Distributed MST: A Smoothed Analysis

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ABSTRACT

We study smoothed analysis of distributed graph algorithms, focusing on the fundamental minimum spanning tree (MST) problem. With the goal of studying the time complexity of distributed MST as a function of the "perturbation" of the input graph, we posit a *smoothing model* that is parameterized by a smoothing parameter $0 \le \epsilon(n) \le 1$ which controls the amount of *random* edges that can be added to an input graph *G* per round. Informally, $\epsilon(n)$ is the probability (typically a small function of *n*, e.g., $n^{-\frac{1}{4}}$) that a random edge can be added to a node per round. The added random edges, once they are added, can be used (only) for communication.

We show upper and lower bounds on the time complexity of distributed MST in the above smoothing model. We present a distributed algorithm that, with high probability,¹ computes an MST and runs in $\tilde{O}(\min\{\frac{1}{\sqrt{\epsilon(n)}}2^{O(\sqrt{\log n})}, D + \sqrt{n}\})$ rounds² where ϵ is the smoothing parameter, D is the network diameter and n is the network size.

To complement our upper bound, we also show a lower bound of $\tilde{\Omega}(\min\{\frac{1}{\sqrt{\epsilon(n)}}, D + \sqrt{n}\})$. We note that the upper and lower bounds

essentially match except for a multiplicative $2^{O(\sqrt{\log n})} \operatorname{polylog}(n)$ factor.

Our work can be considered as a first step in understanding the smoothed complexity of distributed graph algorithms.

CCS CONCEPTS

- Theory of computation \rightarrow Distributed algorithms.

KEYWORDS

distributed algorithms, smoothed analysis, random model, minimum spanning tree, lower bound

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1 INTRODUCTION AND MOTIVATION

Smoothed analysis of algorithms was introduced in a seminal paper by Speilman and Teng [23] to explain why the well-studied simplex algorithm for linear programming does well in practice, despite having an (worst-case) exponential run time in theory. The highlevel idea behind the smoothed analysis of the simplex algorithm is the following:

- perturbing the input data with a *small* amount of *random* noise (e.g., Gaussian noise with mean zero, parameterized by the variance of the noise), and then
- (2) showing that the perturbed input can be solved efficiently by the simplex algorithm, i.e., in polynomial time. In particular, Spielman and Teng quantify the run time as a function of the perturbation; the more the perturbation (i.e., larger the variance of the noise), the faster the run time.

Smoothed analysis is thus different from the *worst-case* analysis of algorithms. It is also different from the *average-case analysis*, which assumes a probability distribution on the set of all possible inputs. Smoothed analysis, on the other hand, is sort of a hybrid between the above two — it considers the worst-case input and then randomly perturbs it. If even small perturbations (say, adding random noise) lead to efficient run time, then this means that the worst-case is quite sensitive to the input parameters. In practice, there will usually be noise and thus the algorithm is likely to avoid the worst-case behavior.

In this paper, we initiate the study of *smoothed analysis of distributed graph algorithms.* Our paper is motivated by the work of Dinitz et al. [5] who initiated the study of smoothed analysis for *dynamic networks* (we refer to Section 1.1 for more details). A main contribution of our paper is positing smoothing models in the context of *distributed graph algorithms* and performing analyses of the models. While many smoothing models are possible for such algorithms, a key goal is to identify models that lead to non-trivial bounds on the distributed complexity (here we focus on time complexity) of fundamental graph algorithms.

We focus on the distributed minimum spanning tree (MST) problem in synchronous *CONGEST* networks (see Section 2 for details on this standard distributed computing model). The worst-case time (round) complexity of distributed MST has been extensively studied for the last three decades and tight bounds are now well established (see, e.g., [17, 18]). There is an optimal distributed MST

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¹Throughout, with high probability (whp) means with probability at least $1 - n^{-c}$, for some fixed, positive constant *c*.

²The notation \tilde{O} hides a polylog(*n*) factor and $\tilde{\Omega}$ hides a $\frac{1}{\text{polylog}(n)}$ factor, where *n* is the number of nodes of the graph.

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algorithm (see, e.g., [16]) that runs in $\tilde{O}(D + \sqrt{n})$ rounds, where *D* is the graph diameter and *n* is the number of nodes in the network. Also, there is a (essentially) matching lower bound of $\tilde{\Omega}(D + \sqrt{n})$ rounds that applies even to randomized Monte-Carlo distributed algorithms [22].

The lower bound is shown by presenting a weighted graph (in particular, a family of graphs) and showing that no distributed algorithm can solve MST faster. This raises a motivating question for smoothed analysis: Is the worst-case bound specific to the choice of the weighted graph (family)? Or more precisely, is it specific to the choice of the graph topology or the edge weights or both? If small perturbations do not change the worst-case bound by too much then we can say that the lower bound is *robust* and, if they do we can say that the bounds are *fragile* [5]. Thus smoothed analysis can lead to a better understanding of the complexity of distributed MST by studying perturbations of the worst-case input. This is one of the motivations in studying smoothed analysis of distributed algorithms.

However, to answer the above questions, one has to first come up with a suitable smoothing model. For example, one possible smoothing model, in the spirit of Speilman and Teng's original smoothing model, would be perturbing the edge weights of the input graph by a small amount. It is apparent that if the perturbation is quite small relative to the weights (since the weights can be wellspaced), then this does not any effect on the lower bound - it remains $\tilde{\Omega}(D + \sqrt{n})$. Another possible model, which we explore in this paper, again in the spirit of original model but now applied to perturbing the topology of the input graph, is smoothing the input graph by adding³ a small number of random edges. While there are a few possible ways to accomplish this, we focus on a particular smoothing model described next. We will discuss other smoothing models (which can be considered variants of this model) in Section 5. A practical motivation for this kind of smoothing, i.e., adding a small number of random edges to a given graph, is that many real-world networks might be better modeled by graphs with some underlying structure with some amount of randomness. For example, it is well know that real-world graphs have power-law degree distribution, but they are not arbitrary (worst-case) powerlaw graphs but can be reasonably modeled by random graphs with power-law degree distribution [7].

We consider a *smoothing model* (see Section 2.2) that is parameterized by a smoothing parameter $0 \le \epsilon = \epsilon(n) \le 1$ that controls the amount of *random* edges that can be added to an input graph G = (V, E) per round. $\epsilon(n)$ is typically a small function of *n*, say, $\epsilon(n) = n^{-\frac{1}{4}}$. More precisely, our smoothing model allows any node to add a random edge with probability $\epsilon(n)$ in each round; the added edges can be used for communication in later rounds. (We note that the added edges, otherwise, do not change the underlying solution with respect to *G*; e.g., for MST, the added edges have weight ∞ and hence don't affect the MST of *G*.) Besides this additional feature, nodes behave as in the standard model, i.e., can communicate using edges of *G*. We formally define the model in Section 2.2. Note that nodes can as well choose *not* to use this additional feature. Depending on $\epsilon(n)$, the number of random edges added per round can be small. (In Section 5, we consider a variant of this model, which essentially gives the same bounds as the ones discussed here.)

An alternate way of thinking about our smoothing model is as follows. Assume that the graph *G* is embedded in a congested clique. The congested clique model has been studied extensively in the distributed computing literature; see, e.g., [2, 3, 10–12, 19–21]). A node — besides using its incident edges in *E* — can also *choose* to use a random edge (not in *G*, but in the clique) with probability ϵ in a round to communicate (once chosen, a random edge can be used subsequently till end of computation). Note that if ϵ is small, say, for example $\epsilon = O(n^{-\frac{1}{4}})$, then the probability of adding a random edge by a node in a round is small. In particular, if $\epsilon = 0$, then this boils down to the traditional model, i.e., working on the given graph *G* with no additional random edges, as ϵ increases, the number of random edges increases with it.

We note that the smoothing model is sort of a hybrid between the traditional model where communication is allowed only along the edges of an arbitrary graph *G* and a model where *G* is a random graph (e.g., Erdos-Renyi graph model [4, 13]) or an expander (see e.g., [1] and the references therein). In the smoothing model we start with an arbitrary graph *G* and add random edges (parameterized by ϵ). In Section 2.2, we further explore relationship between the smoothing model and other distributed computing models.

Our goal is to study how the distributed complexity of MST varies as a function of $\epsilon(n)$ (among other usual graph parameters such as network size, network diameter, etc.). We show upper and lower bounds on the time complexity of distributed MST in the aforementioned smoothing model. We present a distributed algorithm, which (with high probability) computes an MST and runs in $\tilde{O}(\min\{\frac{1}{\sqrt{\epsilon(n)}}2^{O(\sqrt{\log n})}, D+\sqrt{n}\})$ rounds, where ϵ is the smoothing parameter, D is the network diameter, and n is the network size, i.e., the number of nodes in the network.

To complement our upper bound, we also show a lower bound of $\tilde{\Omega}(\min\{\frac{1}{\sqrt{\epsilon}}, D + \sqrt{n}\})$. Our bounds show non-trivial dependence on the smoothing parameter $\epsilon(n)$, and the bounds are essentially match except for a $2^{O(\sqrt{\log n})}$ factor and a polylogarithmic factor.

1.1 Related work

Smoothed analysis was introduced by Spielman and Teng[23] and has since been applied for various algorithms problems in the sequential setting (see, e.g., [24] for a survey).

The only work that we are aware of in the context of smoothed analysis of distributed algorithms is that of Dinitz et al. [5] who study smoothed analysis of distributed algorithms for *dynamic networks*. Their dynamic network model is a dynamic graph $\mathcal{H} =$ G_1, G_2, \ldots that describes an evolving network topology, where G_i is the graph at round *i*. It is assumed that all graphs in \mathcal{H} share the same node set, but the edges can change with some restrictions, e.g., each graph should be connected. They define a smoothing model for a dynamic graph that is parameterized with a smoothing factor $k \in \{1, 2, \ldots, \binom{n}{2}\}$. To *k*-smooth a dynamic graph \mathcal{H} is to replace each static graph G_i in \mathcal{H} with a smoothed graph G'_i sampled uniformly from the space of graphs that are: (1) within *edit distance k* of *G*, and (2) are allowed by the dynamic network model (e.g., smoothing cannot generate disconnected graph). The edit distance

 $^{^3}$ One can also delete edges, although we don't consider this in this paper, see Section 2.2.

is the number of edge additions/deletions needed to transform one graph to another, assuming they share the same node set.

Our smoothing model can also be thought of in terms of choosing a random graph within a *positive* edit distance (i.e., edges are only *added* to the original input graph) where the number of random edges added is proportional to $n\epsilon(n)$ (per round or in total — see Section 5).

Dinitz et al. study three well-known problems that have strong lower bounds in dynamic network models, namely, *flooding*, *random walks*, and *aggregation*. For each problem, they study robustness/fragility of the existing bound by studying how it improves under increasing amounts of smoothing.

2 OUR MODEL

We first discuss the distributed computing model and then discuss our smoothing model.

2.1 Distributed Computing Model

We consider a system of *n* nodes, represented as an undirected, connected graph G = (V, E). Each edge $e \in E$ may have an associated weight w(e), which can be represented using $O(\log n)$ bits. If there is no weight on an edge, then it can be considered to be ∞ . Each node *u* runs an instance of a distributed algorithm and has a unique identifier ID_u of $O(\log n)$ bits.

The computation advances in *synchronous* rounds, where in every round, nodes can send messages, receive messages that were sent in the same round by neighbors in *G*, and perform some local computation.

Our algorithms work in the CONGEST model [16], where in each round a node can send at most one message of size $O(\log n)$ bits via a single edge (whether the edge is in *G* or is a smoothed edge).

2.2 Smoothing Model

Given a (arbitrary) undirected connected graph G(V, E) (throughout n = |V|), the smoothing model allows adding some random edges to the input graph G, thereby "perturbing" graph structure. We call this process *smoothing*, where we add a small number of random edges to the original graph. We describe the process of adding edges which is parameterized by a smoothing parameter $0 \le \epsilon = \epsilon(n) \le 1$ as follows. The smoothing parameter (which in general is a function of n, the network size⁴ controls the amount of random edges that can be added per round. Henceforth, we call this as the ϵ -smoothing model.

More precisely, every node, in every round, with probability ϵ (the smoothing parameter) can *add* an edge to a *random* node (chosen uniformly at random from *V*) in the graph. Let the added random edges form the set *R* (different from the original edge set *E*). Note that we allow multi-edges in the random edge choosing process; however, if there is more than one edge between two nodes, then only one edge (especially, if it belongs to *E*) that matters. The added edge persists for future rounds and can be used henceforth for communication; its weight is ∞ . A distributed algorithm can

potentially exploit these additional edges to improve the time complexity. 5

In this work, we only consider adding edges to the graph; one can also consider deleting edges from the original graph. However, for many problems such as MST, it is arguably more appropriate to (potentially) add edges. In fact, deleting edges can change the graph. Whereas, in the ϵ -smoothing model, since the added edges to the given graph *G* are purely communicating edges (with weight ∞), the MST with respect to *G* is unchanged. In fact, the model allows us to study tradeoffs between the amount of random edges added to the efficiency of computing a solution of *G*.

As mentioned earlier, the ϵ -smoothing model gives a "smooth" tradeoff between the traditional CONGEST model where there no additional random edges ($\epsilon = 0$) in *G* (the input graph) and a model where there is a random graph embedded in *G*. In this sense, it is different from studying distributed computing on (purely) random graph models or expander graph models (e.g., [1, 4, 13]. We note the work of Ghaffari et al[8, 9] embeds a random graph in a given graph *G* and uses this embedding to design algorithms that depend on the mixing time of *G*. This is still the traditional CONGEST model, though we use their result in our algorithms.

As mentioned in Section 1, we can also relate the well-studied *congested clique* model to the ϵ -smoothing model and also give a way to understand computation tradeoffs between the traditional CONGEST model and the congested clique model. Assuming the input graph *G* is embedded in a congested clique, the ϵ parameter controls the power to use the non-graph clique edges. If $\epsilon = 0$, then we have the traditional CONGEST model and for any $\epsilon > 0$, if we spend enough rounds, then one can throw a random edge between every pair of nodes which boils down to the congested clique. Of course, this is costly, which illustrates the power of the congested clique model (where the clique edges can be used for "free"). Studying time and message complexity bounds in terms of ϵ can help us understand the power of the clique edges with respect to solving a problem on a given input graph.

3 DISTRIBUTED MST IN THE SMOOTHING MODEL

For the sake of exposition, we first present a distributed MST algorithm that runs in $\tilde{O}(\min\{\frac{1}{\epsilon}+2^{O(\sqrt{\log n})}, D+\sqrt{n}\})$ rounds. Then we present an improved algorithm that runs in $\tilde{O}(\min\{\frac{1}{\sqrt{\epsilon(n)}}2^{O(\sqrt{\log n})}, D+\sqrt{n}\})$ rounds. The second algorithm is a modification of the first and its time complexity approaches the lower bound of $\tilde{\Omega}(\min\{\frac{1}{\sqrt{\epsilon}}, D+\sqrt{n}\})$ shown in Section 4. Thus up to a multiplicative factor of

 $2^{O(\sqrt{\log n})}$ polylog *n*, the bounds are tight.

We give a high-level overview of our approach of our first algorithm before we get into the technical details. The algorithm can be described in two parts which are described in Sections 3.1 and 3.2 respectively. At the outset we note that if $1/\epsilon$ is larger compared to $\tilde{O}(D + \sqrt{n})$, then we simply run the standard time-optimal MST algorithm ([16]) without doing smoothing.

⁴We sometimes just write ϵ , understanding it to be a function of *n*.

 $^{^5 \}mathrm{In}$ this paper, we focus only on time complexity, but message complexity can also be relevant.

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3.1 Part 1: Constructing an Expander

Initially the algorithm exploits the smoothing model to add about $O(\log n)$ random edges per node. This can be accomplished as follows: each node (in parallel) tries to make a random edge selection for the (first) $\Theta(\frac{\log n}{\epsilon})$ rounds, where ϵ is the smoothing parameter. Since the probability of adding a random edge, i.e., a smoothing edge, is ϵ per round, it is easy to show that with high probability a node will add $\Theta(\log n)$ random edges. Via a union bound, this holds for all nodes.

Now consider the graph R(G) induced *only* by the smoothed (random) edges of *G* after $\Theta(\frac{\log n}{\epsilon})$ rounds. In the following lemma 3.2, we show that R(G) is a graph with $O(\log n)$ *mixing time*.

The proof of this result comes from the relation of ϵ -smoothing model to Erdos-Renyi random graph, which we will show next.

LEMMA 3.1. Consider a graph G(V, E) under ϵ -smoothing. If we invoke smoothing for ℓ rounds (where $\ell \epsilon = o(n)$), then the graph induced by the smoothed edges is an Erdos-Renyi random graph G(n,p) where $p = \Theta\left(\frac{l\epsilon}{n}\right)$.

PROOF. We calculate the probability of a smoothed edge between nodes *u* and *v*. Clearly the edge is present if either *u* or *v* successfully adds the other end during ℓ steps. Hence, $p = 1 - (1 - \frac{\epsilon}{n})^{2\ell} = \Theta\left(\frac{\ell\epsilon}{n}\right)$

Remark 1. The following lemma (Lemma 3.2) applies only to R(G) and *not necessarily* to $G \cup R(G)$.

LEMMA 3.2. Let G = (V, E) be an arbitrary undirected graph and let R(G) = (V, F) be the random graph induced (only) by the set Fof random (smoothed) edges after $\Theta(\frac{\log n}{\epsilon})$ rounds. Then, with high probability, R(G) has mixing time $\tau_{mix}(R) = O(\log n)$.

PROOF. Using Lemma 3.1, where $\ell = \Theta(\frac{\log n}{\epsilon})$, we have R(G) is a Erdos-Renyi random graph $G(n, p = \Theta(\frac{\log n}{n})$. It is well-known that with high probability the this random graph is an expander (i.e., has constant conductance) and thus has $O(\log n)$ mixing time (see e.g., [8]).

3.2 Part 2: Constructing an MST

In the second part, the algorithm uses R(G) as a "communication backbone" to construct an MST in $O(\log^2 n)2^{O(\sqrt{\log n})}$ rounds.

Our algorithm crucially uses a routing result due to Ghaffari et al. [8, 9] who show, given an arbitrary graph G = (V, E), how to do *permutation* (or more generally, *multi-commodity*) routing fast. We briefly describe the problem and the main result here and refer to [9] for the details. Permutation or multi-commodity routing is defined as follows: given source-destination pairs of nodes $(s_i, t_i) \in V \times V$ and suppose s_i wants to communicate with t_i (t_i does not know s_i beforehand, but s_i knows the ID of t_i). The width of the pairs is W if each $v \in V$ appears at most W times as s_i or t_i . The goal is to construct a routing path P_i (not necessarily simple) from s_i to t_i such that the set of routing paths has low *congestion* and low *dilation*. Congestion is the maximum number of times any edge is used in all the paths. Dilation is simply the maximum length of the paths. The main result of [9] is that routing paths P_i with *low congestion and dilation* can be found *efficiently*. Once such low congestion and dilation routing paths are found, using a standard trick of random delay routing, it is easy to establish that messages can be routed between the source and destination efficiently, i.e., proportional to congestion and dilation.

THEOREM 3.3 (EFFICIENT ROUTING). (Theorem 8 from [9]).Suppose we solve a multicommodity routing instance $\{(s_i, t_i)\}_i$ and achieve congestion c and dilation d. Then, in $\tilde{O}(c + d)$ rounds, every node s_i can send one $O(\log n)$ -bit message to every node t_i , and vice versa.

Next, let us formally restate their routing results for ease of discussion. First we state their result on multi-commodity routing on a random graph $G(n, \log n)$, i.e., a random graph where each node has $O(\log n)$ random edges (each endpoint chosen uniformly at random.)

THEOREM 3.4. (Theorem 1 from [9]) Consider a multicommodity routing instance of width $\tilde{O}(1)$. There is a multicommodity routing algorithm on a random graph $G(n, (\log n))$ that achieves congestion and dilation $2^{O(\sqrt{\log n})}$, and runs in time $2^{O(\sqrt{\log n})}$.

Then, by the construction of a random-graph-like hierarchy routing over a given graph G, we have the following result.

LEMMA 3.5. (Lemma 11 from [9]) On any graph G with n nodes and m edges, we can embed a random graph G(m, d) with $d \ge 200 \log n$ into G with congestion $\tilde{O}(\tau_{mix} \cdot d)$ and dilation τ_{mix} in time $\tilde{O}(\tau_{mix} \cdot d)$.

Using Lemma 3.5, we have the following trivial corollary for permutation routing.

COROLLARY 1 (PERMUTATION ROUTING). Consider a graph G = (V, E) and a set of n point-to-point routing requests (s_i, t_i) , where s_i, t_i are IDs of the corresponding source and destination. Each node of G is the source and the destination of exactly one message. Then there is a randomized algorithm that delivers all messages in time $\tau_{mix}(G)2^{O(\sqrt{\log n})}$, w.h.p.

Based on their multicommodity routing algorithm, they showed how to construct an MST of G in $\tau_{mix}(G)2^{O(\sqrt{\log n})}$ rounds (Ghaffari et al. [8]). However, this MST algorithm cannot be directly employed in our setting, i.e., to construct a MST of G, over $G \cup R(G)$.

We briefly summarize the idea from section 4 in their paper [8], to explain why this algorithm is not applicable directly in a black box manner. The main reason for this: while the algorithm of [8] operates on the graph *G*, ours operates on graph $G' = G \cup R(G)$. Applying the algorithm directly to $G \cup R(G)$ can (in general) yield an algorithm running in $\tau_{mix}(G')2^{O(\sqrt{\log n})}$ rounds where $\tau_{mix}(G')$ is the mixing time of *G'*. Note that $\tau_{mix}(G')$ (in general) can be of the same order of $\tau_{mix}(G)$ (even for constant smoothing parameter ϵ) in some graphs⁶. Thus the running time bound does not (in general) depend on ϵ and does not give our desired bound of $\tilde{O}(\log n/\epsilon)$ rounds. We give more details on the approach of [8] and then discuss our algorithm.

The approach of [8] is to modify Boruvka's algorithm [15], where the MST is built by merging tree fragments. In the beginning, each

⁶For example consider two cliques of size n/2 connected to each other via O(n) edges.

node is a fragment by itself. The fragment size grows by merging. To ensure efficient communication within a fragment, they maintain a virtual balanced tree for each fragment. A virtual tree is defined by virtual edges among nodes, where virtual edges are communication paths constructed by the routing algorithm. It follows that for each iteration, fragment merging may increase the number of virtual edges of some node v by $d_G(v)$, where $d_G(v)$ is the degree of v in G. Since Borukva's method takes $O(\log n)$ iteration, the virtual degree of any node v is at most $d_G(v)O(\log n)$. Thus virtual trees communication is feasible via commodity routing by theorem 3.4 which takes $\tau_{mix}(G)2^{O(\sqrt{\log n})}$ rounds. Applying the above approach directly to our setting yields only an MST algorithm running in $\tau_{mix}(G')2^{O(\sqrt{\log n})}$ rounds as mentioned in the last paragraph.

In our setting, to obtain an algorithm running in $\tilde{O}(\log n/\epsilon)$ rounds, we would like to perform routing (only) on R(G) instead of $G \cup R(G)$. However, if we use the same algorithm of [8] in R(G)then during the computing of the MST in G, some node v may become overloaded by $d_G(v)$ virtual edges which causes too much congestion in R(G) (where each node has degree $\Theta(\log n)$ only) when $d_G(v)$ is large. It follows that the routing is infeasible in R(G) and the algorithm fails. To solve the problem we proposed a modified algorithm that uses *aggregate routing*.

The idea of aggregate routing is to perform permutation routing, where some destinations are the same. In other words assume a permutation routing problem where there are only $k \leq n$ (distinct) destinations and t_1, \ldots, t_k and there are *n* sources s_1, \ldots, s_n . Let C_i be the set of sources who have the same destination t_i . In aggregate routing, we would like to aggregate the set of messages in C_i and deliver it to t_i . The aggregate function can be a separable (decomposable) function such as min, max or sum. Then we can modify the permutation routing easily as follows. For a node u that is performing the routing, suppose there are multiple messages arriving at *u* in the same round. Then *u* computes the aggregate of the messages belong to the same set C_i (for every $1 \le i \le k$) and forwards that message according to the routing algorithm. At the destination, the aggregate is computed over any received messages destined for this particular destination. With this intuition, we state the following definitions and lemma.

Definition 3.6 (k-Aggregate Routing). Consider a graph G = (V, E), where nodes are divided into k disjoint partitions C_1, C_2, \ldots, C_k . For each partition C_i , there is a leader l_i , known to all members of C_i . Each node $u \in V$ has one message to deliver to its leader. Let f be a separable aggregate function (such as *min*, *max*, or *sum*). The k-aggregate routing problem is to compute the aggregate f over the nodes in each partition and route it to the corresponding leader of the partition.

We show that *k*-aggregate routing can be solved in the same time bounds as multi-commodity routing.

LEMMA 3.7. Consider a graph G = (V, E) with an instance of k-aggregate routing problem. There is a randomized algorithm that solves the problem in time $\tau_{mix}(G)2^{O(\sqrt{\log n})}$, w.h.p.

PROOF. Let m^u denote the original message at u. We will send the tuple (m^u, l^u) where l^u is the leader of the partition that u belongs to. Let f be the separable aggregate function.

Each node v executes the multi-commodity routing algorithm, with this extra rule: In a round t, suppose v receives multiple messages having the same destination, which is some leader l_i , v computes the aggregate $f_{t,i}$ over those messages, and prepares a tuple $(f_{t,i}, l_i)$. v then forwards the aggregated message to the appropriate next-hop neighbor in the routing path for the next round. vperforms the same reduction for messages targeting other leaders.

It is easy to see that this routing schema is not congested, i.e., it is as fast as the multi-commodity/permutation routing. Observing the local invariance of permutation routing: for each destination u, every node, in each round of the algorithm, sends out at most one message routing towards u. This invariance holds in our kaggregate routing, by the above construction.

At the end of the routing, each leader aggregates over its received messages, which is the aggregate in its partition.

While *k*-aggregate routing can be seen as *upcast* [16], the complementary operation of *downcast*, i.e., sending a message from a source to several destinations, can also be done efficiently as shown below.

LEMMA 3.8 (k-AGGREGATE ROUTING AND DOWNCAST). Consider a graph G = (V, E) with an instance of the k-aggregate routing problem. Furthermore, we require that every member of a partition knows the corresponding aggregate value. There is a randomized algorithm that solve the problem in time $\tau_{mix}(G)2^{O(\sqrt{\log n})}$, w.h.p.

PROOF. Using the algorithm in lemma 3.7, each node also records the source of the incoming messages together with the associated leader ID and round number. Then the routing can be reversed. Starting from the leader of each partition, it sends out the aggregate message with its ID, and each node reverses the aggregate message towards the matching sender. Hence the downcast can be accomplished in the same number of rounds as k-aggregate routing.

We are now ready to implement the MST algorithm.

THEOREM 3.9. Consider a weighted graph G = (V, E) in the ϵ -smoothing model. There exists a randomized distributed algorithm that finds an MST of G in time $\tilde{O}(\frac{1}{\epsilon} + 2^{O(\sqrt{\log n})})$, w.h.p.

PROOF. As discussed in Part 1 (Section 3.1) the algorithm executes $\Theta(\frac{\log n}{\epsilon})$ rounds of random edge selection to construct the random graph R(G) (the graph induced only by the random edges). As shown in Lemma 3.2, R(G) is an expander with constant conductance and hence has $O(\log n)$ mixing time.

We use the permutation routing result of [9] to construct the routing structure on R(G), which allows permutation routing in $\tau_{mix}(R(G))2^{O(\sqrt{\log n})} = O(\log n)2^{O(\sqrt{\log n})} = 2^{O(\sqrt{\log n})}$ rounds.

Our MST algorithm is based on the standard Gallagher-Humblet-Spira (GHS)/Boruvka algorithm, see e.g., [16] which is also used in [8] and many other MST algorithm see e.g., [6, 17]. The main modification compared to the standard GHS algorithm is that growth (diameter) of fragments are controlled during merging (as in controlled GHS algorithm [16]). We summarize the algorithm here and sketch how it is implemented.

Let *T* be the (unique) MST on *G* (we will assume that all weights of edges of *G* are distinct). A *MST fragment* (or simply a *fragment*) *F* of *T* is defined as a connected subgraph of *T*, that is, *F* is a subtree of *T*. An *outgoing edge* of a MST fragment is an edge in *E* where one adjacent node to the edge is in the fragment and the other is not. The *minimum-weight outgoing edge* (*MOE*) of a fragment *F* is the edge with *minimum weight* among all outgoing edges of *F*. As an immediate consequence of the cut property for MST, the MOE of a fragment $F = (V_F, E_F)$ is an edge of the MST.

The GHS algorithm operates in *phases* (see e.g., [16]). In the first phase, the GHS algorithm starts with each individual node as a fragment by itself and continues till there is only one one fragment – the MST. That is, at the beginning, there are |V| fragments, and at the end of the last phase, a single fragment which is the MST. All fragments find their MOE simultaneously in parallel.

In each phase, the algorithm maintains the following invariant: Each MST fragment has a leader and all nodes know their respective parents and children. The root of the tree will be the leader. Initially, each node (a singleton fragment) is a root node; subsequently each fragment will have one root (leader) node. Each fragment is identified by the identifier of its root — called the fragment ID — and each node in the fragment knows its fragment ID.

We describe one phase of the GHS (whp there will be $O(\log n)$ phases as discussed below). Each fragment's operation is coordinated by the respective fragment's root (leader). Each phase consists of two major operations: (1) Finding MOE of all fragments and (2) Merging fragments via their MOEs.

We first describe how to perform the first operation (finding MOE). Let F be the current set of fragments. Each node in V finds its (local) minimum outgoing edge (if any), i.e., an edge to a neighbor belonging to a different fragment that is of least weight. We then execute a |F|-Aggregate Routing and Downcast, using *min* as the aggregate function, with each node being the source and having its fragment leader as its destination. At the end of this step, for each fragment, every member knows the minimum outgoing edge (MOE) of the entire fragment. This MOE edge will be chosen for merging in the second operation (merging fragments). Also, each node keeps the reversed routing paths for further usage.

Once the merging (MST) edges are identified the second operation — merging — is processed. In order to avoid long chains of fragments, a simple randomized trick is used. Each fragment chooses to be a *head* or *tail* with probability 1/2. Only *tail* fragments will merge if their outgoing edge points to a *head* fragment. It can be shown (e.g., see [8] that this merging (still) leads to a constant factor decrease in the number of fragments (on average) and hence the number of phases will be $O(\log n)$ in expectation and with high probability.

We now describe how a merge can be implemented efficiently, There will be no change in the head fragment, but all the tail ones will update to acknowledge the head leader as the new leader. For a tail fragment T, let $v \in T$ be the node that is making the merge, v knows the ID of the head leader by communicating with its neighbor which is a member of the head fragment). v routes this new leader ID to the current leader of T. This is done in parallel by permutation routing. The current leader of T downcasts the new leader ID to all *T* members. This is done via the saved reversed routing paths. The merging is now completed, and time for one iteration is the same as that of permutation routing as before, i.e., $O(\log n)2^{O(\sqrt{\log n})}$.

There are $O(\log n)$ phases and each phase can be implemented in $O(\log n)2^{O(\sqrt{\log n})}$ rounds and hence the total time for Part 2 is $O(\log^2 n)2^{O(\sqrt{\log n})}$.

The total time for MST construction is the number of rounds for Part 1 pluses the number of rounds in Part 2:

$$\Theta(\frac{\log n}{\epsilon}) + O(\log^2 n) 2^{O(\sqrt{\log n})} = \tilde{O}(\frac{1}{\epsilon} + 2^{O(\sqrt{\log n})}).$$

3.3 An Improved Algorithm

We now present an algorithm that is a variant of the previous algorithm and improves upon it. The time complexity of the improved algorithm approaches the lower bound (cf. Section 4). The idea is to use *controlled GHS* algorithm [16] to construct MST fragments of suitable size. Then we apply the smoothing (with a smaller number of rounds compared to previous algorithm, i.e., $\tilde{O}(\frac{1}{\sqrt{\epsilon}})$ instead of $\tilde{O}(\frac{1}{\sqrt{\epsilon}})$) to add an array day area to a sume map.

 $\tilde{O}(\frac{1}{\epsilon})$) to add an expander over the super-graph induced by the MST fragments where each super-node is one fragment (partition). Then we compute the final MST in a similar fashion to Theorem 3.9 on the super-graph.

THEOREM 3.10. Given a weighted graph G(V, E) in the ϵ -smoothing model. Then there exists a randomized distributed algorithm that finds an MST of G in time $\tilde{O}(\frac{1}{\sqrt{\epsilon}})2^{O(\sqrt{\log n})}$, w.h.p.

PROOF. We give the algorithm along with its analysis, as follows. Run $O(\frac{\log n}{\sqrt{\epsilon}})$ rounds of smoothing. Denote by *S* the set of smoothed edges generated. Using Lemma 3.1, the probability that a smoothed edge occurs between two nodes in *G* is $p = \Theta(\frac{\sqrt{\epsilon} \log n}{n})$.

Run *controlled GHS* [16] for $\log \frac{1}{\sqrt{\epsilon}}$ phases. This takes $O(\frac{\log n}{\sqrt{\epsilon}})$ rounds. Every cluster (each of which is an MST fragment) will have size $\Omega(\frac{1}{\sqrt{\epsilon}})$ and diameter $O(\frac{1}{\sqrt{\epsilon}})$, and there will be $O(n\sqrt{\epsilon})$ such clusters [16, Section 7.4]. We call these clusters as *base fragments*. We note that communication within a cluster (i.e., between any node of the cluster and its leader) takes $O(\frac{1}{\sqrt{\epsilon}})$ rounds.

View these clusters as a set of *super-nodes*, denoted by V'. Let $E' \subset E$ be the set of inter-super-node edges, let $S' \subset S$ be the set of inter-super-node smoothed edges. Consider two super-graphs: G'(V', E') and R'(V', S'). It is easy to show that due to the probability p of the random edges introduced by the *smoothing* process, the super-graph R'(V', S') is an Erdős–Rényi random graph or a G(n', p')-random graph, where

$$n' = O(n\sqrt{\epsilon}) \tag{1}$$

and

$$p' \ge \Omega\left(\left(\frac{1}{\sqrt{\epsilon}}\right)^2\right)p = \Omega\left(\frac{\log(n')}{n'}\right)$$
 (2)

Thus, the super-graphs $G' \cup R'$ is equivalent to the smoothed graph of our model in Section 2.

Similar to the previous algorithm, we solve the MST problem using the Boruvka's algorithm on the super-graph G', using the

routing structure of Ghaffari et al. [8, 9] and the aggregate routing over R'. To implement the algorithm on the super-graph, we pipeline messages within all the super-nodes, i.e., inside the base fragments. There are $O(\log n)$ phases of the Boruvka's algorithm and each phase takes $O(\frac{1}{\sqrt{\epsilon}}) \cdot 2^{O(\sqrt{\log n})}$ rounds. The extra term of $O(\frac{1}{\sqrt{\epsilon}})$ is incurred by communication within a super-node. Thus, in total, the second part takes $O(\frac{1}{\sqrt{\epsilon}} \cdot \log n) \cdot 2^{O(\sqrt{\log n})}$ rounds.

Combining the MST edges over the super-graph G', and the MST edges in each super-node, we have the MST for the original graph. Therefore, the total time complexity of the algorithm is $O(\frac{\log n}{\sqrt{\epsilon}}) + O(\frac{\log n}{\sqrt{\epsilon}}) \cdot 2^{O(\sqrt{\log n})} = \tilde{O}(\frac{1}{\sqrt{\epsilon}}) \cdot 2^{O(\sqrt{\log n})}.$

4 LOWER BOUND

In this section, we show the following lower bound result on the ϵ -smoothing model. We note that $\tilde{\Omega}(D + \sqrt{n})$ is an unconditional lower bound (without smoothing) that holds even for randomized Monte-Carlo approximate MST algorithms[22].

THEOREM 4.1 (SMOOTH MST LOWER BOUND). There exists a family of graphs \mathcal{G} , such that, under the ϵ -smoothing model, any distributed MST algorithm must incur a running time of $\tilde{\Omega}(\frac{1}{\sqrt{\epsilon}})$, in expectation.

We will prove the lower bound theorem by using the technique used in [22]. First, we will briefly recall the lower bound poof of $\tilde{\Omega}(\sqrt{n})$ (we assume $D = O(\log n)$) without smoothing. For purposes of exposition, we simplify and slightly modify the technique in the mentioned paper, to show only the bound for exact distributed MST. Then we extend it to the smoothing model. The procedure is to establish a chain of algorithm reductions which is the same as in [22], such that it relates distributed MST to a problem with a known lower bound. The following are the chain of reductions:

• Set Disjointness (SD) to Distributed Set Disjointness (DSD). We first reduce the set disjointness (SE) verification problem, a standard well-studied problem in two-party communication complexity to the problem of distributed set disjointness (DSD) verification. In the set disjointness problem (SD), we have two parties Alice and Bob, who each have a *k*-bit string $-x = (x_1, x_2, \ldots, x_k)$ and $y = (y_1, y_2, \ldots, y_k)$ respectively. The goal is to verify if the set disjointness function is defined to be one if the inner product $\langle x, y \rangle$ is 0 (i.e., there is no *i* such that $x_i = y_i = 1$) and zero otherwise. The goal is to solve SD by *communicating as few bits* as possible between Alice and Bob.

In the distributed set disjointness (DSD) verification, the goal is to solve SD in a given input graph G = (V, E), where two distinguished nodes $s, t \in V$ have the bit vectors x and y respectively. In other words, instead of communicating directly as in the two-party problem (between Alice and Bob), the two nodes s and t (standing respectively for Alice and Bob) have to communicate via the edges of G (in the CONGEST model) to solve SD. The goal is to solve the DSD problem using *as few rounds* as possible in the CONGEST model (where only $O(\log n)$ bits per edge per round are allowed).

- Reduction of Distributed set disjointness (DSD) to connected spanning subgraph (CSS) verification. In the CSS problem, we want to solve a graph verification problem which can be defined as follows. In distributed graph verification, we want to efficiently check whether a given subgraph of a network has a specified property via a distributed algorithm. Formally, given a graph G = (V, E), a subgraph H = (V, E')with $E' \subseteq E$, and a predicate Π , it is required to decide whether H satisfies Π (i.e., when the algorithm terminates, every node knows whether *H* satisfies Π). The predicate Π may specify statements such as "H is connected" or "H is a spanning tree" or "*H* contains a cycle". Each vertex in *G* knows which of its incident edges (if any) belong to H. In the connected spanning subgraph (CSS) verification the goal is to verify whether the given subgraph H is connected and spans all nodes of G, i.e., every node in G is incident to some edge in H. The goal is to solve CCS in as few rounds
- as possible (in the CONGEST model).The last reduction is to reduce CCS verification to computing an MST.

The last two reductions above are done exactly as in the paper of [22] that show the time lower bound of $\tilde{\Omega}(D + \sqrt{n})$ rounds; however, the first reduction from SD to DSD is different in the smoothing model, as the input graph used in the DSD can use the (additional) power of the smoothing model.

We will first briefly discuss the reductions as in [22] and then discuss how to modify the first reduction to work for the smoothing model. Here we first state the bounds that we obtain via these reductions, and refer to [22] for the details. The well-known communication complexity lower bound for the SD problem is $\Omega(k)$ (see e.g.,[14]), where k is the length of the bit vector of Alice and Bob. This lower bound holds even for randomized Monte-Carlo algorithms and even under *shared* public coin.

Due to the graph topology used in the DSD problem (see Figure 1 without the smoothing edges) the value of *k* is to be $\Theta(\sqrt{n})$ (which is the best possible). The reduction from SD to DSD shows that the lower bound for DSD, is $\tilde{\Omega}(\sqrt{n})$ rounds. (Note that the diameter of the lower bound graph is $O(\log n)$ and hence subsumed.) This reduction uses the *Simulation Theorem* (cf. Theorem 3.1 in [22]) which is explained later below.

The reduction from DSD to CCS shows that the same time lower bound of $\tilde{\Omega}(\sqrt{n})$ rounds holds for the CCS problem. This reduction shows that the given subgraph *H* in the CCS problem is spanning connected if and only if the input vectors *x* and *y* are disjoint.

The reduction from CCS to MST problem shows that the time lower bound for CCS verification which is $\tilde{\Omega}(\sqrt{n})$ also holds for the MST problem. This reduction takes as input the CCS problem and assigns weight 1 to the edges in the subgraph *H* and weight *n* to all other edges in *G*. It is easy to show that *H* is spanning connected if and only if the weight of the MST is less than *n*. Hence the same time lower bound that holds for CCS also holds for MST.

4.1 Reduction of SD to DSD: The Simulation Theorem

In this section, we explain the key reduction from SD to DSD which uses the *Simulation Theorem* as in [22]. The reduction idea is as



Figure 1: The lower bound graph.

follows. Assuming that we have an algorithm for DSD that finished in *r* rounds, Alice and Bob will simulate this algorithm in the two party model by sending as few bits as possible. The Simulation theorem accomplishes this. The Simulation theorem in [22] et al that shows the simulation in the standard graph (without smoothing) uses constant number of bits per round to do the simulation. Thus if the DSD algorithm finishes in *r* rounds, then the Alice and Bob would have solved SD by exchanging O(r) bits. If r = o(k) (where *k* is the length of the input bit string), then this will contradict the lower bound of set disjointness in the two party model which is $\Omega(k)$ (even for randomized algorithms).

We note that the reduction is similar to that in [22] as the input graph G(x, y) for the DSD is the same (see Figure 1). However, in the smoothing model, the algorithm has the additional power of using the smoothing edges and hence the lower bound will be smaller as we will show below. We will first briefly describe the idea behind the Simulation Theorem as it applicable to *G* without the smoothing. The lower bound graph G(x, y) used in the Simulation Theorem is as follows. Note that the input graph *G* for DSD has two distinguished nodes *s* and *t* that have the inputs *x* and *y* (corresponding to Alice and Bob) respectively.

4.1.1 The lower bound graph (family) for MST. G(x, y) is depicted as in Figure 1, where $|x| = |y| = \sqrt{n}$. G(.) includes \sqrt{n} paths, and a full binary tree (to reduce the diameter to $O(\log n)$). Each path has length of \sqrt{n} ; these are called the path edges. The full binary tree has \sqrt{n} leaves, and hence, has the height of $p = \frac{\log n}{2}$. We number the leaves and the path nodes from left to right $-0, 1, \ldots, \ell$. Note that leaf numbered 0 is node *s* and leaf numbered ℓ is *t*. Consider each leaf $0 < j < \ell$, let it connect to all the nodes *j* in all paths, we call these spoke edges. Note that the binary tree edges, the path edges, and the spoke edges from leaf nodes other than *s* and *t* are present in every graph of the family. It is straightforward that G(x, y) has $\Theta(n)$ nodes, and diameter $D = \Theta(\log n)$.

In the reduction from SD to DSD, Alice and Bob wants to solve SD problem with Alice having input vector $x = (x_1, x_2, \ldots, x_k)$ and Bob having input vector $y = (y_1, y_2, \ldots, y_k)$ where we fix $k = \sqrt{n}$. Depending on x and y, Alice and Bob will fix the spoke edges from nodes s and t. For the SD problem, Alice will add edges from s to the first node in path i if and only if $x_i = 0$. Similarly Bob will add edges from t to the last node in the path i if and only if y_i is 0.

Now we are ready to describe the Simulation Theorem which really gives an algorithm for Alice and Bob to simulate any given algorithm for DSD problem. If the DSD algorithm runs in r rounds, the Simulation theorem will show how to solve the SD problem by exchanging O(r) bits. We sketch the main idea here, which is quite simple, and leave the full details which can be found in [22].

How will Alice and Bob start the simulation? Note that they want to solve the SD problem on their respective inputs x and y in the two-party model. They will then use their respective inputs to construct G(x, y) as described above. Note that Alice will be able to construct all edges of G except the spoke edges of t (since that depends on Bob's input) and vice versa for Bob. Then, assuming that there is an algorithm for the DSD problem on the same inputs that runs in r rounds, Alice and Bob will simulate the DSD algorithm whose output will also give the output for the SD problem (by definition).

The main idea is for Alice and Bob to keep the simulation going as long as possible. If one *disregards the binary tree edges*, all paths are of length $\ell = \Theta(\sqrt{n})$ and hence it is easy to keep the simulation going for $\ell/2$ rounds (say). This is because, Alice has all the information needed to simulate the DSD algorithm till $\ell/2$ steps (i.e., the middle of the path). Why? Because Alice knows her own input and all other nodes in *G* does not have any input. She does not know Bob's input, but does not matter for $\ell/2$ steps since in so many rounds nothing from Bob's part of the graph reaches the "middle" of the path. But of course, the above is not true because of the binary tree edges which has smaller diameter. So to keep the simulation going for Bob, Alice sends the minimum amount of information needed by him. Note that after *i* rounds the computation from Alice's side (which are the set of nodes numbered u_0^p and

 $v_0^1, v_0^2, \ldots, v_0^{\sqrt{n}}$ will have reached nodes at distance *i* on the path. We define the R_i (intuitively *i*-Right) set as follows: R_i includes all nodes on the paths with subscript $j \ge i$, all leaf nodes u_j^p where $j \ge i$, and all ancestor of these leaves, see Figure 1.

In round *i*, Bob needs to keep the correct computation for R_i . To achieve that, Alice sends only the messages sent by the tree nodes in R_{i-1} crossing into the tree nodes in R_i (see Lemma 3.4 in [22]). (A similar observation applies for Alice to do her simulation). Hence only messages sent by at most $O(\log n)$ nodes in the binary tree are needed. Hence in every round at most $O(\log^2 n)$ bits need to be exchanged by Alice and Bob to keep the simulation going. Thus if the DSD algorithm finishes in $o(\ell/\log^2 n) = o(\sqrt{n}/\log^2 n)$ rounds, then the simulation will also end successfully.

Thus we can show the following Simulation Theorem.

THEOREM 4.2 (SIMULATION THEOREM). (Simplified version of Theorem 3.1 in [22]) Given the DSD problem with input size $\Theta(\sqrt{n})$, encoded as G(x, y) (i.e., the x and y are bit vectors of length $\Theta(\sqrt{n})$), if there is a distributed algorithm that solves the DSD problem in time at most T rounds, using messages of size $O(\log n)$. Then there is an algorithm in the two-party communication complexity model that decides SD problem while exchanging at most $O(T \log^2 n)$ bits.

4.2 Lower bound with smoothing

Under the ϵ -smoothing model, we will use the same lower bound graph G(x, y). However, the smoothing model gives the algorithm additional power to add random edges in G(x, y) during the course of the simulation. We will show how to modify the Simulation Theorem to apply to the smoothing model. Naturally, since the algorithm has additional power, it can finish faster, and hence the corresponding lower bound will be smaller.

We focus on Bob and show how he can keep his simulation going. (A similar argument applies for Alice.) We consider how Bob maintains correct states for R_i in round *i*. Beside the required messages as discussed in Theorem 4.2, Bob needs to know the messages sent over (potential) *smoothed edges* crossing $V \setminus R_i$ into R_i . Since there are more messages to keep track, Bob cannot keep the simulation longer than that of the non-smoothing case. Let δ be the number of rounds the simulation is valid without exceeding $\Theta(\sqrt{n})$ bits of communication. Since the smoothing process is randomized, we bound δ in expectation.

We will use a pessimistic estimation to estimate the number of smoothed edges that "affect" Bob from Alice's side. Let $C_i = V \setminus R_i$, let S_i be the expected number of smoothed edges crossing C_i and R_i . S_i indicates the number of extra messages Bob needs to know, in round *i*. Notice that $|C_i| = i\Theta(\sqrt{n})$.

$$S_i = 2\epsilon i\Theta(\sqrt{n})\frac{(n-i\Theta(\sqrt{n}))}{n}$$
(3)

Since $\delta < \Theta(\sqrt{n})$, for every round *i*, $S_i = \Theta(\epsilon \delta \sqrt{n})$. After δ rounds, the expected number of messages over the smoothed edges is thus: $\Theta(\epsilon \delta^2 \sqrt{n})$. Also, by Theorem 4.2, we need to keep track of $(\delta \log n)$ messages. To stay within the budget of \sqrt{n} bits for DSD communication we require: $\epsilon \delta^2 \Theta(\sqrt{n}) + \delta \Theta(\log n) < \frac{\sqrt{n}}{2}$.

communication, we require: $\epsilon \delta^2 \Theta(\sqrt{n}) + \delta \Theta(\log n) \le \frac{\sqrt{n}}{B}$. With $B = \Theta(\log n)$ (the message size), we have $\delta \le \Theta(\frac{1}{\sqrt{\epsilon \log n}})$. Thus, we can keep the simulation going for up to $\Theta(\frac{1}{\sqrt{\epsilon \log n}})$ rounds.

To complete the lower bound argument, the same lower bound applies for MST in the ϵ -smoothing model by the chain of reductions.

5 OTHER SMOOTHING MODELS

In this section, we discuss some of the other plausible smoothing models for the distributed MST problem. The most natural smoothing model that comes into the mind first in respect to a numerical-valued computing problem and that is similar in spirit to the original Spielman-Teng smoothing [23] is where one "perturbs" the edge-weights in the given input graph. However, as we have already noted (see Section 1) this may not make for anything interesting if the perturbations are too small with respect to the original edge-weights.

These considerations motivate us to explore an alternate avenue where the *graph topology* rather than the edge-weights are perturbed. In particular, we consider smoothing models where the perturbation process adds more edges to the original input graphs.Below we describe two such models where additional edges characterize the perturbed graph compared to the original input graph. We call these models the *k*-smoothing models. We distinguish these two models based on whether or not the smoothed edges are known to the algorithm.

5.1 The *k*-smoothing model with known smoothed edges

The main model that we follow in this paper, the ϵ -smoothing model (see Section 2.2) adds smooth edges by node-local computation during the course of an algorithm. We can look at models where smooth edges are added all at once, by some external process, prior to the commencement of the algorithm.

- Consider a smoothing model *M*(*k*, *) where the "perturbation" process *adds k* additional edges to the original input graph. In our notation, * denotes the fact that the smoothed (i.e., additional) edges are known to the algorithm. These *k* additional edges are chosen uniformly at random from all the {n / 2} possible edges of the graph.
- (2) Consider another smoothing model M(δ, *) where the "perturbation" process adds – independently for each of the ⁿ₂ possible edges – an edge with probability δ.

Remark 2. It is not difficult to see that the model $\mathcal{M}(\epsilon, *)$ is essentially equivalent to the model $\mathcal{M}(k, *)$ for the case when $\epsilon = \frac{k}{n}$.

Remark 3. By Lemma 3.1, consider the ϵ -smoothing model with ℓ rounds of smoothing, then: $\mathcal{M}(k = 2\ell \epsilon n, *)$ and $\mathcal{M}(\delta = \frac{2\ell \epsilon}{n}, *)$ are the equivalent models.

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5.2 The *k*-smoothing model with unknown smoothed edges

Essentially, we have the counterparts of $\mathcal{M}(k, *)$ and $\mathcal{M}(\delta, *)$ for the particular case when the smoothed edges are *not* known to the algorithm. We denoted these models $\mathcal{M}(k, \times)$ and $\mathcal{M}(\delta, \times)$.

We also note that both $\mathcal{M}(k, \times)$ and $\mathcal{M}(\epsilon, \times)$ are essentially equivalent to the smoothing model proposed by Dinitz et al. [5], where the new, perturbed input graph is chosen uniformly at random from the set of all possible graphs whose edge-sets are at a (positive) edit-distance *k* from the original input graph. The only subtle difference is that, in their model [5], the edit-distance can be positive as well as negative. We, however, consider only *positive* edit-distances here.

We note that the algorithms specified in this paper do not (at least directly) work when the smoothing edges are not known. However, note that the lower bound holds (for appropriate choice of k in terms of ϵ).

6 CONCLUSION

In this paper, we study smoothed analysis of distributed graph algorithms focusing on the well-studied distributed MST problem. Our work can be considered as a first step in understanding the smoothed complexity of distributed graph algorithms.

We present a smoothing model, and upper and lower bounds for the time complexity of distributed MST in this model. These bounds quantify the time bounds in terms of the smoothing parameter ϵ . The bounds are within a factor of $2^{O(\sqrt{\log n})}$ polylog *n* and a key open problem is whether this gap can be closed.

While we focus on one specific smoothing model, our results also apply to other related smoothing models (discussed in Section 5). A commonality among these models, besides adding random edges, is that the added edges are *known* to the nodes. This knowledge of the random edges are crucial to obtaining our upper bounds. Of course, our lower bounds apply *regardless of* this knowledge.

An important open problem is to show non-trivial bounds when the random edges are *unknown* to the nodes; i.e., the input graph consists of the original graph G plus the random edges and the nodes cannot distinguish between edges in G and the added random edges.

It would also be interesting to explore other fundamental distributed graph problems such as leader election, shortest paths, minimum cut etc., in the smoothing model.

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