Local Algorithms for Estimating Effective Resistance

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ABSTRACT
Effective resistance is an important metric that measures the similarity of two vertices in a graph. It has found applications in graph clustering, recommendation systems and network reliability, among others. In spite of the importance of the effective resistances, we still lack efficient algorithms to exactly compute or approximate them on massive graphs.

In this work, we design several local algorithms for estimating effective resistances, which are algorithms that only read a small portion of the input while still having provable performance guarantees. To illustrate, our main algorithm approximates the effective resistance between any vertex pair \( s, t \) with an arbitrarily small additive error \( \varepsilon \) in time \( O(\text{poly}(\log n/\varepsilon)) \), whenever the underlying graph has bounded mixing time. We perform an extensive empirical study on several benchmark datasets, validating the performance of our algorithms.

CCS CONCEPTS
• Mathematics of computing → Graph algorithms; • Theory of computation → Sketching and sampling.

KEYWORDS
Graph algorithms, Random walks, Effective resistances

1 INTRODUCTION
Metrics that capture the similarity between vertices in a graph have played a pivotal role in the quest for understanding the structure of large-scale networks. Typical examples include personalized PageRank (PPR) [32], Katz similarity [20] and SimRank [19], each of which can be thought of as a random walk-based measure on graphs. These metrics have found applications in recommender systems [21], link prediction [27, 35], etc.

A remarkably important random walk-based metric for measuring vertex similarity is the effective resistance. Given a graph \( G \) treated as a resistor network, the effective resistance \( R_G(s, t) \) between two vertices \( s, t \) in \( G \) is the energy dissipation in the network when routing one unit of current from \( s \) to \( t \). It is well known that the effective resistance is inherently related to the behaviour of random walks on graphs\(^1\). Concretely, the effective resistance between \( s \) and \( t \) is proportional to the commute time \( \kappa(s, t) \), defined as the expected number of steps a random walk starting at \( s \) visits vertex \( t \) and then goes back to \( s \) [9]. Using this interpretation, we can deduce that the smaller \( R_G(s, t) \) is, the more similar two vertices \( s, t \) are.

Indeed, effective resistance has proven ubiquitous in numerous applications including graph clustering [2, 16], recommender systems [22], measuring robustness of networks [15], spectral sparsification [36], graph convolutional networks [1], location-based advertising [37], among others. Moreover, in the theoretical computer science community, the use of effective resistance has led to a breakthrough line of work for provably speeding up the running time of many flow-based problems in combinatorial optimization [3, 11, 29].

Despite the importance of effective resistance, we still lack efficient methods to compute or approximate them on massive graphs. For any two vertices \( s, t \) and approximation parameter \( \varepsilon > 0 \), one can \((1+\varepsilon)\)-approximate \( R_G(s, t) \) in \( \tilde{O}(m \log(1/\varepsilon)) \) time [13], where \( m \) denotes the number of edges in a graph. There exists an algorithm that \((1+\varepsilon)\)-approximates all-pairs effective resistances in \( \tilde{O}(n^2/\varepsilon) \) time [18]. These results, though theoretically competitive, require access to the entire input graph. Given the rapid growth of modern networks, such polynomial time algorithms (even those running in near linear time in the number of vertices and edges) are prohibitively costly. This motivates the following question:

Can we obtain a competitive estimation to \( R_G(s, t) \) while exploring only a small portion of the graph?

We address this question by exploiting the paradigm of local or sub-linear algorithms. This computational model is particularly desirable in applications where one requires the effective resistances amongst only a few number of vertex pairs. Despite that the effective resistance is a key tool in large-scale graph analytics, designing local algorithms for estimating it is a largely unexplored topic.

In this paper, we provide several local algorithms for estimating pairwise effective resistances with provable performance guarantees. For any specified vertex pair \( s, t \), our algorithms output an estimate of \( R_G(s, t) \) with an arbitrarily small constant additive error, while exploring a small portion of the graph. To formally state our results, we utilize the well-known adjacency list model [33], which assumes query access to the input graph \( G \) and supports the following types of queries in constant time:

\[^1\text{We only consider simple random walks in the paper: suppose we are at vertex } v, \text{ we jump to a neighbor of } v \text{ with probability } 1/\deg(v), \text{ where } \deg(v) \text{ is the degree of vertex } v.\]

\[^2\text{Throughout the paper, we use } \tilde{O} \text{ to hide polylogarithmic factors, i.e., } \tilde{O}(f(n)) = O(f(n) \cdot \text{poly log } f(n)).\]
• **degree query:** for any specified vertex \( v \), the algorithm can get the degree \( \deg(v) \) of \( v \);
• **neighbor query:** for any specified vertex \( v \) and index \( i \leq \deg(v) \), the algorithm can get the \( i \)-th neighbor of \( v \);
• **uniform sampling:** the algorithm can sample a vertex \( x \) of \( G \) uniformly at random.

Our main objective is to find a good estimate of the pairwise effective resistance \( R_G(s, t) \) for any specified vertex pair \( s, t \) by making as few queries as possible to the graph while achieving fast running time.

**Our contributions.** We give a systemic study of local algorithms for estimating \( s \)-\( t \) effective resistances for general graphs.

- Theoretically, we provide three types of local algorithms for estimating effective resistances. All of them are based on random walks, but vary from their connections to effective resistances: (i) the first type is based on approximating the pseudo inverse of the Laplacian matrix, (ii) the second type is based on commute times, (iii) the third type is based on the number of spanning trees.
- We empirically demonstrate the competitiveness of our algorithms on popular benchmarks for graph data. In particular, for certain real-world networks, we will see that our algorithms run \( 10^2 \) to \( 10^6 \) faster than existing polynomial-time methods and estimate effective resistance to within a multiplicative error of 0.1.

To illustrate, our main local algorithm approximates \( R_G(s, t) \) with an arbitrarily small additive error \( \epsilon \) in time \( O(\text{poly} \log n / \epsilon) \), whenever the underlying graph has bounded mixing time, which is justified in real-world networks. Previously, the only work on this problem was by Andoni et al. [4], and it achieves \((1 + \epsilon)\)-approximation to \( R_G(s, t) \) in \( O(\frac{1}{\epsilon^2} \text{poly} \log \frac{1}{\epsilon}) \) time for \( d \)-regular expander graphs. Indeed, one of our algorithms for general graphs is based on [4].

Using the fact that the length of shortest paths and effective resistances are exactly the same on tree graphs, we can observe that graphs with large mixing time do not admit efficient local algorithms. Concretely, let us consider a path graph on \( n \) vertices. It is known that the path graph has large mixing time, and there is no local algorithm that makes a sub-linear number of queries and approximates the length of shortest paths within a constant multiplicative factor or additive error, thus giving us the same impossibility result for effective resistances. This suggests that our bounded mixing time assumption is necessary to design local algorithms with sublinear number of queries and running time.

### 2 RELATED WORK

In this section, we discuss some related work.

Hayashi et al. [17] gave an algorithm for approximating the effective resistances of vertex pairs that are endpoints of edges. Their algorithm is based on sampling spanning trees uniformly at random, and it \((1 + \epsilon)\)-approximates \( R_G(s, t) \) for any \((s, t) \in E \) in expected running time \( \frac{\log(2m/\delta)}{2\epsilon^2} \cdot \sum_{u \in V} \pi_G(u) K_G(u, r) \), where \( \pi_G(u) \) denotes the stationary probability at a vertex \( u \in V \) of a random walk on \( G \), \( K_G(u, v) \) denotes the commute time between two vertices \( u, v \in V \) and \( r \in V \) is some vertex.

There also exist several local algorithms for other random walk based quantities, such as the stationary distribution, PageRank, Personalized PageRank and transition probabilities. The stationary distribution. Lee et al. [23] and Bressan et al. [8] studied the question of computing the stationary distribution \( \pi \) of a Markov Chain locally. These algorithms take as input any state \( v \), and answer if the stationary probability of \( v \) exceeds some \( \Delta \in (0, 1) \) and/or output an estimate of \( \pi(v) \). They only make use of a local neighborhood of \( v \) on the graph induced by the Markov chain and run in sublinear time for some families of Markov Chains.

PageRank. Borgs et al. presented a method for identifying all vertices whose PageRank is larger than some threshold [6]. Specifically, for a threshold value \( \Delta \geq 1 \) and a constant \( c > 3 \), with high probability, their algorithm returns a set \( S \subseteq V \) such that \( S \) contains all vertices with PageRank at least \( \Delta \) and no vertex with PageRank at least \( \Delta/c \). The algorithm runs in \( O(\frac{n}{\epsilon^2}) \) time.

Bressan et al. developed a sub-linear time algorithm that employs local graph exploration [7]. Their algorithm \((1 + \epsilon)\)-approximates the PageRank of a vertex on a directed graph. For constant \( \epsilon > 0 \), the algorithm runs in \( O(\min(m^{1/4} \cdot L/\epsilon, 1/\epsilon^3, m^{1/7} d^{-3/7})) \), where \( L \) and \( d \) are respectively the maximum and average outdegree.

Personalized PageRank (PPR). The PPR \( \pi_G(t) \) of a start vertex \( s \) and target vertex \( t \) measures the frequency of visiting \( t \) via short random-walks from \( s \). For a given threshold \( \delta \) such that \( \pi_G(t) > \delta \), Lofgren et al. solved this with small relative error and an expected running time of \( O(\sqrt{d/\delta}) \) [25], where \( d \) is the average in-degree of the graph. Their algorithm is based on a bi-directional search technique and an improved implementation was presented in [24].

Transition probabilities. Another problem related to effective resistance is estimating transition probabilities in a Markov chain. Specifically, given transition matrix \( P \), initial source distribution \( \sigma \), target state \( t \), and a fixed length \( t \), the goal is to estimate the probability \( p \) that an \( t \)-step random walk starting from distribution \( \sigma \) ends at \( t \). Banerjee and Lofgren developed an algorithm that can estimate such a probability with respect to a minimum threshold \( \delta \) such that \( p \geq \delta \) by employing a bi-directional approach [5]. Specifically, their algorithms returns an estimator \( \hat{p} \) of \( p \) such that with high probability \( |\hat{p} - p| < \max(\epsilon p, \delta) \) for any \( \epsilon > 0 \).

### 3 PRELIMINARIES

Let \( G = (V, E) \) be an undirected graph. For any \( v \in V \), we let \( \deg(v) \) denote the degree of \( v \). The volume of a set \( S \) of vertices, denoted \( \text{vol}(S) \), is the sum of their degrees. Furthermore, for a set \( S \subseteq V \), the conductance of \( S \), denoted \( \phi_G(S) \), is the number of edges with one endpoint in \( S \) and the other in \( V \setminus S \) divided by \( \text{vol}(S) \). The conductance of \( G \), denoted \( \phi(G) \), is defined to be \( \min_{S \subseteq V, 0 < \text{vol}(S) \leq \text{vol}(V)} \phi_G(S) \). A graph \( G \) is called an expander if \( \phi(G) \geq \phi \) for some universal constant \( \phi \in (0, 1) \).

Let \( A \) denote its adjacency matrix and let \( D \) denote the degree diagonal matrix. Let \( L = D - A \) denote the Laplacian matrix of \( G \). Let \( L^{1/2} \) denote the Moore-Penrose pseudo-inverse of the Laplacian of \( G \). Let \( 1_u \in \mathbb{R}^V \) denote the (row) indicator vector of vertex \( u \) such that \( 1_u(v) = 1 \) if \( v = u \) and 0 otherwise. Let \( \chi_{at} = 1_u - 1_t \).
Definition 3.1. Given any two vertices \( u, v \in V \), the \( s-t \) effective resistance is defined as
\[
R_G(s, t) := \chi_{s,t}L^1{\chi_{s,t}}^\top = \chi_{s,t}L^1 + 2L^1_{s,t} - 2L^1_{s,t} + L^1_{s,t}.
\]

Random walks. Given a graph \( G \), we consider the random walk on \( G \): suppose we are currently at \( v \), then we jump to a neighbor \( u \) with probability \( \frac{1}{\deg(u)} \). We use \( P := D^{-1}A \) to denote the random walk transition matrix. Let \( \lambda = \max(|\lambda_2|, |\lambda_n|) \), where \( \lambda_i \) is the \( i \)-th largest eigenvalue of the matrix \( P \).

Definition 3.2. The commute time \( \kappa(s, t) \) between vertices \( s, t \) is the expected number of steps in a random walk that starts at vertex \( s \) visits vertex \( t \) and then comes back to \( s \).

Random walks on graphs are a type of Markov Chain. A Markov chain is said to be positive recurrent if, starting in any state \( i \), the expected time until the process returns to state \( i \) is finite. A Markov chain is said to be aperiodic if for any state \( i \) there are no restrictions on when it is possible for the process to enter state \( i \).

Definition 3.3. A Markov chain is said to be ergodic if it is aperiodic and positive recurrent.

Informally, the mixing time of the graph \( G \) refers to the number of steps needed before a random walk on \( G \) converges to its stationary distribution. We refer to [34] for a formal definition. It is known that the spectral gap \( 1 - \lambda \) is intimately related to the mixing time of \( G \). That is, the larger \( 1 - \lambda \) is, the smaller mixing time is, and vice versa.

4 THE LOCAL ALGORITHMS

4.1 Algorithms based on approximating Laplacian inverse

In this section, we provide local algorithms for effective resistances by approximating the Laplacian pseudo-inverse \( L^1 \) of the graph.

High-level idea. Our algorithm works for general graphs and is based on the aforementioned sublinear-time algorithm for \( d \)-regular graphs [4]. The basic idea is as follows. Recall that by definition of effective resistance, \( R_G(s, t) = \chi_{s,t}L^1{\chi_{s,t}}^\top \) and \( P := D^{-1}A \) is the random walk transition matrix. Using the Neumann series of the matrix \( L^1 \) (see Lemma 4.3), we can write
\[
R_G(s, t) = \chi_{s,t} \sum_{k=0}^{\infty} P^kD^{-1}{\chi_{s,t}}^\top
= \chi_{s,t}P^{\ell-1}{\chi_{s,t}}^\top + \chi_{s,t} \sum_{k=\ell+1}^{\infty} P^kD^{-1}{\chi_{s,t}}^\top,
\]
for any \( \ell > 0 \). For graphs with large spectral gap (i.e., expanders graphs or graphs with low random walk mixing time), we can show that for any additive error \( \epsilon \), we can choose \( \ell \) appropriately such that the second term is at most \( \epsilon/2 \), and the first term can be approximated within additive error \( \epsilon/2 \). For the latter, we use a simple Monte Carlo approach (i.e., to use the empirical distribution of the endpoints of a small number of random walks) to approximate the quantity \( 1_{\epsilon}P^\ell \), the (transition) probability that a length-\( \ell \) random walk starting from \( s \) ends at \( t \), for any \( \ell \geq 1 \).

Now we introduce one assumption, building upon which we present and analyze two local algorithms.

Assumption 4.1. Let \( G \) be a connected graph with minimum vertex degree at least 1. Further, assume that the Markov chain corresponding to the random walk on \( G \) is ergodic.

The first algorithm: \texttt{EstEff-TranProb}. We first present an algorithm that uses the above idea. Recall that \( \lambda = \max(|\lambda_2|, |\lambda_n|) \), where \( \lambda_i \) is the \( i \)-th largest eigenvalue of the matrix \( P \).

Theorem 4.2. Under Assumption 4.1, there is an algorithm \texttt{EstEff-TranProb}(\( G, s, t \)) (see Algorithm 1) that outputs an estimate \( \hat{\delta}_{s,t} \) such that with probability at least 9/10, it holds that
\[
|\hat{\delta}_{s,t} - \delta_{s,t}| \leq \epsilon.
\]

The running time and query complexity of the algorithm are \( O(t^4(\log t)/\epsilon^2) \) for \( t = \frac{\log(4(1 - \epsilon)\lambda)}{\log(1/\delta)} \).

The above algorithm is very efficient, if the graph has small \( \lambda \), or has low mixing time, a property that is satisfied by many real networks. Now we present the algorithm \texttt{EstEff-TranProb}.

Algorithm 1: \texttt{EstEff-TranProb}(\( G, s, t \))

1. \[ t = \frac{\log(4(1 - \epsilon)\lambda)}{\log(1/\delta)} \]
2. \[ r \leftarrow \frac{40t^2(\log(80t))}{\epsilon^2} \]
3. for \( i = 0, 1, \ldots, t - 1 \) do
4. \[ \text{Perform } r \text{ independent random walks of length } i \]
5. \[ \text{starting at } s, \text{ and let } X_{i,s} \text{ (resp., } X_{i,t} \text{)} \]
6. \[ \text{be the number of walks that end at } s \text{ (resp., } t \text{).} \]
7. \[ \text{Perform } r \text{ independent random walks of length } i \]
8. \[ \text{starting at } t, \text{ and let } Y_{i,s} \text{ (resp., } Y_{i,t} \text{)} \]
9. \[ \text{be the number of walks that end at } s \text{ (resp., } t \text{).} \]
10. \[ \text{Set } \hat{\delta}_{s,t}^{(i)} = \frac{X_{i,s}}{r\deg(s)} - \frac{X_{i,t}}{r\deg(t)} = \frac{Y_{i,s}}{r\deg(s)} + \frac{Y_{i,t}}{r\deg(t)} \]
11. \[ \text{return } \hat{\delta}_{s,t} = \sum_{i=0}^{t-1} \hat{\delta}_{s,t}^{(i)} \]

Proof of Theorem 4.2. We first note that the running time and query complexity of the algorithm are \( O(t^4) = O(t^4(\log t)/\epsilon^2) \).

In the following, we prove the correctness of the algorithm. We first present a basic property of effective resistance. Let \( Q = D^{-1/2}AD^{-1/2} \). Recall that \( L = D - A = D^{1/2}(I - Q)D^{1/2} \) and that \( R_G(s, t) = \chi_{s,t}L^1{\chi_{s,t}}^\top \). Note that \( Q = D^{-1/2}AD^{-1/2} = D^{1/2}PD^{-1/2} \) is symmetric and is similar to \( P \) (as the diagonal matrix \( D \) is invertible, which in turn follows from Assumption 4.1). We let \( \lambda_1 \geq \lambda_2 \geq \lambda_3 \geq \cdots \geq \lambda_n \) be the eigenvalues of \( Q \) (and also \( P \)) by the similarity of \( P \) and \( Q \), with corresponding (row) orthonormal eigenvectors \( w_1, w_2, w_3, \ldots, w_n \), i.e., \( w_i^TQ = \lambda_i w_i \). It is known that \( \lambda_1 = 1 \) and \( w_1 = 1_{\epsilon}D^{1/2}/\sqrt{2m} \).

Lemma 4.3. It holds that
\[
R_G(s, t) = \chi_{s,t} \sum_{i=0}^{\infty} P^iD^{-1}{\chi_{s,t}}^\top.
\]

Proof. By the spectral decomposition of \( Q \), we have that for any integer \( i \geq 0 \),
\[
Q^i = \sum_{j=1}^{n} \lambda_j^i w_j \otimes w_j = w_1^i w_1 + \sum_{j=2}^{n} \lambda_j^i w_j \otimes w_j.
\]

Since \( L = D - A = D^{1/2}(I - Q)D^{1/2} \), we have that \( L^1 = D^{-1/2}(I - Q)^{1/2} \).

\[
\]
Claim 4.4. It holds that \( |\mathcal{R}_G(s, t) - \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} | \leq \epsilon. \)

Proof. It holds that
\[
|\mathcal{R}_G(s, t) - \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} | = \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} - \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} \]
\[
\leq \sum_{i=0}^{\ell-1} \sum_{j=0}^{n} \alpha_j \beta_j \leq \frac{2 \lambda \ell}{1 - \lambda} \leq \frac{\epsilon}{2},
\]
where the last inequality follows from \( \ell = \frac{\log(4/(\epsilon - \lambda \ell))}{\log(1/\lambda)}. \)

Claim 4.5. With probability at least 9/10,
\[
|\hat{\delta}_{s, t} - \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} | \leq \frac{\epsilon}{2}.
\]

Proof. We observe that for any \( i \geq 0, \)
\[
\chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} = (1s - 1t) P_i \cdot D^{-1} (1s - 1t) = 1P_i 1^T_{\deg(s)} - 1P_i 1^T_{\deg(t)} - 1P_i 1^T_{\deg(s)} + 1P_i 1^T_{\deg(t)}.
\]

Note that for any \( 0 \leq i \leq \ell - 1, \) in the algorithm, we perform \( r \) random walks of length \( i \) from \( s. \) Since \( X_{s, t} \) is the number of walks that end at \( s \) and \( 1P_i 1^T_{\deg(s)} \) is exactly the probability of a random walk of length \( i \) from \( s \) to \( s, \) we have that \( \mathbb{E} X_{s, t} = r \cdot 1P_i 1^T_{\deg(s)}. \)

Furthermore, by the Chernoff-Hoeffding bound,
\[
P\left| \frac{X_{s, t}}{r \deg(s)} - \frac{1P_i 1^T_{\deg(s)}}{\deg(s)} \right| \geq \frac{\epsilon}{8\ell} \leq \frac{r}{40\ell},
\]
where the last inequality follows from \( r = 40\ell^2(\log(80\ell))/\epsilon^2. \) Similarly,
\[
P\left| \frac{X_{s, t}}{r \deg(t)} - \frac{1P_i 1^T_{\deg(t)}}{\deg(t)} \right| \leq \frac{\epsilon}{8\ell},
\]
\[
P\left| \frac{Y_{s, t}}{r \deg(s)} - \frac{1P_i 1^T_{\deg(s)}}{\deg(s)} \right| \leq \frac{\epsilon}{8\ell},
\]
\[
P\left| \frac{Y_{s, t}}{r \deg(t)} - \frac{1P_i 1^T_{\deg(t)}}{\deg(t)} \right| \leq \frac{\epsilon}{8\ell}.
\]

Thus by a union bound, it holds that
\[
|\hat{\delta}_{s, t} - \sum_{i=0}^{\ell-1} \chi_{s, t} P_i \cdot D^{-1} \chi_{s, t} | \leq \frac{\epsilon}{2},
\]
with probability \( 1 - 4\cdot \frac{\epsilon}{8\ell} = \frac{9}{10}. \)

Therefore, with probability at least 9/10, it holds that
\[
|\mathcal{R}_G(s, t) - \hat{\delta}_{s, t} | \leq \epsilon.
\]

This finishes the proof of Theorem 4.2.

The second algorithm: EstEff-TransProb-Collision. In the previous algorithm, we used the simple Monte Carlo approach to approximate the transition probabilities (which correspond to Line 4 and 5 in Algorithm 1). Now we give a more efficient procedure to estimate the transition probability \( 1P_i 1^T. \) Such an algorithm is based on the idea of treating the term \( 1P_i 1^T. \) as a collision probability of two random walks of length \( i/2, \) starting from \( s \) and \( t, \) respectively. In particular, if \( p = 1P_i 1^T, \) then for typical vertices \( s, t, \) we can approximate the probability \( p \) in \( O(1/\sqrt{p}) \) time, in contrast to the \( O(1/p) \) time from the Monte Carlo approach. This idea of approximating transition probability has been given in [5]. We use this idea to present a new algorithm whose performance guarantee is given in the following theorem.

Theorem 4.6. Suppose that Assumption 4.1 holds. Suppose further that for any \( i \leq \ell, \)
\[
\|1P_i D^{-1/2}\|_2^2 \|1P_i D^{-1/2}\|_2^2 \leq \beta_i,
\]
for some parameters \( \beta_i \)'s. The Algorithm 2 (i.e., EstEff-TransProb-Collision(G, s, t)) outputs an estimate \( \hat{\delta}_{s, t} \) such that with probability
Thus, the running time and query complexity of the algorithm are $O(\sum_{i=0}^{3} r_i) = O(\frac{\log(4/\epsilon)}{\epsilon^{2}})$.

On the choice of $\beta_i$: Note that the algorithm is parametrized by $\beta_i$. We note that for expander graphs or graphs with low mixing time, it holds that $\beta_i$ is a number that exponentially decreases in terms of $i$, i.e., $\beta_i \leq \epsilon^i$ for some constant $\epsilon < 1$, as long as $\ell$ is not too large. The reason is that in an expander graph $G$ with $\phi(G) \geq \phi$ for some constant $\phi$, it holds that $||1 \cdot P^i \cdot D^{-1/2}||_2 \leq \frac{1}{\text{vol}(G)} + (1 - \frac{\phi}{4})^{2i}$ for any starting vertex $s$ (see e.g., [12]). Therefore, in the case, the running time in Theorem 4.6 will be dominated by $O(\frac{\log(4/\epsilon)}{\epsilon^{2}})$, which is faster than Algorithm 1.

Algorithm 2: EstEff-TransProb-Collision($G, \epsilon, s, t$)

```
1 \ell \leftarrow \frac{\log(4/(\epsilon(1-\ell)))}{\log(1/\ell)}
2 for i := 0, 1, \ldots, \ell - 1 do
3 \quad r_i \leftarrow 20000(\sqrt{\frac{D_{s,t}}{\epsilon^2}} + \frac{\epsilon^i}{\ell})
4 \quad Perform \, r_i \, independent \, random \, walks \, of \, length
5 \quad i_1 := [i/2] \, starting \, at \, s \, (resp., \, t), \, and \, let \, X_{s,t} \in \mathbb{R}^V
6 \quad (resp., \, Y_{s,t} \in \mathbb{R}^V) \, be \, a \, row \, vector \, whose \, \ell \, th \, component \, is \, the \, fraction \, of \, random \, walks \, from \, s \, (resp., \, t) \, that
7 \quad end \, up \, at \, \alpha, \, divided \, by \, \deg(\alpha)
8 \quad Perform \, r \, independent \, random \, walks \, of \, length
9 \quad i_2 := [i/2] \, starting \, at \, s \, (resp., \, t), \, and \, let \, Y_{s,t} \in \mathbb{R}^V
10 \quad (resp., \, X_{s,t} \in \mathbb{R}^V) \, be \, a \, row \, vector \, whose \, \ell \, th \, component \, is \, the \, fraction \, of \, random \, walks \, from \, s \, (resp., \, t) \, that
11 \quad end \, up \, at \, \alpha, \, divided \, by \, \deg(\alpha)
12 Set \delta(i)_{s,t} = X_{s,t} \cdot Y_{s,t} - X_{s,t} \cdot Y_{t} - X_{t} \cdot Y_{s,t} + X_{t} \cdot Y_{t}
13 return \delta_{s,t} = \sum_{i=0}^{\ell-1} \delta(i)_{s,t}
```

Proof of Theorem 4.6: W.l.o.g. we consider the case that the length $i$ of the random walk is even. Note that for any $s, t$,

$$1 \cdot P^i \cdot D^{-1} = 1 \cdot (D^{-1} A)^i \cdot 1^T = 1 \cdot (D^{-1} A)^i \cdot (D^{-1} A)^{1/2} \cdot D^{1/2}$$

Thus,

$$\frac{1 \cdot P^i \cdot D^{-1/2}}{\deg(t)} = (1 \cdot P^i \cdot D^{-1/2}, 1 \cdot P^i \cdot D^{-1/2}).$$

Note that for any vertex $u$, the quantity $\{1 \cdot P^i \cdot D^{-1/2}(v)\}$ is the probability of a length-$i(2)$ random walk that starts from $s$ and ends at vertex $v$, divided by $\deg(u)$; and the quantity $\{1 \cdot P^i \cdot D^{-1/2}\}$ is the probability of a length-$i(2)$ random walk that starts from $s$ and ends at vertex $u$, divided by $\deg(u)$.

Now we use the argument in the proof of Lemma 19 in [10]. Specifically, let $Z_{s,t} = \tilde{X}_{s,t} \cdot \tilde{Y}_{s,t}$, where $\tilde{X}_{s,t} \cdot \tilde{Y}_{s,t}$ are defined in Algorithm 2. Then $E(Z_{s,t}) = (D^{-1/2} P^i \cdot s_t)^T (D^{-1/2} P^i \cdot t)$. By Chebyshev’s inequality and Lemma 19 in [10], we get $\Pr[|Z_{s,t} - E(Z_{s,t})| > \varepsilon] < (\varepsilon^2) \cdot (\frac{\beta_i}{\ell^2} + \frac{\epsilon^i}{\ell}) \leq \frac{1}{2\ell}$, as we have chosen $\ell = 20000(\sqrt{\frac{D_{s,t}}{\epsilon^2}} + \frac{\epsilon^i}{\ell})$ in the algorithm. Then the statement of the theorem follows by analogous argument from the proof of Theorem 4.2.

Finally, we remark that the success probabilities of both algorithms EstEff-TransProb and EstEff-TransProb-Collision can be boosted to $1 - \frac{1}{\text{poly}(n)}$ by standard median trick, i.e., repeatedly run these algorithms $O(\log(n))$ times and output the median. On graphs with bounded mixing time, which correspond to graphs such that $1 - \varepsilon \geq \frac{1}{\text{poly}(\log(n))}$, the algorithms run in $O(\text{poly}(\log(n)/\varepsilon))$ time.

4.2 Algorithms based on commute times of random walks

In this section, we provide two algorithms based on the well known connections between effective resistances and commute time/visiting probability in random walks. Let $\gamma > 0$ be a threshold parameter.

The first algorithm: EstEff-MC: We can use the commute time $\kappa(s, t)$ to approximate $R_{s,t}(s, t)$. We make use of the following results.

Lemma 4.7 ([9, 30]): It holds that $\kappa(s, t) = 2mR_{s,t}(s, t)$.

Lemma 4.8 (Proposition 2.3 in [26]): The probability that a random walk starting at $s$ visits $t$ before returning to $s$ is $\kappa(s, t) \cdot \pi(s)$, where $\pi(s) = \frac{\deg(s)}{2m}$ is the stationary probability of $s$.

We obtain the following corollary by Lemmas 4.7 and 4.8.

Corollary 4.9: The probability $p(s, t)$ that a random walk starting at $s$ visits $t$ before returning to $s$ is $\frac{1}{R_{s,t}(s, t) \cdot \deg(s)}$. In particular, if $R_{s,t}(s, t) \leq 1$, then $p(s, t) \geq \frac{1}{\gamma \deg(s)}$.

The corollary above suggests the Monte Carlo algorithm below. The algorithm performs a number of random walks, starting at vertex $s$. Then it essentially count how many times the random walk traverses from $s$ to $t$ and back.

Algorithm 3: EstEff-MC($G, s, t, y, \gamma$)

```
1 W.l.o.g., suppose that $\deg(s) \leq \deg(t)$
2 $N_0 \leftarrow 3\log\gamma \cdot \deg(s), X \leftarrow 0$
3 for $i = 1, \ldots, N_0$ do
4 \quad Perform a random walk from $s$, and stop the walk
5 \quad (1) if the walk has visited $t$ and then returns to $s$.
6 \quad (2) or if the walk has return to $s$ before visiting $t$.
7 \quad If the item (1) occurs, $X \leftarrow X + 1$
8 return $\frac{N_0}{\deg(s) \cdot X}$
```
Theorem 4.10. Assume that that \( R_G(s, t) \leq \gamma \). Let \( N_0 = \frac{3 \ln \gamma \cdot \deg(s)}{\varepsilon^2} \). Then with probability 2/3, Algorithm 3 (i.e., EstEff-MC) returns an \((1 + \varepsilon)\)-approximation for \( R_G(s, t) \). The running time of the algorithm is \( O\left( \frac{\ln \gamma \cdot \deg(s)}{\varepsilon^2} \right) \).

We remark that the above algorithm runs in sublinear time if \( \gamma = o(1) \), i.e., \( R_G(s, t) \) is small enough. In other words, when the two vertices \( s, t \) are "similar" enough, our algorithm will be fast.

Proof of Theorem 4.10. In Algorithm 3, let \( X_i \) be the indicator variable that denotes the \( i \)-th random walk to be successful (where we do not abort the walk because of its length). Then \( P(X_i = 1) = p(s, t) \) where \( p(s, t) \) is as defined in Corollary 4.9. Furthermore, let \( X = \sum_{i=1}^{N_0} X_i \). Observe that \( E(X) = N_0 \cdot p(s, t) = \frac{R_{\text{eff}}(s, t) \cdot \deg(s)}{\varepsilon^2} \), where we have used that \( p(s, t) = \frac{1}{R_{\text{eff}}(s, t) \cdot \deg(s)} \).

Next, assume that \( R_{\text{eff}}(s, t) \leq \gamma \); let \( N_0 = \frac{\ln(1/\delta) \cdot 3 \cdot \gamma \cdot \deg(s)}{\varepsilon^2} \). Then, by using Chernoff and union bounds we find that \( P\left[ |X - E(X)| > \varepsilon' \cdot E(X) \right] < 2 \cdot e^{-\frac{\varepsilon'^2 \cdot N_0 \gamma}{\deg(s) \cdot \varepsilon^2}} \leq 2 \cdot \delta \) for any \( \varepsilon' > 0 \). Thus, we find that with probability at least 1 - \( 2 \delta \), \( (1 - 2\varepsilon') R_{\text{eff}}(s, t) \leq \frac{N_0}{\deg(s)} \). Now, choosing \( \varepsilon' = \varepsilon/2 \) yields the desired approximation ratio.

As a second step, we will show that each of the random walks in Algorithm 3 is expected to terminate within at most \( 2m \gamma \) steps. Consider the two cases in which the walks terminates. Let \( y_i, i \in \{1, 2\} \) denote the number of steps taken in random walk in some iteration, such that \( i = 1 \) if the first termination criterion of the loop is fulfilled and \( i = 2 \) in the other case. Then clearly, the number of steps taken in a random walk is \( \min\{y_1, y_2\} \). Furthermore, it holds that \( \min\{y_1, y_2\} \leq y_1 \). Note that \( E(y_i) \) is the commute time \( \kappa(s, t) \).

Therefore, we find that \( E(\min\{y_1, y_2\}) \leq E(y_1) = \kappa(s, t) \).

Finally, let \( \delta = 1/3 \). Then we find that Algorithm 3

- runs in expected time \( R_{\text{eff}}(s, t) \cdot N_0 \) is \( O\left( \frac{\ln \gamma \cdot \deg(s)}{\varepsilon^2} \right) \) and
- with probability at least \( 1 - \delta = 2/3 \), \( \frac{N_0}{\deg(s)} \) is an \((1 + \varepsilon)\)-approximation of \( R_{\text{eff}}(s, t) \).

This concludes the proof.

The second algorithm: EstEff-MC2. For the special case that there is an edge \((s, t)\) between the two specified vertices \( s, t \), we can also make use of the following probabilistic interpretation of effective resistance.

Lemma 4.11 ([31]). Consider an edge \((s, t)\). Then \( R_G(s, t) \) is the probability that a random walk from \( s \) visits \( t \) for the first time using \((s, t)\).

This suggests the following Monte Carlo algorithm.

Theorem 4.12. For \( R_G(s, t) > \gamma \), Algorithm 4 (i.e., EstEff-MC2) returns with probability \((1 - \delta)\) a \((1 + \varepsilon)\)-approximation of \( R_G(s, t) \).

The proof of the above theorem is deferred to Appendix B. Note that in contrast to Algorithm 3, the random walks in Algorithm 4 stop as soon as we have reached the destination vertex \( t \). Hence, one can expect that Algorithm 4 runs faster than Algorithm 3. Experimental comparisons of the running times can be found in Section 5.1.
Algorithm 6: EstEff-SpanTree(G, ε, δ, u, v)

1.  \( a \leftarrow \text{AppNumST}(G, \frac{\epsilon}{2}, \frac{\delta}{2}) \)
2.  \( b \leftarrow \text{AppNumST}(G, \frac{\epsilon}{2}, \frac{\delta}{2}) \)
3.  \( \text{return } \frac{a^n - b^n}{2} \)

Theorem 4.15. Algorithm 6 returns with probability at least \( 1 - \delta \) an estimator \( X \) such that

\[
\epsilon^{-\alpha} R_G(s, t) \leq X \leq \epsilon^{\alpha} R_G(s, t).
\]

The algorithm uses \( \hat{O}(\epsilon^{-5} + \epsilon^{-2} \log^2 n) \log \delta^{-1} \) queries.

We give the proof of Theorem 4.15 in Appendix B and remark that the above algorithm seems of theoretical interest only, as it does not perform well in practice.

5 EXPERIMENTS

In this section, we show our experimental results. The experiments were conducted on a Linux server with Intel Xeon E5-2643 (3.4GHz) and 768GB of main memory, and all the programs were implemented in C++ and compiled with g++ 4.8.4. The graphs used in the experiments are taken from SNAP\(^3\) and basic information about the graphs is given in Table 1. We generated query pairs by randomly sampling edges 1,000 times with replacements.

| Dataset  | \( n \) | \( m \) |
|----------|--------|--------|
| Facebook | 4,039  | 88,233 |
| DBLP     | 317,080| 1,049,869|
| YouTube  | 1,134,891| 2,987,627|

We implemented the following algorithms:

- **Exact**: This method first applies the QR decomposition to the Laplacian as preprocessing and computes effective resistance according to its definition, i.e., \( R_G(s, t) := \chi_{X_s}^t L^* \chi_{X_s}^t \).
- **HAY** [17]: This method computes effective resistances of all the edges at once by sampling spanning trees and it is still state-of-the-art for this problem. We fixed the number of sampled spanning trees to 10,000.
- **TP**: Implementation of Algorithm 1. We set \( \epsilon = \lambda = 0.1 \).
- **TP-C**: Implementation of Algorithm 2. We set \( \epsilon = \lambda = 0.1 \).
- **MC**: Implementation of Algorithm 3. We set \( \epsilon = \gamma = 0.1 \).
- **MC2**: Implementation of Algorithm 4. We set \( \epsilon = \gamma = 0.1 \).
- **ST**: Implementation of Algorithm 6. We set \( \epsilon = 0.1 \).

To implement each algorithm, we used the same number of random walks as the one given in the corresponding pseudocode.

5.1 Running time

Figure 1 shows the running time of each method. For our methods, we plotted the running time for the 1,000 queries in increasing order. For Exact and HAY, we plotted their preprocessing time.

We do not show the running time of Exact on DBLP and YouTube because it did not terminate in 8 hours.

We can first observe that MC and ST on Facebook are as slow as previous (polynomial-time) algorithms, and hence we do not consider those algorithms for other graphs.

We can observe that TP, TP-C, and MC2 are much faster than the existing methods. Note that the running time of MC2 depends on the queried edge \((s, t)\) because it runs until the random walk starting at \( s \) reaches \( t \). In contrast, the running time of TP and TP-C is almost independent of the queried edge, which is preferable. A reason that TP-C is slower than TP is that we need to compute inner products in TP-C (Line 6 of Algorithm 2).

5.2 Accuracy

Figure 2 shows the accuracy of existing and our methods. For each method, we computed the relative error as \( |R - \tilde{R}|/R \) for each query, where \( R \) is the exact effective resistance for Facebook using Exact and the one estimated by HAY for DBLP and YouTube, and \( \tilde{R} \) is the estimated effective resistance. Then, we plotted the 1,000 relative errors after sorting them in increasing order. Except for ST, the relative error of our methods are within 0.1 for most of the queries, as expected from the choice \( \epsilon = 0.1 \). Also, the results for Facebook justifies the use of HAY on DBLP and YouTube as the baseline method. In Figure 2(a), the results of TP and TP-C are very close such that their lines overlap. The fact that the lines change concavity twice appears to be a universal phenomenon for probabilistic distributions.

In Figure 3, each blue point represents \((R, \tilde{R})\) for a query, where \( R \) and \( \tilde{R} \) are as defined in the paragraph above. MC2 shows the best accuracy on DBLP and YouTube. Accuracy of TP-C is comparable to that of TP. Recalling that TP runs faster than TP-C, we can conclude that TP is superior to TP-C.

6 CONCLUSION

In this paper, we developed a number of local algorithms for estimating the pairwise effective resistances, a fundamental metric for measuring the similarity of vertices in a graph. Our algorithms explore only a small portion of the graph while provides a good approximation to \( R_G(s, t) \) for any specified \( s, t \). Our algorithms are desirable in applications where the effective resistances of a small number of vertex pairs are needed. Our experiments on benchmark datasets validate the performance of these local algorithms.

ACKNOWLEDGMENTS

YY. is supported in part by JSPS KAKENHI Grant Number 17H04676, 18H05291, and 20H05965.

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Figure 1: Running time

(a) Facebook

(b) DBLP

(c) YouTube

Figure 2: Relative error

(a) Facebook

(b) DBLP

(c) YouTube

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Figure 3: Quality of estimated effective resistance

We present here the proofs of two theorems in Section 4.

Theorem A.1 (The Chernoff-Hoeffding bound). Let $s \geq 1$. Let $X = \sum_{1 \leq i \leq s} X_i$, where $X_i, 1 \leq i \leq s$, are independently distributed in $[0,1]$. Then for all $t > 0$,

$$
\mathbb{P}[|X - \mathbb{E}[X]| > t] \leq e^{-2t^2/s}.
$$

A THE CHERNOFF-HOEFFDING BOUND

We make use of the following Chernoff-Hoeffding bound (see Theorem 1.1 in [14]).

B MISSING PROOFS OF SECTION 4

We present here the proofs of two theorems in Section 4.
Proof of Theorem 4.12. In Algorithm 4, let $X_i$ be the indicator variable that denotes the $i$-th random walk to be successful (regardless of its length). Then $P(X_i = 1) = R_G(s,t)$. Furthermore, let $X = \sum_1^M X_i$. Observe that $E(X) = M_0 \cdot R_G(s,t)$.

Next, assume that $R_G(s,t) > \gamma$; let $M_0 = \ln(1/\delta')/3 \cdot \gamma^2 \cdot \ln(1/\delta')$. Using Chernoff and union bounds we find that $P[|X - E(X)| > \epsilon \cdot E(X)] < 2 \cdot e^{-\lambda M_0 R_G(s,t)/3} < 2 \cdot \delta'$ for any $\epsilon, \delta' > 0$. Thus, we find that with probability at least $1 - 2\delta'$, $(1 - \epsilon)R_G(s,t) \leq X/M_0 \leq (1 + \epsilon)R_G(s,t)$. Now, choosing $\delta' = \frac{\delta}{2}$ yields the desired approximation ratio.

Proof of Theorem 4.15. By Lemma 4.14, with probability $1 - \frac{\delta}{2}$, the $a$ returned in line 1 of Algorithm 6 satisfies $|a - \frac{\log(T(G'))}{n-1}| \leq \frac{\epsilon}{2}$. Similarly, with the same probability, $|b - \frac{\log(T(G))}{n}| \leq \frac{\epsilon}{2}$. By the union bound, this implies that with probability $1 - \delta$, $a(n-1) - bn \leq \frac{\epsilon}{2}(2n-1) - \log(T(G)) + \log(T(G'))$. Thus, we find that $e^{a(n-1)} - e^{\epsilon n} \leq e^{\epsilon n} \cdot \frac{T(G')}{T(G)}$. Similarly, it holds that $e^{b n} \geq e^{\epsilon n} \cdot \frac{T(G)}{T(G')}$. Let $X = e^{a(n-1)}$. Then by Lemma 4.13, $e^{\epsilon n} R_G(s,t) \leq X \leq e^{\epsilon n} R_G(s,t)$. This yields the desired approximation ratio. \qed