

A fast distributed approximation algorithm for minimum spanning trees

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Received: 31 December 2006 / Accepted: 21 August 2007 / Published online: 30 October 2007
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Abstract We present a distributed algorithm that constructs an $O(\log n)$ -approximate minimum spanning tree (MST) in any arbitrary network. This algorithm runs in time $\tilde{O}(D(G) + L(G, w))$ where $L(G, w)$ is a parameter called the *local shortest path diameter* and $D(G)$ is the (unweighted) diameter of the graph. Our algorithm is existentially optimal (up to polylogarithmic factors), i.e., there exist graphs which need $\Omega(D(G) + L(G, w))$ time to compute an H -approximation to the MST for any $H \in [1, \Theta(\log n)]$. Our result also shows that there can be a significant time gap between exact and approximate MST computation: there exists graphs in which the running time of our approximation algorithm is exponentially faster than the *time-optimal* distributed algorithm that computes the MST. Finally, we show that our algorithm can be used to find an approximate MST in wireless networks and in random weighted networks in almost optimal $\tilde{O}(D(G))$ time.

Keywords Minimum spanning tree · Distributed approximation algorithm · Randomized algorithm

1 Introduction

1.1 Background and previous work

The distributed minimum spanning tree (MST) problem is one of the most important problems in the area of distributed

computing. There has been a long line of research to develop efficient distributed algorithms for the MST problem starting with the seminal paper of Gallager et al. [8] that constructs the MST in $O(n \log n)$ time and $O(|E| + n \log n)$ messages. The communication (message) complexity of the Gallager et al. algorithm is optimal, but its time complexity is not. Hence further research concentrated on improving the time complexity. The time complexity was first improved to $O(n \log \log n)$ by Chin and Ting [2], further improved to $O(n \log^* n)$ by Gafni [7], and then improved to *existentially optimal* running time of $O(n)$ by Awerbuch [1]. The $O(n)$ bound is existentially optimal in the sense that there exists graphs where no distributed MST algorithm can do better than $\Omega(n)$ time. This was the state of the art till the mid-nineties when Garay, Kutten, and Peleg [9] raised the question of identifying graph parameters that can better capture the complexity (motivated by “universal” complexity) of distributed MST computation. For many existing networks G , their diameter $D(G)$ (or D for short) is significantly smaller than the number of vertices n and therefore is a good candidate to design protocols whose running time is bounded in terms of $D(G)$ rather than n . Garay, Kutten, and Peleg [9] gave the first such distributed algorithm for the MST problem with running time $O(D(G) + n^{0.61})$, which was later improved by Kutten and Peleg [16] to $O(D(G) + \sqrt{n} \log^* n)$. Elkin [4] refined this result further and argued that a parameter called “MST-radius” captures the complexity of distributed MST algorithms better. He devised a distributed protocol that constructs the MST in $\tilde{O}(\mu(G, w) + \sqrt{n})$ time, where $\mu(G, w)$ is the “MST-radius” of the graph [4] (is a function of the graph topology as well as the edge weights). The ratio between diameter and MST-radius can be as large as $\Theta(n)$, and consequently, on some inputs, this protocol is faster than the protocol of [16] by a factor of $\Omega(\sqrt{n})$. However, a drawback of this protocol (unlike the previous MST protocols

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[16,9,2,7,8]) is that it cannot detect the termination of the algorithm in that time (unless $\mu(G, w)$ is given as part of the input). Finally, we note that the time-efficient algorithms of [16,4,9] are not message-optimal [i.e., they require asymptotically much more than $O(|E| + n \log n)$ messages, e.g., the protocol of [16] takes $O(|E| + n^{1.5})$ messages].

The lack of progress in improving the result of [16], and in particular breaking the \sqrt{n} barrier, led to work on lower bounds for the distributed MST problem. Peleg and Rabinovich [21] showed that $\tilde{\Omega}(\sqrt{n})$ time is required for constructing an MST even on graphs of small diameter (for any $D = \Omega(\log n)$) and showed that this result establishes the asymptotic near-tight (existential) optimality of the protocol of [16]. Faster algorithms can be developed for networks with very small constant diameter. An $O(\log \log n)$ time algorithm for networks with $D = 1$ was given in [19], and an $O(\log n)$ time algorithm for networks with $D = 2$ was given in [18]. However, there exist a significant gap in the time complexity of the distributed MST algorithms between the cases of network diameters 2 and 3. In [18], it was shown that the time complexity of any distributed MST algorithm is $\Omega(\sqrt[4]{n}/\sqrt{B})$ for networks of diameter 3 and $\Omega(\sqrt[3]{n}/\sqrt{B})$ for networks of diameter 4.

While the previous distributed protocols deal with computing the exact MST, the next important question addressed in the literature concerns the study of distributed *approximation* of the MST, i.e., constructing a spanning tree whose total weight is near-minimum. From a practical perspective, given that the construction of an MST can take as much as $\tilde{\Omega}(\sqrt{n})$ time, it is worth investigating whether one can design distributed algorithms that run faster and output a near-minimum spanning tree. Peleg and Rabinovich [21] was one of the first to raise the question of devising faster algorithms that construct an approximation to the MST and left it open for further study. To quote their paper: “To the best of our knowledge, nothing nontrivial is known about this problem. . .”. Since then, the most important new result is the *hardness* results shown by Elkin [6]. This result shows that *approximating* the MST problem on graphs of small diameter (e.g., $O(\log n)$) within a ratio H requires essentially $\Omega(\sqrt{n}/HB)$ time (assuming B bits can be sent through each edge in one round), i.e., this gives a time-approximation trade-off for the distributed MST problem: $T^2 H = \Omega(\sqrt{n}/B)$. However, not much progress has been made on designing time-efficient distributed approximation algorithms for the MST problem. To quote Elkin’s survey paper [5]: “There is no satisfactory approximation algorithm known for the MST problem”. To the best of our knowledge, the only known distributed approximation algorithm for the MST problem is given by Elkin in [6]. This algorithm gives an H -approximation to the MST with running time $O(D(G) + \frac{\omega_{\max}}{H-1} \cdot \log^* n)$, where ω_{\max} is the ratio between the maximum and minimum weights of the edges in the input graph G . Thus, this algorithm is not

independent of the edge weights and its running time can be quite large.

1.2 Distributed computing model and our results

We present a fast distributed approximation algorithm for the MST problem. First, we briefly describe the distributed computing model that is used by our algorithm (as well as the previous MST algorithms [2,8,9,16,1,7,4] mentioned above) which is now standard in the distributed computing literature (see e.g., the book by Peleg [20]).

Distributed computing model. We are given a network modeled as an undirected weighted graph $G = (V, E, w)$ where V is the set of the nodes (vertices) and E is the set of the communication links between them and $w(e)$ is the weight of the edge $e \in E$. Without loss of generality, we assume that G is connected. Each node hosts a processor with limited initial knowledge. Specifically, we make the common assumption that each node has unique identity numbers (this is not really essential, but simplifies presentation) and at the beginning of computation, each vertex v accepts as input its own identity number and the weights of the edges adjacent to v . Thus, a node has only *local* knowledge limited to itself and its neighbors. The vertices are allowed to communicate through the edges of the graph G . We assume that the communication is synchronous and occurs in discrete pulses (time steps). (This assumption is not essential for our time complexity analysis. One can use a *synchronizer* to obtain the same time bound in an asynchronous network at the cost of some increase in the message complexity [20].) In each time step, each node v can send an arbitrary message of size $O(\log n)$ through each edge $e = (v, u)$ that is adjacent to v , and the message arrives at u by the end of this time step. (If unbounded-size messages are allowed, the MST problem can be trivially solved in $O(D(G))$ time [20].) The weights of the edges are at most polynomial in the number of vertices n , and therefore, the weight of a single edge can be communicated in one time step. This model of distributed computation is called the CONGEST($\log n$) model or simply the CONGEST model [20] (the previous results on the distributed MST problem cited in Sect. 1.1 are for this model). We note that more generally, CONGEST(B) model allows messages of size at most $O(B)$ to be transmitted in a single time step across an edge. Our algorithm can straightforwardly be applied to this model also. We will assume $B = \log n$ throughout this paper.

Overview of the results. Our main contribution is an almost existentially optimal (in both time and communication complexity) distributed approximation algorithm that constructs an $O(\log n)$ -approximate minimum spanning tree, i.e., whose cost is within an $O(\log n)$ factor of the MST. The running

time¹ of our algorithm is $\tilde{O}(D(G) + L(G, w))$, where $L(G, w)$ is a parameter called the *local shortest path diameter* (we defer the definition of $L(G, w)$ to Sect. 2.2). Like the MST-radius, $L(G, w)$ depends both on the graph topology as well as on the edge weights. $L(G, w)$ always lies between 1 and n . The parameter $L(G, w)$ can be smaller or larger than the diameter and typically it can be much smaller than \sqrt{n} [recall that this is essentially a lower bound on distributed (exact) MST computation]. In fact, we show that there exist some graphs for which any distributed algorithm for computing an MST will take $\tilde{\Omega}(\sqrt{n})$ time, while our algorithm will compute a near-optimal MST in $\tilde{O}(1)$ time, since $L(G, w) = \tilde{O}(1)$ and $D = \tilde{O}(1)$ for these graphs. Thus there exists an exponential gap between exact MST and $O(\log n)$ -approximate MST computation. However, in some graphs $L(G, w)$ can be asymptotically larger than both the diameter and \sqrt{n} . By combining the MST algorithm of Kutten and Peleg [16] with our algorithm in an obvious way, we can obtain an algorithm with the same approximation guarantee but with running time $\tilde{O}(D(G) + \min(L(G, w), \sqrt{n}))$.

The parameter $L(G, w)$ is not arbitrary. We show that it captures the hardness of distributed approximation quite precisely: there exists a family of n -vertex graphs where $\Omega(L(G, w))$ time is needed by any distributed approximation algorithm to approximate the MST within an H -factor, $1 \leq H \leq O(\log n)$ (cf. Theorem 5). This implies that our algorithm is existentially optimal (upto a polylogarithmic factor) and in general, no other algorithm can do better. We note that the existential optimality of our algorithm is with respect to $L(G, w)$ instead of n as in the case of Awerbuch's algorithm [1]. Our algorithm is also existentially optimal (upto a polylogarithmic factor) with respect to the communication (message) complexity—it takes $\tilde{O}(|E|)$ messages—since $\Omega(|E|)$ messages are clearly needed in some graphs to construct any spanning tree [15, 23].

One of the motivations for this work is to investigate whether a fast distributed algorithm that constructs a (near-optimal) MST can be developed for some special classes of networks. An important consequence of our results is that networks with low $L(G, w)$ value (compared to $O(D(G))$) admit a distributed $\tilde{O}(D(G))$ time $O(\log n)$ -approximation algorithm. In particular, unit disk graphs have $L(G, w) = 1$. Unit disk graphs are commonly used to model wireless networks. We also show that $L(G, w) = O(\log n)$ with high probability in any arbitrary network whose edge weights are chosen independently at random following any distribution with constant mean (cf. Theorem 8).

2 Distributed approximate MST algorithm

2.1 Nearest neighbor tree scheme

The main objective of our approach is to construct a spanning tree, called the *Nearest Neighbor Tree (NNT)*, efficiently in a distributed fashion. In our previous work [13], we introduced the Nearest Neighbor Tree and showed that its cost is within an $O(\log n)$ factor of the cost of the MST. The scheme is used to construct an NNT (henceforth called *NNT scheme*) as follows: (1) each node first chooses a unique *rank* from a totally ordered set; a ranking of the nodes corresponds to a permutation of the nodes; (2) each node (except the one with the highest rank) connects (via the *shortest path*) to the *nearest* node of higher rank. We show that the NNT scheme constructs a spanning subgraph in any weighted graph whose cost is at most $O(\log n)$ times that of the MST, irrespective of how the ranks are selected (as long as they are distinct). Note that some cycles can be introduced in step 2, and hence to get a spanning tree we need to remove some edges to break the cycles. Our NNT scheme is closely related to the approximation algorithm for the *traveling salesman problem* (coincidentally called Nearest Neighbor algorithm) analyzed in a classic paper of Rosenkrantz, Lewis, and Stearns [22]. Imase and Waxman [12] also used a scheme based on [22] (their algorithm can also be considered a variant of the NNT scheme) to show that it can maintain an $O(\log n)$ -approximate Steiner tree dynamically (assuming only node additions, but not deletions.) However, their algorithm will not work in a distributed setting (unlike our NNT scheme) because one cannot connect to the nearest node (they can do that since the nodes are added one by one) as this can introduce cycles.

The main advantage of the NNT scheme is that each node, individually, has the task of finding its nearest node of higher rank to connect to, and hence no explicit coordination is needed among the nodes. However, despite the simplicity of the NNT scheme, it is not clear how to efficiently implement the scheme in a general weighted graph. In our previous work [13], we showed how the NNT scheme can be implemented in a *complete metric* graph G (i.e., $D(G) = 1$). This algorithm takes only $O(n \log n)$ messages to construct an $O(\log n)$ -approximate MST as opposed to the $\Omega(n^2)$ lower bound (shown by Korach et al. [14]) on the number of messages needed by any distributed MST algorithm in this model. If the time complexity needs to be optimized, then the NNT scheme can easily be implemented in $O(1)$ time by using $O(n^2)$ messages, as opposed to the best known time bound of $O(\log \log n)$ for the (exact) MST [17]. These results suggest that the NNT scheme can yield faster and more communication-efficient algorithms than the algorithms that compute the exact MST. However, an efficient implementation in a general weighted graph is non-trivial and was

¹ We use the notations $\tilde{O}(f(n))$ and $\tilde{\Omega}(f(n))$ to denote $O(f(n) \cdot \text{polylog}(f(n)))$ and $\Omega(f(n)/\text{polylog}(f(n)))$, respectively.

left open in [13]. Thus, a main contribution of this paper is an efficient implementation of the scheme in a general network. The main difficulties are avoiding congestion when finding the nearest node of higher rank efficiently in a distributed fashion (since many nodes are trying to search at the same time) and avoiding cycle formation. We use a technique of “incremental” neighborhood exploration that avoids congestion and cycle formation, and is explained in detail in Sect. 2.3.

2.2 Preliminaries

We use the following definitions and notations concerning an undirected weighted graph $G = (V, E, w)$. We say that u and v are *neighbors* of each other if $(u, v) \in E$.

Notations:

- $|Q(u, v)|$ or simply $|Q|$ —is the number of edges in path Q from u to v . We call $|Q|$ the *length of the path Q* .
- $w(Q(u, v))$ or $w(Q)$ —is the weight of the path Q , which is defined as the sum of the weights of the edges in path Q , i.e., $w(Q) = \sum_{(x,y) \in Q} w(x, y)$.
- $P(u, v)$ —is a shortest path (in the weighted sense) from u to v .
- $d(u, v)$ —is the (weighted) *distance* between u and v , and defined by $d(u, v) = w(P(u, v))$.
- $N_\rho(v)$ —is the set of all *neighbors* of v within distance ρ , i.e., $N_\rho(v) = \{u \mid (u, v) \in E \wedge w(u, v) \leq \rho\}$.
- $W(v)$ —is the weight of the largest edge adjacent to v , e.g., $W(v) = \max_{(v,x) \in E} w(v, x)$.
- $l(u, v)$ —is the number of edges in the minimum-length shortest path from u to v . Note that there may be more than one shortest path from u to v . Thus, $l(u, v)$ is the number of edges of the shortest path having the least number of edges, i.e.,

$$l(u, v) = \min\{|P(u, v)| \mid P(u, v) \text{ is a shortest path}\}.$$

Definition 1 (ρ -neighborhood) ρ -neighborhood of a node v , denoted by $\Gamma_\rho(v)$, is the set of the nodes that are within distance ρ from v . $\Gamma_\rho(v) = \{u \mid d(u, v) \leq \rho\}$.

Definition 2 $((\rho, \lambda)$ -neighborhood) (ρ, λ) -neighborhood of a node v , denoted by $\Gamma_{\rho, \lambda}(v)$, is the set of all nodes u such that there is a path $Q(v, u)$ such that $w(Q) \leq \rho$ and $|Q| \leq \lambda$. Clearly, $\Gamma_{\rho, \lambda}(v) \subseteq \Gamma_\rho(v)$.

Definition 3 (Shortest path diameter (SPD)) SPD is denoted by $S(G, w)$ (or S for short) and defined by

$$S = \max_{u, v \in V} l(u, v).$$

Definition 4 (Local Shortest Path Diameter (LSPD)) LSPD is denoted by $L(G, w)$ (or L for short) and defined by $L = \max_{v \in V} L(v)$, where $L(v) = \max_{u \in \Gamma_{W(v)}(v)} l(u, v)$.

Notice that $1 \leq L \leq S \leq n$ in any graph. However, there exists graphs, where L is significantly smaller than both S and the (unweighted) diameter of the graph, D . For example, in a chain of n nodes (all edges with weight 1), $S = n$, $D = n$, and $L = 1$.

2.3 Distributed NNT algorithm

We recall the basic NNT scheme as follows. Each node v selects a unique rank $r(v)$. Then each node finds the nearest node of higher rank and connects to it via the shortest path. Now we describe each of these steps in detail.

Rank selection. The nodes select unique ranks as follows. First, a leader is elected by using a leader election algorithm. Let s be the leader node. The leader picks a number $p(s)$ from the range $[b - 1, b]$, where b is a number arbitrarily chosen by s , and sends this number $p(s)$ along with its identity number $ID(s)$ to all of its neighbors. A neighbor v of the leader s , after receiving $p(s)$, picks a number $p(v)$ from the open interval $[p(s) - 1, p(s))$, thus $p(v)$ is less than $p(s)$, and then transmits $p(v)$ and $ID(v)$ to all of its neighbors. This process is repeated by every node in the graph. Notice that at some point, every node in the graph will receive a message from at least one of its neighbors since the given graph is connected; some nodes may receive more than one message. As soon as a node u receives the first message from a neighbor v , it picks a number $p(u)$ from $[p(v) - 1, p(v))$, so that it is smaller than $p(v)$, and transmit $p(u)$ and $ID(u)$ to the neighbors. If u receives another message later from another neighbor v' , u simply stores $p(v')$ and $ID(v')$, and does nothing else. $p(u)$ and $ID(u)$ constitute u 's rank $r(u)$ as follows.

Definition 5 *Rank.* The rank of a node u is defined as $r(u) = (p(u), ID(u))$ and for any two nodes u and v ,

$$r(u) < r(v) \text{ iff}$$

$$p(u) < p(v) \quad \text{or} \quad [p(u) = p(v) \text{ and } ID(u) < ID(v)].$$

At the end of execution of the above procedure of rank selection, it is easy to make the following observations.

Observation 1 *Each node knows the ranks of all of its neighbors.*

Proof Once a node receives the rank from one of its neighbors, it selects its own rank and sends it to all of its neighbors. Since the underlying graph G is connected, eventually (within time D , where D is the diameter of the graph), each node receives the messages containing the ranks from all of its neighbors. \square

Observation 2 *Each node u , except the leader s , has at least one neighbor v , i.e., $(u, v) \in E$, such that $r(u) < r(v)$.*

Proof Each node $u \neq s$, selects its own rank $r(u)$ such that $r(u) < r(v)$ only after receiving $r(v)$ from some neighbor v . \square

Observation 3 The leader s has the highest rank among all nodes in the graph.

Proof Since the leader s is the initiator of this rank selection process, we have $r(s) > r(v)$ for any $v \in V$ where $v \neq s$. \square

Connecting to a higher-ranked node. Each node v (except the leader s) executes the following algorithm simultaneously to find the nearest node of higher rank and connect to it. By Observation 2, we can conclude that for any node v , exploring the nodes in $\Gamma_{W(v)}(v)$ is sufficient to find a node of higher rank.

Each node v executes the algorithm in *phases*. In the first phase, v sets $\rho = 1$. In the subsequent phases, it doubles the value of ρ ; that is, in the i th phase, $\rho = 2^{i-1}$. In a phase of the algorithm, v explores the nodes in $\Gamma_\rho(v)$ to find a node u (if any) such that $r(u) > r(v)$. If such a node with higher rank is not found, v continues to the next phase with ρ doubled. By Observation 2, v needs to increase ρ to at most $W(v)$. Each phase of the algorithm consists of one or more *rounds*. In the first round, v sets $\lambda = 1$. In the subsequent rounds, the values for λ are doubled, i.e., in the j th round, $\lambda = 2^{j-1}$. In a particular round, v explores all nodes in $\Gamma_{\rho,\lambda}(v)$. At the end of each round, v counts the number of the nodes it has explored. If the number of nodes remain the same in two successive rounds of the same phase (that is, v already explored all nodes in $\Gamma_\rho(v)$), v doubles ρ and starts the next phase. If at any point of time, v finds a node of higher rank, it terminates its exploration.

Since all of the nodes explore their neighborhoods simultaneously, many nodes may have overlapping ρ -neighborhoods. This might create congestion of the messages in some edges that may result in increased running time of the algorithm, in some cases by a factor of $\Theta(n)$. Consider the network given in Fig. 1. If $r(v) < r(u_i)$ for all i , when $\rho \geq 2$ and $\lambda \geq 2$, an exploration message sent to v by any u_i will be forwarded to all other u_i s. Note that the values for ρ and λ for all u_i s may not necessarily be the same at a particular time. Thus, the congestion at any edge (v, u_i) can be as much as the number of such nodes u_i , which can be, in fact, $\Theta(n)$ in some graphs. However, to improve the running time of the algorithm, we keep congestions on all edges bounded by $O(1)$ by sacrificing the quality of the NNT, but only by a constant factor. To do so, v decides that some lower ranked u_i s can connect to some higher ranked u_i s and informs them instead of forwarding their message to the other nodes (details are given below). Thus, v forwards messages from only one u_i and this avoids the congestion. As a result, a node may not connect to the nearest node of higher rank. However,

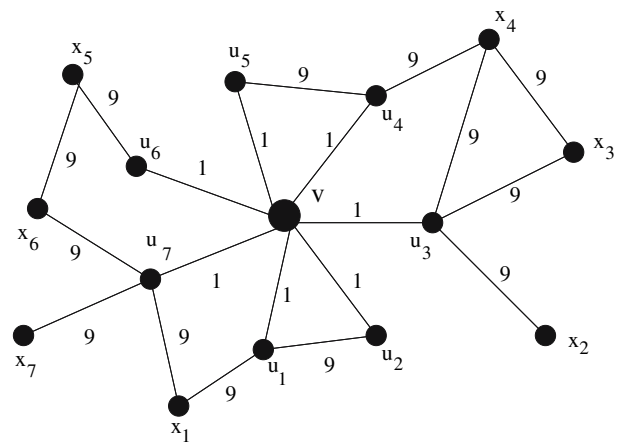


Fig. 1 A network with possible congestion in the edges adjacent to v . The weight of the edge (v, u_i) is one for every i , and nine for the rest of the edges. Assume $r(v) < r(u_i)$ for all i

our algorithm guarantees that the distance to the connecting node is not larger than four times the distance to the nearest node of higher rank. The detailed description is given below.

1. Exploration of ρ -neighborhood to find a node of higher rank:

Initiating exploration. Initially, each node v sets the radius $\rho \leftarrow 1$ and $\lambda \leftarrow 1$. The node v explores the nodes in $\Gamma_{\rho,\lambda}(v)$ in a BFS-like manner to find if there is a node $x \in \Gamma_{\rho,\lambda}(v)$ such that $r(v) < r(x)$. v sends *explore* messages $\langle \text{explore}, v, r(v), \rho, \lambda, pd, l \rangle$ to all $u \in N_\rho(v)$. In the message $\langle \text{explore}, v, r(v), \rho, \lambda, pd, l \rangle$, v is the originator of the *explore* message; $r(v)$ is its rank, ρ is its current phase value; λ is its current round number in this phase; pd is the weight of the path traveled by this message so far (from v to the current node), and l is the number of links that the message can travel further. Before v sends the message to its neighbor u , v sets $pd \leftarrow w(v, u)$ and $l \leftarrow \lambda - 1$.

Forwarding explore messages. Any node y may receive more than one *explore* message from the same originator v via different paths for the same round. Any subsequent message is forwarded only if the later message arrives through a shorter path than the previous one. Any node y , after receiving the message $\langle \text{explore}, v, r(v), \rho, \lambda, pd, l \rangle$ from one of its neighbors, say z , checks if it previously received another message $\langle \text{explore}, v, r(v), \rho, \lambda, pd', l' \rangle$ from z' with the same originator v such that $pd' \leq pd$. If so, y sends back a *count* message to z with $\text{count} = 0$. The purpose of the *count* messages is to determine the number of nodes explored by v in this round. Otherwise, if $r(v) < r(y)$, y sends back a *found* message to v containing y 's rank. Otherwise, If $N_{\rho-pd}(y) - \{z\} = \emptyset$ or $l = 0$, y sends back a *count* message with $\text{count} = 1$ and sets a marker $\text{counted}(v, \rho, \lambda) \leftarrow \text{TRUE}$. The purpose of the marker $\text{counted}(v, \rho, \lambda)$ is to make sure that y is counted only once for the same source

v and in the same phase and round of the algorithm. If $r(v) > r(y)$, $l > 0$, and $N_{\rho-pd}(y) - \{z\} \neq \emptyset$, y forwards the *explore* message to all of its neighbors $u \in N_{\rho-pd}(y) - \{z\}$ after setting $pd \leftarrow pd + w(y, u)$ and $l \leftarrow l - 1$.

Controlling congestion. If at any time step, a node v receives more than one, say $k < 1$, *explore* messages from different originators u_i , $1 \leq i \leq k$, v forwards only one *explore* message and replies back to the other u_i s as follows. Let $\langle \text{explore}, u_i, r(u_i), \rho_i, \lambda_i, pd_i, l_i \rangle$ be the *explore* message from originator u_i . If there is a u_j such that $r(u_i) < r(u_j)$ and $pd_j \leq \rho_i$, v sends back a *found* message to u_i telling that u_i can connect to u_j where the weight of the connecting path $w(Q(u_i, u_j)) = pd_i + pd_j \leq 2\rho_i$. In this way, some of the u_i s are replied back a *found* message and their *explore* messages will not be forwarded by v .

Now, there are at least one u_i left, to which v did not send the *found* message back. If there is exactly one such u_i , v forwards its *explore* message; otherwise, v takes the following actions. Let u_s be the node with lowest rank among the rest of the u_i s (i.e., those u_i s which were not sent a *found* message by v), and u_t , with $t \neq s$, be an arbitrary node among the rest of u_i s. Now, it must be the case that ρ_s is strictly smaller than ρ_t (otherwise, v would send a *found* message to u_s), i.e., u_s is in an earlier phase than u_t . This can happen if in some previous phase, u_t exhausted its ρ -value with smaller λ -value leading to a smaller number of rounds in that phase and a quick transition to the next phase. In such a case, we keep u_t waiting for at least one round without affecting the overall running time of the algorithm. To do this, v forwards *explore* message of u_s only and sends back *wait* messages to all u_t .

Each *explore* message triggers exactly one reply (either *found*, *wait*, or *count* message). These reply back messages move in similar fashion as of *explore* messages but in the reverse direction and they are aggregated (convergecast) on the way back as described next. Thus those reply messages also do not create any congestion in any edge.

Convergecast of the replies of the *explore* messages. If any node y forwards the *explore* message $\langle \text{explore}, v, r(v), \rho, \lambda, pd, l \rangle$ received from z for the originator v to its neighbors in $N_{\rho-pd}(y) - \{z\}$, eventually, at some point later, y will receive replies to these *explore* messages, from the nodes in $N_{\rho-pd}(y) - \{z\}$. Each of these replies is either a *count* message, a *wait* message, or a *found* message. Once y receives replies from all nodes in $N_{\rho-pd}(y) - \{z\}$, it takes the following actions. If at least one of the replies is a *found* message, y ignores all *wait* and *count* messages, and sends the *found* message to z toward the originator v . If there is more than one *found* message, select the one with the minimum path weight and ignore the rest. Now, if there is no *found* message and at least one *wait* message, y sends back only one *wait* message to z toward the originator v and ignore the

count messages. If all of the replies are *count* messages, y adds the count values of these messages and sends a single *count* message to v with the aggregated count. Also, y adds itself to the count if the marker $\text{counted}(v, \rho, \lambda) = \text{FALSE}$ and sets $\text{counted}(v, \rho, \lambda) \leftarrow \text{TRUE}$. At the very beginning, y initializes $\text{counted}(v, \rho, \lambda) \leftarrow \text{FALSE}$. The *count* messages (also the *wait* and *found* messages) travel in the opposite direction of the *explore* messages using the same paths toward v . Thus, these reply back messages form a convergecast as opposed to the (controlled) broadcast of the *explore* messages.

Actions of the originator after receiving the replies of the *explore* messages. At some time step, v receives replies of the *explore* messages originated by itself from all nodes in $N_\rho(v)$. Each of these replies is either a *count* message, a *wait* message, or a *found* message. If at least one of the replies is a *found* message, v is done with the exploration and makes the connection as described in Item 2 below. Otherwise, if there is a *wait* message, v again initiates exploration with the same ρ and λ . If all of them are *count* messages, v calculates the total count by adding the count values of these messages and does the following:

- if $\lambda = 1$, v initiates exploration with $\lambda \leftarrow 2$ and the same ρ (2nd round of the same phase);
- if $\lambda > 1$ and the total count for this round is larger than that of the previous round, v initiates exploration with $\lambda \leftarrow 2\lambda$ and the same ρ (next round of the same phase);
- otherwise, v initiates exploration with $\lambda \leftarrow 1$ and $\rho \leftarrow 2\rho$ (first round of the next phase).

2. Making connection:

Let u be a node with higher rank that v found by exploration. If v finds more than one node with rank higher than the rank of itself, then v selects the nearest one among them (break the ties arbitrarily). Let $Q(v, u)$ be the path from v to u . The path $Q(v, u)$ is discovered when u is found in the exploration process initiated by v . During the exploration process, the intermediate nodes in the path simply keep track of the predecessor and successor nodes in the path $Q(v, u)$ for this originator v . The edges in $Q(v, u)$ are added in the resulting spanning tree as follows. To add the edges in $Q(v, u)$, v sends a *connect* message to u along this path. Let $Q(v, u) = \langle v, \dots, x, y, \dots, u \rangle$. Note that by our choice of u , all of the intermediate nodes in this path have rank lower than $r(v)$. When the *connect* message passes through the edge (x, y) , node x uses (x, y) as its connecting edge regardless of the ranks of x and y . If x is still doing exploration to find a higher ranked node, x stops the exploration process as the edge (x, y) serves as x 's connection. If x is already connected using a path, say $\langle x, x_1, x_2, \dots, x_k \rangle$, the edge (x, x_1) is removed from the tree, but the rest of the

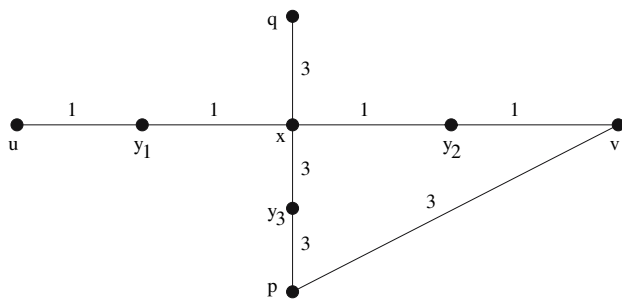


Fig. 2 A possible scenario of creating cycle and avoiding it. Nodes are marked with letters. Edge weights are given in the figure. Let $r(u) = 11$, $r(v) = 12$, $r(p) = 13$, $r(q) = 14$, and ranks of the rest of the nodes are smaller than 11. Node u connects to v , v connects to p , and p connects to q

edges in this path still remains in the tree. Once u receives the *connect* message originated by v , u sends a *rank-update* message back to v . All nodes in the path $Q(v, u)$ including v upgrade their ranks to $r(u)$; i.e., they assume a new rank which is equal to the rank of u .

It might happen that in between exploration and connection, some node x in the path $Q(v, u)$ changed its rank due to a connection by some originator other than v . In such a case, when the *connect* message originated by v travels through x , if x 's current rank is larger than $r(v)$, x accepts the connection as the last node in the path and returns a *rank-update* message with $r(x)$ toward v instead of forwarding the *connect* message to the next node (i.e., y) toward u . This is necessary to avoid cycle creation.

Each node has a unique rank and it can connect only to a node with higher rank. Thus if each node can connect to a node of higher rank using a direct edge (as in a complete graph), it is easy to see that there cannot be any cycle. However, in the above algorithm, a node u connects to a node of higher rank, v , $r(u) < r(v)$, using a path $Q(u, v)$, which may contain more than one edge and in such a path, ranks of the intermediate nodes are smaller than $r(u)$. Thus the only possibility of creating a cycle is when some other connecting path goes through these intermediate nodes. For example, in Fig. 2, the paths $P(u, v)$ and $P(p, q)$ both go through a lower ranked node x .

In Fig. 2, if p connects to q using path $\langle p, x, q \rangle$ before u makes its connection, x gets a new rank which is equal to $r(q)$. Thus u finds a higher ranked node, x , at a closer distance than v and connects to x instead of v . Note that if x is already connected to some node, it releases such connection and takes $\langle x, q \rangle$ as its new connection, i.e., q is x 's new parent. Now y_2 uses either (y_2, x) or (y_2, v) , but not both, for its connection. Thus there is no cycle in the resulting graph.

Now, assume that u already made its connection to v , but p is not connected yet. At this moment, x 's rank is upgraded to $r(v)$ which is still smaller than $r(p)$. Thus p finds q

as its nearest node of higher rank and connects using path $\langle p, x, q \rangle$. In this connection process, x removes its old connecting edge (x, y_2) and gets (x, q) as its new connecting edge. Again, there cannot be any cycle in the resulting graph.

If x receives the connection request messages from both u (toward v) and p (toward q) at the same time, x only forwards the message for the destination with highest rank; here it is q . u 's connection only goes up to x . Note that x already knows the ranks of both q and v from previous exploration steps. In the next section, a formal and robust proof is given to show that there is no cycle in the resulting NNT (Lemma 4).

2.4 Analysis of algorithm

In this section, we analyze the correctness and performance of the distributed NNT algorithm. The following lemmas and theorems show our results.

Lemma 1 Assume that during the exploration, v found a higher ranked node u and the path $Q(v, u)$. If v 's nearest node of higher rank is u' , then $w(Q) \leq 4d(v, u')$.

Proof Assume that u is found when v explored the (ρ, λ) -neighborhood for some ρ and λ . Then $d(v, u') > \rho/2$, otherwise, v would find u' as a node of higher rank in the previous phase and would not explore the ρ -neighborhood. Now, u could be found by v in two ways. i) The *explore* message originated by v reaches u and u sends back a *found* message. In this case, $w(Q) \leq \rho$. ii) Some node y receives two *explore* messages originated by v and u via the paths $R(v, y)$ and $S(u, y)$, respectively, where $r(v) < r(u)$ and $w(S) \leq \rho$; and y (on behalf of u) sent a *found* message to v (see "Controlling Congestion" in Item 1). In this case, $w(Q) = w(R) + w(S) \leq 2\rho$, since $w(R) \leq \rho$. Thus, in both cases, we have $w(Q) \leq 4d(v, u')$. \square

Lemma 2 The algorithm adds exactly $n - 1$ edges to the NNT.

Proof Let a node v connect to another node u using the path $Q(v, u) = \langle v, \dots, x, y, z, \dots, u \rangle$. When a *connect* message goes through an edge, say (x, y) (from x to y), in this path, the edge (x, y) is added to the tree. We say the edge (x, y) is associated to node x (not to y) based on the direction of the flow of the *connect* message. If, previously, x was associated to some other edge, say (x, y') , the edge (x, y') was removed from the tree. Thus each node is associated to at most one edge.

Except the leader s , each node x must make a connection and thus at least one *connect* message must go through or from x . Then, each node, except s , is associated to some edge in the tree.

Thus each node, except s , is associated to exactly one edge in the NNT; and s cannot be associated to any node since a

connect message cannot be originated by or go through s ; s can only be the destination (the last node in the path) since s has the highest rank.

Now, to complete the proof, we need to show that no two nodes are associated to the same edge. To show this, we use the following lemma.

Lemma 3 *Whenever x is associated to the edge (x, y) , at that point of time, $r(x) \leq r(y)$.*

Proof Node x become associated to the edge (x, y) only after a *connect* message passes through (x, y) from x to y . When the *connect* message went through (x, y) from x to y , $r(x)$ and $r(y)$ became equal. Later if another *connect* message increases $r(x)$, then either $r(y)$ is also increased to the same value or x become associated to some edge other than (x, y) . Thus, while keeping (x, y) associated to x , it must be true that $r(x) \leq r(y)$. \square

Only the nodes x and y can be associated to the edge (x, y) . Let x be associated to the edge (x, y) . By Lemma 3, $r(x) \leq r(y)$. Then any new *connect* message that might make (x, y) associated to y , by passing the *connect* message from y to x , must pass through x toward some node with rank higher than $r(y)$ (i.e., this connect message cannot terminate at x). This will make x associated to some edge other than (x, y) . Therefore, no two nodes are associated to the same edge. \square

Lemma 4 *The edges in the NNT added by the given distributed algorithm do not create any cycle.*

Proof Suppose to the contrary that

$\langle v_0, v_1, v_3, \dots, v_k, v_0 \rangle$ be a cycle created by the edges added to the NNT. Then either one of the following must be true.

- v_i is associated to (v_i, v_{i+1}) for $0 \leq i \leq k-1$, and v_k is associated to (v_k, v_0) .
- v_i is associated to (v_i, v_{i-1}) for $1 \leq i \leq k$, and v_0 is associated to (v_0, v_k) .

For both cases, by Lemma 3, we have $r(v_0) = r(v_1) \leq \dots = r(v_k)$. Here, we have a contradiction. The ranks of all nodes in this cycle can never be the same. Initially, the ranks are distinct. Later, when a node u connects to the node v via the *connecting path* $Q(u, v)$, the ranks of the nodes in the path $Q(u, v)$ are upgraded to $r(v)$. Notice that the rank of a node cannot be decreased. It can only be increased. It is easy to see that a single connecting path cannot contain any cycle. The above cycle must be created by at least two connecting paths. Let $Q(u, v)$ be the last connecting path that completes this cycle, and v_i and v_j be the first and last node in the path $Q(u, v)$ among the nodes that are common both in this

path and the cycle. The path $Q(u, v)$ goes beyond v_i ; that means $r(v) > r(v_i)$. Since $r(v_i) = r(v_j)$, $r(v) > r(v_j)$; this implies that $v \neq v_j$ and the path $Q(u, v)$ goes beyond v_j . As a result, this connecting path (u, v) will upgrade the ranks of v_i and v_j to $r(v)$, which is higher than the ranks of the other nodes in the cycle. This leads to a contradiction. Thus, there cannot be any cycle in the NNT. \square

From Lemmas 2 and 4 we have the following theorem.

Theorem 1 *The above algorithm produces a tree spanning all nodes in the graph.*

We next show that the spanning tree found by our algorithm is an $O(\log n)$ -approximation to the MST (Theorem 2).

Theorem 2 *Let the NNT be the spanning tree produced by the above algorithm. Then the cost of the tree $c(NNT) \leq 4\lceil \log n \rceil c(MST)$.*

Proof Let $H = (V_H, E_H)$ be a complete graph constructed from $G = (V, E)$ as follows. $V_H = V$ and weight of the edge $(u, v) \in E_H$ is the weight of the shortest path $P(u, v)$ in G . Now, the weights of the edges in H satisfy the triangle inequality. Let NNT_H be a nearest neighbor tree and MST_H be a minimum spanning tree on H . We can show that $c(NNT_H) \leq \lceil \log n \rceil c(MST_H)$ (a proof is given in the appendix).

Let NNT' be a spanning tree on G , where each node connects to the nearest node of higher rank via a shortest path. By Lemma 1, we have $c(NNT) \leq 4c(NNT')$. Further, it is easy to show that $c(NNT') \leq c(NNT_H)$ and $c(MST_H) \leq c(MST)$. Thus, we have

$$\begin{aligned} c(NNT) &\leq 4c(NNT_H) \\ &\leq 4\lceil \log n \rceil c(MST_H) \\ &\leq 4\lceil \log n \rceil c(MST). \end{aligned}$$

\square

Remark With the help of Theorem 2.13 in [10], an alternative upper bound of $12\lceil \log \frac{w_{max}}{w_{min}} \rceil c(MST)$ for $c(NNT)$ can be achieved, where w_{max} and w_{min} are the maximum and minimum edge weights, respectively. This bound is independent of the number of nodes n , but depends on the weights of the edges.

Theorem 3 *The running time of the above algorithm is*

$$O(D + L \log n).$$

Proof Time to elect leader is $O(D)$. The rank choosing scheme takes also $O(D)$ time.

In the exploration process, ρ can increase to at most $2W$; because, within distance W , it is guaranteed that there is a node of higher rank (Observation 2). Thus, the number of phases in the algorithm is at most $O(\log W) = O(\log n)$.

In each phase, λ can grow to at most $4L$. When $L \leq \lambda < 2L$ and $2L \leq \lambda < 4L$, in both rounds, the count of the number of nodes explored will be the same. As a result, the node will move to the next phase.

Now, in each round, a node takes at most $O(\lambda)$ time; because the messages travel at most λ edges back and forth and at any time the congestion in any edge is $O(1)$. Thus any round takes time at most

$$\sum_{\lambda=1}^{\log(4L)} O(\lambda) = O(L).$$

Thus, time for the exploration process is $O(L \log W)$. Total time of the algorithm for leader election, rank selection, and exploration is $O(D + D + L \log n) = O(D + L \log n)$. \square

Theorem 4 *The message complexity of the algorithm is*

$$O(|E| \log L \log n) = O(|E| \log^2 n).$$

Proof The number of phases in the algorithm is at most $O(\log L)$. In each phase, each node executes at most $O(\log W) = O(\log n)$ rounds. In each round, each edge carries $O(1)$ messages. That is, number of messages in each round is $O(|E|)$. Thus total messages is $O(|E| \log L \log n)$. \square

3 Exact versus approximate MST and near-optimality of NNT algorithm

Comparison with distributed algorithms for (exact) MST.

There can be a large gap between the local shortest path diameter L and $\tilde{\Omega}(\sqrt{n})$, which is the lower bound for exact MST computation. In particular, we can show that there exists a family of graphs where NNT algorithm takes $\tilde{O}(1)$ time, but any distributed algorithm for computing (exact) MST will take $\tilde{\Omega}(\sqrt{n})$ time. To show this we consider the parameterized (weighted) family of graphs called \mathcal{J}_m^K defined in Peleg and Rabinovich [21] (see Sects. 4.1, 5.3 in [21] for a description of how to construct \mathcal{J}_m^K). (One can also show a similar result using the family of graphs defined by Elkin in [6].) The size of \mathcal{J}_m^K is $n = \Theta(m^{2K})$ and its diameter $\Theta(Km) = \Theta(Kn^{1/(2K)})$. For every $K \geq 2$, Peleg and Rabinovich show that any distributed algorithm for the MST problem will take $\Omega(\sqrt{n}/BK)$ time on some graphs belonging to the family. The graphs of this family have $L = \Theta(m^K) = \sqrt{n}$. We modify this construction as follows: the weights on all the highway edges except the first highway (H^1) is changed to 0.5 (originally they were all zero); all other weights remain the same. This makes $L = \Theta(Km)$, i.e., same order as the diameter. One can check that the proof of Peleg and Rabinovich is still valid, i.e., the lower bound for MST will take $\Omega(\sqrt{n}/BK)$ time on some graphs of this

family, but NNT algorithm will take only $\tilde{\Omega}(L)$ time. Thus we can state:

Theorem 5 *For every $K \geq 2$, there exists a family of n -vertex graphs in which NNT algorithm takes $O(Kn^{1/(2K)})$ time while any distributed algorithm for computing the exact MST requires $\tilde{\Omega}(\sqrt{n})$ time. In particular, for every $n \geq 2$, there exists a family of graphs in which NNT algorithm takes $\tilde{O}(1)$ time whereas any distributed MST algorithm will take $\tilde{\Omega}(\sqrt{n})$ time.*

Such a large gap between NNT and any distributed MST algorithm can be also shown for constant diameter graphs, using a similar modification of a lower bound construction given in Elkin [6] (which generalizes and improves the results of Lotker et al. [18]).

Near (existential) optimality of NNT algorithm. We show that there exists a family of graphs such that any distributed algorithm to find a $H(\leq \log n)$ -approximate MST takes $\Omega(L)$ time (where L is the local shortest path diameter) on some of these graphs. Since NNT algorithm takes $\tilde{O}(D+L)$, this shows the near-tight optimality of NNT (i.e., tight up to a $\text{polylog}(n)$ factor). This type of optimality is called *existential optimality* which shows that our algorithm cannot be improved in general. To show our lower bound we look closely at the hardness of distributed approximation of MST shown by Elkin [6]. Elkin constructed a family of weighted graphs \mathcal{G}^ω (Fig. 1, Sect. 3.1 in [6]) to show a lower bound on the time complexity of any H -approximation distributed MST algorithm (whether deterministic or randomized). We briefly describe this result and show that this lower bound is *precisely the local shortest path diameter L of the graph*. The graph family $\mathcal{G}^\omega(\tau, m, p)$ is parameterized by 3 integers τ, m , and p , where $p \leq \log n$. The size of the graph $n = \Theta(\tau m)$, the diameter is $D = \Theta(p)$ and the local shortest path diameter can be easily checked to be $L = \Theta(m)$. Note that graphs of different size, diameter, and $LSPD$ can be obtained by varying the parameters τ, m , and p . (We refer to [6] for the detailed description of the graph family and the assignment of weights.) We now slightly restate the results of [6] (assuming the CONGEST(B) model):

Theorem 6 [6]

1. *There exist some graphs belonging to the family $\mathcal{G}^\omega(\tau, m, p)$ having diameter at most D for $D \in 4, 6, 8, \dots$ and $LSPD L = \Theta(m)$ such that any randomized H -approximation algorithm for the MST problem on these graphs takes $T = \Theta(L) = \Omega((\frac{n}{H \cdot D \cdot B})^{\frac{1}{2} - \frac{1}{2(D-1)}})$ distributed time.*
2. *If $D = O(\log n)$ then the lower bound can be strengthened to $\Theta(L) = \Omega(\sqrt{\frac{n}{H \cdot B \cdot \log n}})$.*

Using a slightly different weighted family $\tilde{\mathcal{G}}^\omega(\tau, m)$ parameterized by two parameters τ and m , where size $n = \tau m^2$, diameter $D = \Omega(m)$ and LSPD $L = \Theta(m^2)$, one can strengthen the lower bound of the above theorem by a factor of $\sqrt{\log n}$ for graphs of diameter $\Omega(n^\delta)$.

The above results show the following two important facts:

1. There are graphs having diameter $D \ll L$ where any H -approximation algorithm requires $\Omega(L)$ time.
2. More importantly, for graphs with very different diameters—varying from a constant (including 1, i.e., exact MST) to logarithmic to polynomial in the size of n —the lower bound of distributed approximate-MST is captured by the local shortest path parameter. In conjunction with our upper bound given by the NNT algorithm which takes $\tilde{O}(D + L)$ time, this implies that the LPSD L captures in a better fashion the complexity of distributed $O(\log n)$ -approximate-MST computation.

4 Special classes of graphs

We show that in unit disk graphs (a commonly used model for wireless networks) $L = 1$, and in random weighted graphs, $L = O((\log n))$ with high probability. Thus our algorithm will run in near-optimal time of $\tilde{O}(D(G))$ on these graphs.

Unit disk graph (UDG). Unit disk graph is an euclidian graph where there is an edge between two nodes u and v if and only if $\text{dist}(u, v) \leq R$ for some R (R is typically taken to be 1). Here $\text{dist}(u, v)$ is the euclidian distance between u and v ; that is the weight of the edge (u, v) . Theorem 7 shows that for any UDG, $L = 1$. For a 2-dimensional UDG, the diameter D can be as large as $\Theta(\sqrt{n})$.

Theorem 7 *In any UDG, the local shortest path diameter L is 1.*

Proof For any node v , $W(v) \leq R$ by definition of UDG. Now if there is a node u such that $d(u, v) \leq R$, then $\text{dist}(u, v) \leq R$ by the triangle inequality. Thus, (v, u) is in E and the edge (v, u) is the shortest path from v to u ; as a result, $l(v, u) = 1$. Therefore, for any v ,

$$L(v) = \max_{u \in \Gamma_{W(v)}(v)} l(v, u) = 1,$$

$$L = \max_{v \in V} L(v) = 1.$$

□

Graph with random edge weights. Consider any graph G (topology can be arbitrary) with edge weights chosen randomly from $[0, 1]$ following any arbitrary distribution (i.e., each edge weight is chosen i.i.d from the distribution). The following theorem shows that L and S is small compared to the diameter for such a graph.

Theorem 8 *Consider a graph G where the edge weights are chosen randomly from $[0, 1]$ following any (arbitrary) distribution with a constant (independent of n) mean. Then: (1) $L = O(\log n)$ with high probability (whp), i.e., probability at least $1 - 1/n^{\Omega(1)}$; and (2) the shortest path diameter $S = O(\log n)$ if $D < \log n$ and $S = O(D)$ if $D \geq \log n$ whp.*

Proof Let the edge weights are randomly drawn from $[0, 1]$ with mean μ . For any node v , $W(v) \leq 1$. Consider any path with $m = k \log n$ edges, for some constant k . Let the weights of the edges in this path be w_1, w_2, \dots, w_m . For any i , $E[w_i] = \mu$. Since $\frac{1}{2}\mu k \log n \geq 1$ for sufficiently large k , we have

$$\begin{aligned} \Pr \left\{ \sum_{i=1}^m w_i \leq 1 \right\} &\leq \Pr \left\{ \sum_{i=1}^m w_i \leq \frac{1}{2} \mu k \log n \right\} \\ &= \Pr \left\{ \mu - \frac{1}{m} \sum_{i=1}^m w_i \geq \frac{1}{2} \mu \right\}. \end{aligned}$$

Using Hoeffding bound [11] and putting $k = \frac{6}{\mu^2}$,

$$\Pr \left\{ \mu - \frac{1}{m} \sum_{i=1}^m w_i \geq \frac{1}{2} \mu \right\} \leq e^{-m\mu^2/2} = \frac{1}{n^3}.$$

Thus if it is given that the weight of a path is at most 1, then the probability that the number of edges $\leq \frac{6}{\mu^2} \log n$ is at most $\frac{1}{n^3}$. Now consider all nodes u such that $d(v, u) \leq W(v)$. There are at most $n - 1$ such nodes and thus there are at most $n - 1$ shortest paths leading to those nodes from v . Thus using union bound,

$$\Pr \left\{ L(v) \geq \frac{6}{\mu^2} \log n \right\} \leq n \times \frac{1}{n^3} = \frac{1}{n^2}.$$

Using $L = \max\{L(v)\}$ and union bound,

$$\Pr \left\{ L \geq \frac{6}{\mu^2} \log n \right\} \leq n \times \frac{1}{n^2} = \frac{1}{n}.$$

Therefore, with probability at least $1 - \frac{1}{n}$, L is smaller than or equal to $\frac{6}{\mu^2} \log n$. Proof of part 2 is similar. □

5 Conclusion and future work

We presented and analyzed a simple approximation algorithm for constructing a low-weight spanning tree. We also presented its efficient implementation in an arbitrary network of processors.

The local nature of the NNT-scheme seems to be suitable for designing an efficient distributed *dynamic* algorithm, where the goal is to maintain an NNT of good quality, as nodes are added or deleted. Moreover, it is interesting to see whether the ideas in this paper can be extended to design

an efficient distributed algorithm for the more challenging problem of finding a k -connected subgraph. These look promising for future work.

Acknowledgments We are very grateful to the referees for their careful reading of the paper and detailed comments which helped greatly in improving the presentation of the paper.

Appendix

Cost of an NNT in a complete metric graph

For a complete metric graph H , Lemma 6 shows an upper bound on the cost of the nearest neighbor tree NNT_H with respect to the cost of a minimum spanning tree MST_H on H . To prove Lemma 6, we use the following lemma concerning the traveling salesman problem (TSP) that appears in Rosenkrantz, Stearns, and Lewis [22, Lemma 1].

Lemma 5 [22] *Let $G = (V, E)$ be a weighted metric (complete) graph on n nodes. Let $d(p, q)$ be the weight of the edge between nodes p and q . Suppose there is a mapping l assigning each node p a number l_p such that the following two conditions hold:*

- (a) $d(p, q) \geq \min(l_p, l_q)$ for all nodes p and q .
- (b) $l_p \leq \frac{1}{2}c(TSP)$ for all nodes p , where $c(TSP)$ is the cost of a optimal (shortest) traveling salesman tour in G .

Then

$$\sum_{p \in V} l_p \leq \frac{1}{2}([\log n] + 1)c(TSP).$$

Lemma 6 *For a complete metric graph H ,*

$$c(NNT_H) \leq [\log n]c(MST_H).$$

Proof It can be shown that (e.g., see [3])

$$c(TSP_H) \leq 2c(MST_H). \quad (1)$$

To apply Lemma 5, based on the NNT scheme, we define a mapping l as follows:

$l_p = d(p, nnt(p))$ if $nnt(p)$ (the node that p connects to) exists; otherwise (if p is the highest-ranked node), $l_p = \frac{1}{2}c(TSP_H)$.

l satisfies condition (a): Let p and q be any two nodes, and without loss of generality, assume $rank(p) < rank(q)$. Then by definition of $nnt()$, $d(p, q) \geq d(p, nnt(p)) = l_p$ (note that p cannot be the highest-ranked node).

l satisfies also condition (b): It is trivially true for the highest-ranked node. For any other node p , $l_p = d(p, nnt$

$(p))$. There are exactly two disjoint paths between p and $nnt(p)$ in the TSP route. Let S_1 and S_2 be these two paths. Then

$$c(S_1) + c(S_2) = c(TSP_H),$$

and by triangle inequality, we have $d(p, nnt(p)) \leq c(S_1)$ and also $d(p, nnt(p)) \leq c(S_2)$. Thus

$$d(p, nnt(p)) \leq \frac{1}{2}c(TSP_H).$$

Let p_0 be the highest-ranked node. By the construction of NNT and applying Lemma 5, we have

$$c(NNT_H) = \sum_{p \in V} l_p - l_{p_0} \leq \frac{1}{2}[\log n]c(TSP_H). \quad (2)$$

The lemma follows from Inequality 1 and 2. \square

Remark Lemma 6 can also alternatively be derived by a reduction of Theorem 2 in Imase and Waxman [12] (which also uses Lemma 1 of [22]). Imase and Waxman gave an $O(\log n)$ -approximation algorithm that maintains a Steiner tree dynamically under node additions only (not deletions). Theorem 2 in [12] can be used to show our Lemma 6 if we make the following observation: add the nodes one by one in decreasing order of their ranks.

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