$.

   (b) Generate $r_e$ from $\text{Binomial}(2^i, p_e)$.

   (c) If $r_e > 0$, add edge $e$ to $G_e$ with weight $r_e/p_e$.

Figure 1: Our sparsification algorithm
Theorem 6. \(G_S \in (1 \pm \varepsilon/3)G\) with probability at least \(1 - 4/n\).

In the second stage, we prove the following theorem.

Theorem 7. \(G_e \in (1 \pm \varepsilon/3)G_S\) with probability at least \(1 - 4/n\).

Combining the above two theorems and using the union bound, we obtain Theorem 5 (Observe that since \(\varepsilon \leq 1\), \((1 + \varepsilon/3)^2 \leq 1 + \varepsilon\) and \((1 - \varepsilon/3)^2 \geq 1 - \varepsilon\).

The following property is key to proving both Theorem 6 and Theorem 7.

Lemma 3. For any \(i \geq 0\), any edge \(e \in Y_i\) is \(k_i\)-heavy in \(G_i = (V, X_i)\), where \(k_i = \rho \cdot 2^{i+1}\).

Proof. For \(i = 0\), all edges in \(Y_i\) are in NI forests \(T_{2p+1}, T_{2p+2}, \ldots\) of \(G_i = G\). The proof follows from Lemma 2.

We now prove the lemma for \(i \geq 1\). Let \(G_e = (V_e, E_e)\) be the component of \(G_i\) containing \(e\). We will show that \(e\) is \(k_i\)-heavy in \(G_e\); since \(G_e\) is a subgraph of \(G_i\), the lemma follows. In the execution of the else block of step 12 on \(G_e\), there are multiple contraction operations, each of them comprising the contraction of a set of edges. We show that any such contracted edge is \(k_i\)-heavy in \(G_e\); it follows that \(e\) is \(k_i\)-heavy in \(G_e\).

Let \(G_e\) have \(t\) contraction phases and let the graph produced after contraction phase \(r\) be \(G_{e,r}\). We now prove that all edges contracted in phase \(r\) must be \(k_i\)-heavy in \(G_e\) by induction on \(r\). For \(r = 1\), since \(e\) appears in the \((k_i + 1)\)st NI forest of phase 1, \(e\) is \(k_i\)-heavy in \(G_e\) by Lemma 2. For the inductive step, assume that the property holds for phases 1, 2, \ldots, \(r\). Any edge that is contracted in phase \(r + 1\) appears in the \((k_i + 1)\)st NI forest of phase \(r + 1\); therefore, \(e\) is \(k_i\)-connected in \(G_{e,r}\) by Lemma 2. By the inductive hypothesis, all edges of \(G_e\) contracted in previous phases are \(k_i\)-heavy in \(G_{e,r}\); therefore, an edge that is \(k_i\)-heavy in \(G_{e,r}\) must have been \(k_i\)-heavy in \(G_e\).

Proof of Theorem 6. The next lemma follows from Lemma 1.

Lemma 4. With probability at least \(1 - 4/n^2\), for every cut \(C\) in \(G_P\), \(|2 x^{(C)}_{i+1} + f^{(C)}_i - x^{(C)}_i| \leq \varepsilon/13 + \varepsilon/2^{i+1}\).

Proof. Use the following parameters in Lemma 1.

- \(R = Y_i; Q = X_i; \bar{R} = 2X_{i+1}\)
- \(\delta = \varepsilon/13; \rho = 1/2; \pi = \rho \times 2^{i+1}\).

Lemma 2 ensures that \(R\) is \(\pi\)-heavy in \((V, Q)\); also, it can be verified that \(\delta^2 \cdot \rho \cdot \pi = 6 \ln n\).

We use the above lemma to prove the following lemma.

Lemma 5. Let \(S_j = \left(\bigcup\nolimits_{i=j}^{K-1} 2^{i-j} F_i\right) \cup 2^{K-j} Y_K\) for any \(j \geq 0\). Then, \(S_j \in (1 \pm (\varepsilon/3)2^{-i/2})G_j\) with probability at least \(1 - 4/n\), where \(G_j = (V, X_j)\).

To prove this lemma, we need to use the following fact.

Fact 1. Let \(x \in (0, 1)\) and \(r_i = 13 \cdot 2^{i/2}\). Then, for any \(k \geq 0\),

\[
\prod_{i=0}^{k} (1 + x/r_i) \leq 1 + x/3
\]

\[
\prod_{i=0}^{k} (1 - x/r_i) \geq 1 - x/3.
\]
Proof. We prove by induction on \( k \). For \( k = 0 \), the property trivially holds. Suppose the property holds for \( k - 1 \). Then,

\[
\prod_{i=0}^{k}(1+x/r_i) = \prod_{i=0}^{k}(1 + \frac{x}{13 \cdot 2^{i/2}})
\]

\[
= (1+x/13) \cdot \prod_{i=1}^{k} \left(1 + \frac{x/\sqrt{2}}{13 \cdot 2^{(i-1)/2}}\right)
\]

\[
\leq (1+x/13) \cdot \left(1 + \frac{x}{3\sqrt{2}}\right)
\]

\[
\leq 1 + x/3
\]

\[
\prod_{i=0}^{k}(1-x/r_i) = \prod_{i=0}^{k}(1 - \frac{x}{13 \cdot 2^{i/2}})
\]

\[
= (1-x/13) \cdot \prod_{i=1}^{k} \left(1 - \frac{x/\sqrt{2}}{13 \cdot 2^{(i-1)/2}}\right)
\]

\[
\geq (1-x/13) \cdot \left(1 - \frac{x}{3\sqrt{2}}\right)
\]

\[
\geq 1 - x/3.
\]

\[\square\]

Proof of Lemma. For any cut \( C \) in \( G \), let the edges crossing \( C \) in \( S_j \) be \( S_j^C \), and let their total weight be \( s_j^C \). Also, let \( X_i^C, Y_i^C \) and \( F_i^C \) be the set of edges crossing \( C \) in \( X_i, Y_i \) and \( F_i \) respectively, and let their total weights be \( x_j^C, y_j^C \) and \( f_j^C \). (Recall that all edges in \( X_i, Y_i \) and \( F_i \) are unweighted; therefore \( x_j^C = |X_j^C|, y_j^C = |Y_j^C| \) and \( f_j^C = |F_j^C| \).

Since \( K \leq n - 1 \), we can use the union bound on Lemma 4 to conclude that with probability at least \( 1 - 4/n \), for every \( 0 \leq i \leq K \) and for all cuts \( C \),

\[
2x_{i+1}^C + f_i^C \leq (1 + \varepsilon/r_i)x_i^C
\]

\[
2x_i^C + f_i^C \geq (1 - \varepsilon/r_i)x_i^C
\]

where \( r_i = 13 \cdot 2^{i/2} \). Then,

\[
s_j^C = 2^{K-i}y_k^C + 2^{K-j}f_k^C + 2^{K-1-j}f_{K-1}^C + \ldots + f_j^C
\]

\[
= 2^{K-i}y_k^C + 2^{K-1-j}f_{K-1}^C + \ldots + f_j^C \quad \text{since } y_k^C + f_k^C = x_k^C
\]

\[
= 2^{K-1-i}(2x_k^C + f_k^C) + (2^{K-2-j}f_{K-2} + \ldots + f_j^C)
\]

\[
\leq (1 + \varepsilon/r_{K-1})2^{K-1-i}x_{K-1}^C + (2^{K-2-j}f_{K-2} + \ldots + f_j^C)
\]

\[
\leq (1 + \varepsilon/r_{K-1})(2^{K-1-i}x_{K-1}^C + 2^{K-2-j}f_{K-2} + \ldots + f_j^C)
\]

\[
\ldots
\]

\[
\leq (1 + \varepsilon/r_{K-1})(1 + \varepsilon/r_{K-2}) \ldots (1 + \varepsilon/r_j)x_j^C
\]

\[
\leq (1 + (\varepsilon/2^{j/2})/r_{K-1-j})(1 + (\varepsilon/2^{j/2})/r_{K-2-j}) \ldots (1 + (\varepsilon/2^{j/2})/r_0)x_j^C \quad \text{since } r_{j+1} = r_j \cdot 2^{j/2}
\]

\[
\leq (1 + (\varepsilon/3)2^{-j/2})x_j^C \quad \text{by Fact 1}.
\]
Similarly,

$$s_j^C = 2^{K-j}x_j^C + 2^{K-j}f_j^C + 2^{K-j-1}f_{K-1}^C + \ldots + f_j^C$$

$$= 2^{K-j}x_j^C + \sum_{i=j}^{K-1} f_i^C$$ since $x_j^C + f_j^C = x_j^C$.

$$= 2^{K-j}x_j^C + f_j^C + 2^{K-j}f_{K-1}^C + \ldots + f_j^C$$

$$\geq (1 - \varepsilon / r_{K-1}) 2^{K-j}x_j^C + (2^{K-j}f_{K-2}^C + \ldots + f_j^C)$$

$$\geq (1 - \varepsilon / r_{K-1}) (2^{K-j}x_j^C + f_j^C + 2^{K-j}f_{K-2}^C + \ldots + f_j^C)$$

$$\vdots$$

$$\geq (1 - \varepsilon / r_{K-1}) (1 - \varepsilon / r_{K-2}) \ldots (1 - \varepsilon / r_j) x_j^C$$

$$\geq (1 - (\varepsilon 2^{-j/2}) / r_{K-1}) \ldots (1 - (\varepsilon 2^{-j/2}) / r_0) x_j^C$$

since $r_{j+1} = r_j \cdot 2^{j/2}$

$$\geq (1 - (\varepsilon / 3) 2^{-j/2}) x_j^C$$ by Fact[1] \qed

Theorem[3] now follows as a corollary of the above lemma for $j = 0$.

**Proof of Theorem[7]** Now, we use the sparsification framework developed in [9] and outlined previously in [2] to prove Theorem[7]. Observe that edges $F_0 \cup 2^K Y_K$ are identical in $G_{\varepsilon}$ and $G_e$. Therefore, we do not consider these edges in the analysis below.

For any $i \geq 1$, let $\psi(i)$ be such that $2^{\psi(i)} \leq \rho \cdot 4^i \leq 2^{\psi(i)+1}$. Note that for any $j$, $\psi(i) = j$ for at most one value of $i$. Then, for any $j \geq 1$, $R_j = F_i$ if $j = \psi(i)$ and $R_j = \emptyset$ if there is no $i$ such that $j = \psi(i)$. We set $\alpha = 32/3$; $\pi = \rho \cdot 4^K$; for any $j \geq 1$, $Q_j = (V, W_j)$ where $W_j = \cup_{i-1 \leq r \leq K} 4^{K-r+1} 2^r F_r$ if $R_j \neq \emptyset$ and $j = \psi(i)$, and $W_j = \emptyset$ if $R_j = \emptyset$.

The following lemma proves $\pi$-connectivity.

**Lemma 6.** With probability at least $1 - 4/n$, every edge $e \in F_i = R_{\psi(i)}$ for each $i \geq 1$ is $\rho \cdot 4^K$-heavy in $Q_{\psi(i)}$.

**Proof.** Consider any edge $e \in F_i$. Since $F_i \subseteq Y_{i-1}$, Lemma[3] ensures that $e$ is $\rho \cdot 2^i$-heavy in $G_{i-1} = (V, X_{i-1})$, and therefore $\rho \cdot 2^{i-1}$-heavy in $(V, 2^{i-1} X_{i-1})$. Since $\varepsilon \leq 1$, Lemma[5] ensures that with probability at least $1 - 4/n$, the weight of each cut in $(V, 2^{i-1} X_{i-1})$ is preserved up to a factor of 2 in $Z_{i} = (V, \cup_{i-1 \leq r \leq K} 2^r F_r)$. Thus, $e$ is $\rho \cdot 4^{i-1}$-heavy in $Z_i$.

Consider any cut $C$ containing $e \in F_i$. We need to show that the weight of this cut in $Q_{\psi(i)}$ is at least $4^K$.

Let the maximum $\lambda_d$ of an edge in $C$ be $\rho \cdot 4^{K}$, for some $k_c \geq i$. By the above proof, $a$ is $\rho \cdot 4^{k_c-1}$-heavy in $Z_{k_c}$. Then, the total weight of edges crossing cut $C$ in $Q_{\psi(k_c)}$ is at least $\rho \cdot 4^{K-k_c+1} + \rho \cdot 4^K$. Since $k_c \geq i$, $\psi(k_c) \geq \psi(i)$ and $Q_{\psi(k_c)}$ is a subgraph of $Q_{\psi(i)}$. Therefore, the the total weight of edges crossing cut $C$ in $Q_{\psi(i)}$ is at least $\rho \cdot 4^K$. \qed

We now prove the $\alpha$-overlap property. For any cut $C$, let $f_j^C$ and $w_j^C$ respectively denote the total weight of edges crossing cut $C$ in $F_j$ and $W_{\psi(i)}$ respectively for any $i \geq 0$. Further, let the number of edges
crossing cut C in \( \cup_{i=1}^{K} 2^{j} F_i \) be \( f(C) \). Then,

\[
\sum_{i=1}^{K} \frac{w_i(C)}{2^{\psi(i)-1}} \leq \sum_{i=1}^{K} \frac{w_i(C) \cdot 4^i}{2 \rho \cdot 4^k} = \sum_{i=1}^{K} \frac{w_i(C)}{2 \cdot 4^{K-i}} = \sum_{i=1}^{K} \frac{w_i(C)}{2 \cdot 4^{K-i}} = \sum_{i=1}^{K} \sum_{r=1}^{K} \frac{f_r(C) \cdot 2^{r-4^{K-r+1}}}{2 \cdot 4^{K-i}}
\]

Using Theorem 4 we conclude the proof of Theorem 7.

**Size of the skeleton graph.** We now prove that the expected number of edges in \( G_\varepsilon \) is \( O(n \log n / \varepsilon^2) \).

For \( i \geq 1 \), define \( D_i \) to be the set of connected components in the graph \( G_i = (V, X_i) \); let \( D_0 \) be the single connected component in \( G \). For any \( i \geq 1 \), if any connected component in \( D_i \) remains intact in \( D_{i+1} \), then there is no edge from that connected component in \( F_i \). On the other hand, if a component in \( D_i \) splits into \( \eta \) components in \( D_{i+1} \), then the algorithm explicitly ensures that \( \sum_{r \in \mathcal{F}_i} w_r \) from that connected component is

\[
\sum_{r \in \mathcal{F}_i} \frac{w_r}{2^{\rho - 2^r}} \leq \left( \frac{\rho^{2^{\psi(i)-1}}}{\rho^{2^0}} \right)^\eta = 4 \eta \leq 8(\eta - 1).
\]

Therefore, if \( d_i = |D_i| \), then

\[
\sum_{i=1}^{K} \sum_{r \in \mathcal{F}_i} \frac{w_r}{2^{\rho - 2^r}} \leq \sum_{i=1}^{K} 8(d_{i+1} - d_i) \leq 8n,
\]

since we can have at most \( n \) singleton components. It follows from Theorem 4 that the expected number of edges added to \( G_\varepsilon \) by the sampling is \( O(n \log n / \varepsilon^2) \). Since the number of edges added to \( G_\varepsilon \) in steps 6 and 7 of the algorithm is \( O(n \log n / \varepsilon^2) \), the total number of edges in \( G_\varepsilon \) is \( O(n \log n / \varepsilon^2) \).

**Time complexity of the algorithm.** If \( m \leq 2\rho n \), the algorithm terminates after the first step which takes \( O(m) \) time. Otherwise, we prove that the expected running time of the algorithm is \( O(m + n \log n / \varepsilon^2) = O(m) \) since \( \rho = \Theta(\log n / \varepsilon^2) \). First, observe that phase 1 takes \( O(m + n \log n) \) time. We will show that iteration \( i \) of phase 2 takes \( O(|Y_{i-1}|) \) time. Since \( Y_i \subseteq X_i \) and \( E[|X_i|] = E[|X_{i-1}|] / 2 \), and \( |Y_0| \leq m \), it follows that the expected overall time complexity of phase 2 is \( O(m) \). Finally, the time complexity of phase 3 is \( O(m + n \log n / \varepsilon^2) \) (see e.g. [10]).

In iteration \( i \) of phase 2, the first step takes \( |Y_{i-1}| \) time. We show that all the remaining steps take \( O(|X_i| + n \log n) \) time. Since \( X_i \subseteq Y_{i-1} \) and the steps are executed only if \( Y_{i-1} = \Omega(n \log n / \varepsilon^2) \), it follows that the total time complexity of iteration \( i \) of phase 2 is \( O(|Y_{i-1}|) \).

First, observe that step 8 and the if block of step 12 take \( O(|X_i|) \) time. So, we are left with the repeated invocations of the else block of step 12. Each invocation of the else block takes \( O(|V_c| \log n + |E_c|) \) time for the current \( V_c, E_c \). So, the last invocation of the else block takes at most \( O(|X_i| + n \log n) \) time. In any other invocation, \( |E_c| = \Omega(|V_c| \log n) \) and hence the time spent is \( O(|E_c|) \). We show that \( |E_c| \) decreases by a factor of 2 from one invocation of the else block to the next; then the total time over all invocations of the else block is \( O(|X_i| + n \log n) \).

To see that the \( |E_c| \) halves from one invocation of the else block to the next, consider an iteration that begins with \( |E_c| > 2k_i \cdot |V_c| \). By Lemma 2, \( E_c \) for the next iteration (denoted by \( E'_c \)) comprises only edges in the first \( k_i \) NI forests constructed in the current iteration. So \( |E'_c| \leq k_i \cdot |V_c| < |E_c| / 2 \).
4 Future Work

The obvious open question is whether these results can be extended to weighted graphs, at least if the weights are polynomially bounded in $n$. Another possibility is to extend these results to the semi-streaming model for unweighted graphs. A more ambitious open problem is to obtain an efficient (i.e. near-linear in $m$) algorithm that constructs a skeleton containing $o(n \log n)$ edges while approximately preserving the weights of all cuts with high probability.

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A Proof of Lemma 1

To prove Lemma 1 we will need two theorems from [9]. The first theorem is a non-uniform extension of Chernoff bounds.

**Theorem 8** (Hariharan-Panigrahi[9]). Consider any subset $C$ of unweighted edges, where each edge $e \in C$ is sampled independently with probability $p_e$ for some $p_e \in [0, 1]$ and given weight $1/p_e$ if selected in the sample. Let the random variable $X_e$ denote the weight of edge $e$ in the sample; if $e$ is not selected in the sample, then $X_e = 0$. Then, for any $p$ such that $p \leq p_e$ for all edges $e$, any $\epsilon \in (0, 1]$, and any $N \geq |C|$, the following bound holds:

$$\Pr \left[ \left| \sum_i X_e - |C| \right| > \epsilon N \right] < 2e^{-0.38\epsilon^2 p_N}.$$  

To state the second theorem, we need the following definitions.

**Definition 3.** For any undirected graph $G$ and for any $k > 0$, the $k$-projection of any cut $C$ is the set of $k$-heavy edges in $C$.

**Definition 4.** The edge connectivity of an undirected graph $G$ is the minimum weight of a cut in $G$.

The theorem counts the number of distinct $k$-projections in cuts of weight $\alpha k$ for any $k \geq c$, where $c$ is the edge connectivity of the graph.

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2For Chernoff bounds, see e.g. [15].

3For any event $\delta$, $\Pr[\delta]$ represents the probability of event $\delta$.  

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Theorem 9 (Hariharan-Panigrahi). For any undirected graph with edge connectivity \( c \) and for any \( k \geq c \) and any \( \alpha \geq 1 \), the number of distinct \( k \)-projections of cuts of weight at most \( \alpha k \) is at most \( n^{2\alpha} \).

Using the above two theorems, we now prove Lemma 1.

Proof of Lemma 1. Let \( \mathcal{C}_j \) be the set of all cuts \( C \) such that \( 2^j \pi \leq r(C) \leq 2^{j+1} \pi - 1 \), \( j \geq 0 \). We will prove that with probability at least \( 1 - 2e^{-0.38 \cdot 2^j \cdot \pi \cdot 2^j} \leq 2e^{-6 \cdot 2^j \cdot \ln n} \), all cuts in \( \mathcal{C}_j \) satisfy the property of the lemma. Then, the lemma follows by using the union bound over \( j \) since \( 2n^{-2} \cdot 2^{n-4} + \ldots + 2n^{-2j} + \ldots \leq 4n^{-2} \).

We now prove the property of the lemma for cuts \( C \in \mathcal{C}_j \). Since each edge \( e \in R(C) \) is sampled with probability \( p \) in obtaining \( \hat{R}(C) \), we can use Theorem 8 with sampling probability \( p \). Then, for any \( R(C) \) where \( C \in \mathcal{C}_j \), by Theorem 8 we have

\[
P \left[ \left| \hat{r}(C) - r(C) \right| > \delta q(C) \right] < 2e^{-0.38 \cdot 2^j \cdot p \cdot q(C)} \leq 2e^{-6 \cdot 2^j \cdot \ln n} = 2n^{-6 \cdot 2^j},
\]

since \( q(C) \geq \pi \cdot 2^j \) for any \( C \in \mathcal{C}_j \). Since each edge in \( R(C) \) is \( \pi \)-heavy in \((V, Q)\), Theorem 9 ensures that the number of distinct \( R(C) \) sets for cuts \( C \in \mathcal{C}_j \) is at most \( n^{2^j(\pi/2^j + 1)} = n^{4 \cdot 2^j} \). Using the union bound over these distinct \( R(C) \) edge sets, we conclude that with probability at least \( 1 - 2n^{-2j+1} \), all cuts in \( \mathcal{C}_j \) satisfy the property of the lemma. \( \square \)