

# Memory-Aware Framework for Efficient Second-Order Random Walk on Large Graphs

Yingxia Shao<sup>★</sup>, Shiyue Huang<sup>§</sup>, Xupeng Miao<sup>§</sup>, Bin Cui<sup>§</sup>, Lei Chen<sup>△</sup>

<sup>★</sup>Beijing Key Lab of Intelligent Telecommunications Software and Multimedia, BUPT

<sup>§</sup>School of EECS & Key Laboratory of High Confidence Software Technologies (MOE), Peking University

<sup>△</sup>Hong Kong University of Science and Technology

shaoyx@bupt.edu.cn, {huangshiyue, xupeng.miao, bin.cui}@pku.edu.cn, leichen@cse.ust.hk

## ABSTRACT

Second-order random walk is an important technique for graph analysis. Many applications use it to capture higher-order patterns in the graph, thus improving the model accuracy. However, the memory explosion problem of this technique hinders it from analyzing large graphs. When processing a billion-edge graph like Twitter, existing solutions (e.g., alias method) of the second-order random walk may take up 1796TB memory. Such high memory overhead comes from the memory-unaware strategies for node sampling across the graph.

In this paper, to clearly study the efficiency of various node sampling methods in the context of second-order random walk, we design a cost model, and then propose a new node sampling method following the acceptance-rejection paradigm to achieve a better balance between memory and time cost. Further, to guarantee the efficiency of the second-order random walk within arbitrary memory budgets, we propose a memory-aware framework on the basis of the cost model. The framework applies a cost-based optimizer to assign desirable node sampling method for each node in the graph within a memory budget while minimizing the time cost. Finally, we provide general programming interfaces for users to benefit from the memory-aware framework easily. The empirical studies demonstrate that our memory-aware framework is robust with respect to memory and is able to achieve considerable efficiency by reducing 90% of the memory cost.

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## CCS CONCEPTS

- Theory of computation → Graph algorithms analysis;
- Computing methodologies → Parallel algorithms.

## KEYWORDS

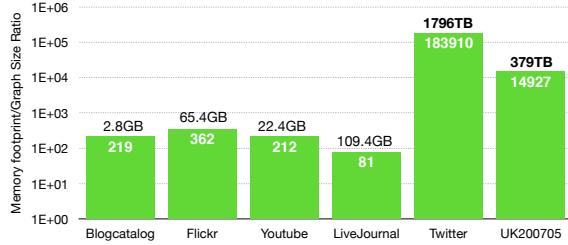
Random walk; Memory efficient; Graph algorithm; Large-scale

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## 1 INTRODUCTION

Random walk is widely used for graph analysis like proximity computation [17, 32, 38, 42], aggregate estimation [16, 24] and graph embedding [3, 10, 25, 35]. Most existing models use the first-order random walk [14], which assumes the next node to be visited only depends on the current node. However, the first-order random walk abstracts an oversimplification of real-world systems. An example is user trails on the Web [40], where nodes are Web pages and interactions are users navigating from one Web page to another. A user's next page visit not only depends on the last page but also is influenced by the sequence of previous clicks, which are called higher-order dependencies. The first-order random walk fails to capture such higher-order dependencies. Second-order random walk is designed to model the higher-order dependencies, thus improving the accuracy of applications. One popular and recent application of second-order random walk is to learn high-quality embeddings for graph analysis tasks, like node classification [1, 36], link prediction [18, 39]. Node2vec [9, 44] is one of the most successful graph embedding models. It uses the second-order random walk to encode nodes from the same community closely together and outperforms the deepwalk [25], a model using first-order random walk. Besides the graph embedding, second-order



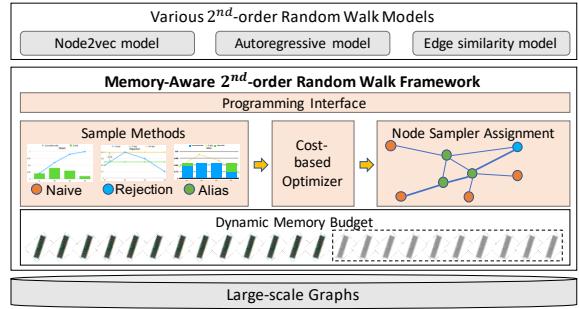
**Figure 1: The ratio of total memory footprint to the memory size of the graph when running node2vec. The number at the top of each bar is the corresponding total memory footprint.**

random walk is also successfully applied to find meaningful communities [2, 30] and model sequential data [38, 40].

Unlike the first-order random walk, the second-order one visits the next node depending on both the current node and the previous node. Given that the previous and current nodes are  $u$  and  $v$ , the probability of the next node  $z$  is  $p(z|vu)$ . We call  $p$  as *edge-to-edge* transition probability (*e2e* for short). Then for a graph  $G(V, E)$ , there are in total  $|E|d$  different *e2e* probabilities, where  $d$  is the average degree of  $G$ . The management of  $|E|d$  *e2e* probabilities is the bottleneck for running second-order random walks on large graphs.

A straightforward solution is to build the *e2e* distribution on demand, however, this approach has low efficiency and the sampling time cost of each node is linear to the node degree. To improve the sampling efficiency, current solutions store the whole distribution in memory with sophisticated data structures (e.g., alias table [37]). But this incurs the memory explosion problem [35, 44]. From Figure 1, we find that when running node2vec over a billion-edge graph like Twitter, which has about 41.6M nodes and 2.4B edges, current solutions require 1796TB memory that is 183910× larger than the graph size of Twitter, and it is really expensive to set up a machine/cluster to have that much memory. Even for much smaller graphs (e.g., Youtube, LiveJournal), the existing empirical studies [35] also demonstrate that node2vec encounters out-of-memory problem on a commodity server. It is obvious that current memory-intensive solutions are not cost-efficient approaches to process large graphs in the real world. Therefore, the research question is *how to maximize the efficiency of executing second-order random walk on large graphs within a memory budget*.

In this paper, we dissect the random walk operation into a sequence of node sampling, where a node generates samples from a discrete (non-)uniform distribution (e.g., *e2e* distribution). Then we design a cost model to reveal the trade-off between time and memory cost for different node sampling methods. Specifically, given a single node  $v$ , the node sampling method in the straightforward solution, called *naive method*, entails  $O(d_v)$  time cost and  $O(1)$  memory cost,



**Figure 2: Overview of the memory-aware framework.**

where  $d_v$  is the degree of node  $v$ . The aforementioned efficient node sampling with alias table, called *alias method*, entails  $O(1)$  time complexity and  $O(d_v^2)$  memory complexity. Inspired by the acceptance-rejection paradigm, we introduce a new sample method called *rejection method*. The rejection method uses the first-order random walk to sample node, accepts the sampled node with probability  $\beta$ , named acceptance ratio, and repeats the sampling until generating an accepted node. The rejection method entails  $O(C_v)$  time complexity and  $O(d_v)$  memory complexity. Here  $C_v$  is a model-related parameter and we prove  $C_v$  is smaller than  $d_v$ . Compared with the naive method and the alias method, rejection method achieves a better balance between memory and time cost.

From the above cost analysis, we find that these sampling methods have advantages. However, they are fragile and memory-unaware, i.e., they occupy fixed memory footprint, and cannot boost the efficiency when the available memory is larger than the required one or ensure the success of the task when available memory is insufficient. Therefore, we propose a memory-aware second-order random walk framework to guarantee the high efficiency on arbitrary memory budgets. The core of the framework is a cost-based optimizer. Based on the cost model, the optimizer assigns proper node sampling method for each node in the graph while minimizing the total time cost. The assignment problem is modeled as a multiple-choice knapsack problem [33], and we develop an effective greedy algorithm to generate the assignment in high efficiency. Further, considering the dynamic memory budgets in practice, we also introduce an adaptive version of the greedy algorithm, so that we can efficiently update the node sampling assignment when the memory budget changes including increase and decrease.

Finally, with our technical contributions, we carefully implement the memory-aware framework (Figure 2) as a middleware for various second-order random walk models. In this work, we implement two popular second-order random walk models – node2vec and autoregressive model [27] for the experiments. Further, to easily benefit from the aforementioned cost-based optimizer, the framework provides flexible

programming interfaces for users to define their own sample methods and the second-order random walk models. We evaluate the framework on six real-world large graphs including two billion-edge graphs. The empirical studies demonstrate that our memory-aware framework is robust with respect to memory and is able to achieve comparable efficiency by saving 90% of the memory cost which is required by the existing solutions.

We summarize our contributions as follows:

- We propose a memory-aware framework for second-order random walk on large graphs.
- We develop a cost-based optimizer to find a node sampler assignment in high efficiency.
- We introduce a rejection method for second-order random walk.
- We conduct extensive experiments on real-world datasets and show the superiority of our framework compared with the existing methods.

**Organization.** In Section 2, we present the backgrounds of this work. In Section ??, we introduce the rejection method for second-order random walk and establish the cost model. Then we elaborate the cost-based optimizer, including node sampler assignment, theoretical analysis and its adaptive version in Section 5, followed by the introduction of the memory-aware framework in Section 5.4. In Section 6, we present the experimental results. Finally, we discuss the related work and give the conclusion in Section 7 and 8.

## 2 PRELIMINARY

A graph  $G = (V, E)$  is defined by a set  $V$  of vertices and a set  $E$  of pairwise relations (edges) among vertices in  $V$ . We use  $v$  to represent a vertex in  $G$ ,  $N(v)$  to represent the neighborhood, and the degree of  $v$  is  $d_v$ . An edge is represented by  $(u, v)$  and the corresponding weight is  $w_{uv}$ .

### 2.1 Second-order Random Walk

Random walk on a graph is a stochastic process, that describes a walk that consists of a succession of random selected vertices. *First-order random walk* selects a vertex  $z$  based on the state of last vertex  $v$ , and it follows the transition distribution  $p(z|v) = \frac{w_{vz}}{W_v}$ , where  $W_v = \sum_{t \in N(v)} w_{vt}$ . *Second-order random walk* selects a vertex  $z$  based on the states of last two vertices  $u$  and  $v$ , and the transition distribution is  $p(z|vu)$ . For simplicity, the distribution  $p(z|v)$  is called *node-edge* distribution (n2e for short), while the one  $p(z|vu)$  is called *edge-edge* distribution (e2e for short).

The concrete e2e distribution is application dependent. Here we introduce two popular second-order random walk models: node2vec model and autoregressive model.

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### Algorithm 1 Second-order random walk generation procedure

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**Input:** graph:  $G(V, E)$ , start node:  $v$ , length:  $maxLen$

**Output:** random walk:  $L$

```

1:  $L = \{v\}$  // an array
2: for  $t$  in  $1..maxLen$  do
3:   if  $t == 1$  then
4:      $u_{t-1} = L[t-1]$ 
5:     Node Sampler: draw node  $u_t$  from  $p(u_t|u_{t-1})$ 
6:   else
7:      $u_{t-1} = L[t-1]$ ,  $u_{t-2} = L[t-2]$ 
8:     Node Sampler: draw node  $u_t$  from  $p(u_t|u_{t-1}, u_{t-2})$ 
9:   end if
10:   $L.append(u_t)$ 
11: end for
12: return  $L$ 

```

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**Node2vec model.** As briefly mentioned before, node2vec is a network embedding algorithm using second-order random walk. The e2e distribution in node2vec defines the probability of walking from  $(u, v)$  to  $(v, z)$  as  $p_{uvz} = \frac{w'_{vz}}{W'_v}$ , where  $W'_v = \sum_{t \in N(v)} w'_{vt}$ , and  $w'_{vz}$  is defined as follows,

$$w'_{vz} = \begin{cases} \frac{w_{vz}}{a} & l_{uz} = 0 \\ \frac{w_{vz}}{b} & l_{uz} = 1 \\ \frac{w_{vz}}{b} & l_{uz} = 2 \end{cases} \quad (1)$$

where  $l_{uz}$  is the unweighted distance between vertices  $u$  and  $z$ , and  $a$  and  $b$  are two hyperparameters controlling the affinity of local structure. Both  $a$  and  $b$  are positive real values.

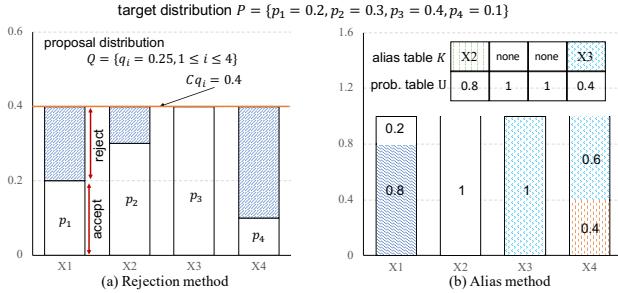
**Autoregressive model.** The autoregressive model [27] is used to compute second-order pagerank [38]. The e2e distribution is  $p_{uvz} = \frac{p'_{uvz}}{\sum_{t \in N(v)} p'_{uvt}}$ , where  $p'_{uvz} = (1 - \alpha)p_{vz} + \alpha p_{uz}$ ,  $p_{vz} = \frac{w_{vz}}{W_v}$ , and  $0 \leq \alpha < 1$ .

Note that in addition to the two models introduced above, there are many other second-order random walk models, such as the one utilizing network flow data [38], edge similarity model [19]. In this paper, we concentrate on the two models discussed here as representatives.

**2.1.1 Node Sampler.** Algorithm 1 lists the general procedure of producing a second-order random walk on graph  $G$ . From the algorithm, it is easy to see that the core operation is the selection of a node from the neighborhood (Lines 5 and 8). We name it as *node sampler* in this paper. The main logic of node sampler is to apply different sampling methods to sample nodes. The efficiency of node sampler is heavily related to the sampling methods.

### 2.2 Three Sampling Methods

Given a discrete probability distribution  $P = \{p_i\}$ ,  $i = 1..n$ , where  $p_i$  is the probability of element  $i$ , and  $n$  is the number of elements, the sampling method is to draw an element  $x$



**Figure 3: The examples of using rejection method and alias method to sample a target distribution  $P$ .**

on the basis of distribution  $P$ , i.e.,  $P[x = i] = p_i$ . There are many efficient sampling methods to realize the nonuniform sampling. In this paper, we focus on naive method, rejection method, and alias method.

**Naive method** [22]. This method generates a uniform  $[0, 1]$  random variable  $r$ , and put  $x = i$  if

$$\sum_{k=1}^{i-1} p_k < r \leq \sum_{k=1}^i p_k \quad (p_0 = 0). \quad (2)$$

We can apply various search techniques (e.g., linear search, binary search, etc.) to locate the position of  $r$  in the cumulative distribution of  $P$ . The sampling time complexity is  $O(n)$  or  $O(\log(n))$  while the memory complexity is  $O(n)$ .

**Rejection method** [28]. This method samples an element  $i$  from the target distribution  $P$  by instead sampling from a proposal distribution  $Q = \{q_i\}$  and accepting the outcome from  $Q$  with probability  $\frac{p_i}{Cq_i}$ , repeating the draws from  $Q$  until an outcome from  $Q$  is accepted. Usually the proposal distribution  $Q$  is chosen by user and it is easier to sample than the target distribution  $P$ . In addition,  $C$  is called *bounding constant*, and satisfies  $p_i \leq Cq_i$  for all values of  $i$ . The average time complexity of rejection sampling is  $O(C)^1$ .

Figure 3(a) shows an example of using rejection method to sample a target distribution  $P = \{0.2, 0.3, 0.4, 0.1\}$ . We define the proposal distribution  $Q$  to be the uniform distribution, i.e.,  $Q = \{0.25, 0.25, 0.25, 0.25\}$ , and set  $C$  to be 1.6. Then the acceptance probabilities of four elements  $X1, X2, X3, X4$  are 0.5, 0.75, 1, 0.25 respectively.

**Alias method** [37]. Alias method converts target distribution into a uniform distribution over (possibly degenerate) binary outcomes. Since the target distribution is nonuniform, it means that some elements have more than  $\frac{1}{n}$  probability, and some have less. For each element having less than  $\frac{1}{n}$ , alias method constructs an alias-outcome by combining it with a small portion of some probability mass from one of the higher-probability elements, topping up the total probability to  $\frac{1}{n}$ . Internally, alias method maintains two tables, a probability table  $U = \{u_i\}$  and an alias table  $K = \{k_i\}$

(for  $1 \leq i \leq n$ ), where  $u_i$  is the probability of choosing element at index  $i$  as an outcome and  $k_i$  is the alias-outcome with respect to  $i$ . To sample a value, it first generates a uniform  $[1, n]$  random variable  $x$  to determine an index into the two tables. Then it generates a  $[0, 1]$  random variable  $r$ , and on the basis of the probability  $u_x$ , the final outcome is

$$\begin{cases} x & r \leq u_x, \\ k_x & \text{otherwise.} \end{cases}$$

Figure 3(b) illustrates the probability table and alias table when using alias method to sample the target distribution  $P = \{0.2, 0.3, 0.4, 0.1\}$ . The algorithm typically entails  $O(n)$  time complexity to build the two tables, after which random elements can be drawn from the distribution in  $O(1)$  time complexity.

### 3 SECOND-ORDER RANDOM WALK WITH REJECTION

In this section, we introduce the rejection method for second-order random walk. Considering that the time cost of rejection method is related to the bounding constant, we also empirically and theoretically analyze the distribution of the bounding constants, and introduce a simple but effective solution for bounding constant estimation.

#### 3.1 Rejection Method for Node2vec and Autoregressive Models

According to Section 2.2, rejection method consists of target distribution  $P$ , proposal distribution  $Q$ , and bounding constant  $C$ . In the context of second-order random walk, assume the previous and current nodes are  $u$  and  $v$ , then  $P$  is  $\{\frac{w_{uz}}{W_v}\}$ ,  $Q$  is  $\{\frac{w_{uz}}{W'_v}\}$ . According to the condition  $p_i \leq Cq_i$ , we compute the bounding constant  $C_{uv}$  of an edge  $(u, v)$  as follows,

$$C_{uv} = \max\left\{\frac{P}{Q}\right\} = \max_{z \in N(v)}\left\{\frac{w_{uz}}{w_{uz}} \frac{W_v}{W'_v}\right\}. \quad (3)$$

During the sampling, the acceptance ratio  $\beta_{uvz}$  is

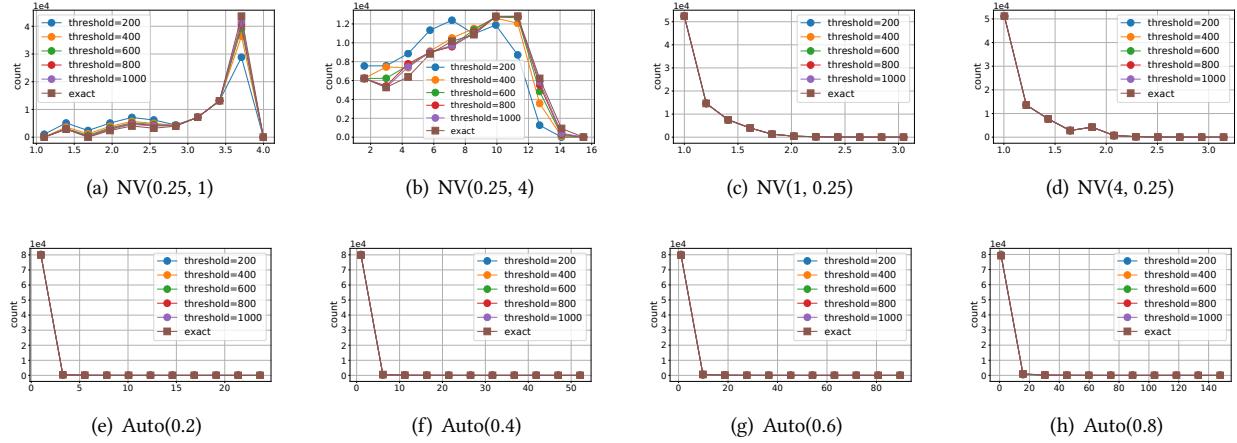
$$\beta_{uvz} = \frac{1}{C_{uv}} \frac{w_{uz}}{w_{uz}} \frac{W_v}{W'_v} = \frac{w_{uz}}{w_{uz}} \min_{t \in N(v)}\left\{\frac{w_{vt}}{w'_{vt}}\right\}. \quad (4)$$

**Rejection for node2vec.** According to the Equation 1, assume the previous node  $u$  and current node  $v$  have common neighbors, the ratio  $\frac{w_{uz}}{w_{uz}}$  belongs to  $\{\frac{1}{a}, \frac{1}{b}, 1\}$ . Then the bounding constant is  $C_{uv} = \frac{W_v}{W'_v} \max\{\frac{1}{a}, \frac{1}{b}, 1\}$ . The acceptance ratio for a vertex  $z$  is  $\beta_{uvz} = \frac{w_{uz}}{w_{uz}} \min\{1, a, b\}$ .

**Rejection for Autoregressive model.** Similarly,  $C_{uv}$  is

$$\text{The acceptance ratio } \beta_{uvz} \text{ is } \frac{(1-\alpha)+\alpha \frac{p_{uz}}{p_{uz}}}{\max_{t \in N(v)}\{(1-\alpha)+\alpha \frac{p_{vt}}{p_{vt}}\}}.$$

<sup>1</sup><http://www.columbia.edu/~ks20/4703-Sigman/4703-07-Notes-ARM.pdf>



**Figure 4: The bounding constant distribution on Flickr.** We visualize the distributions of exact bounding constants and estimated bounding constants with five different thresholds. The range of the bounding constants (x-axis) is uniformly divided into 10 buckets, i.e.,  $\frac{\max - \min}{10}$ . NV(0.25, 1) means  $a = 0.25$  and  $b = 1$  in node2vec model, and Auto(0.2) means  $\alpha = 0.2$  in autoregressive model.

### 3.2 The Distribution of Bounding Constants

The following theorem proves that the bounding constants for any edge in an unweighted graph is bounded and cannot be arbitrary large. The conclusion can be extended to the weighted graph with more complex analysis.

**THEOREM 1.** *Given an edge  $(u, v)$  in an unweighted graph, the bounding constant  $C_{uv}$  in node2vec and autoregressive model is bounded. Specifically, let  $\theta_{uv}$  be the number of common neighbors of  $(u, v)$  ( $\theta_{uv} \geq 1$ ), then the bound of  $C_{uv}$  of node2vec is*

$$C_{uv} \leq \begin{cases} \frac{d_v}{\theta_{uv}} & a \geq 1, b \geq 1, \\ \frac{d_v}{d_v} & 0 < a < 1, b \geq a, \\ \frac{d_v}{d_v - 1 - \theta_{uv}} & 0 < b < 1, a \geq b. \end{cases}$$

The one of autoregressive model is  $C_{uv} \leq \frac{d_v}{\theta_{uv}}$ .

**PROOF.** 1) The bound of  $C_{uv}$  in node2vec. In an unweighted graph,  $W_v = d_v$  and the n2e transition probability is  $\frac{1}{d_v}$ . Then

$$C_{uv} \text{ is } \frac{d_v}{\theta_{uv} + \frac{1}{a} + \frac{d_v - 1 - \theta_{uv}}{b}} \max\{\frac{1}{a}, \frac{1}{b}, 1\}.$$

Next we give the bound with different  $a$  and  $b$ .

$$\text{Case 1 } (a \geq 1, b \geq 1): C_{uv} = \frac{d_v}{\theta_{uv} + \frac{1}{a} + \frac{d_v - 1 - \theta_{uv}}{b}} \leq \frac{d_v}{\theta_{uv}}.$$

$$\text{Case 2 } (1 > a > 0, b \geq a): C_{uv} = \frac{d_v}{a\theta_{uv} + 1 + \frac{d_v - 1 - \theta_{uv}}{b}} \leq d_v.$$

$$\text{Case 3 } (1 > b > 0, a \geq b): C_{uv} = \frac{d_v}{b\theta_{uv} + \frac{b}{a} + (d_v - 1 - \theta_{uv})} \leq \frac{d_v}{d_v - 1 - \theta_{uv}}.$$

2) The bound of  $C_{uv}$  in autoregressive model. Let  $p_{vz} = \frac{1}{d_v}$  and  $p_{uz} = \frac{1}{d_u}$ , then  $C_{uv}$  is  $C_{uv} = \frac{(1-\alpha)+\alpha \frac{d_v}{d_u}}{(1-\alpha)+\alpha \frac{\theta_{uv}}{d_u}} \leq \frac{d_v}{\theta_{uv}}$ .

**Discussion.** There are two special cases (i.e.,  $\theta_{uv} = 0$  and  $d_v = 1$ ) for the bounds. When  $\theta_{uv} = 0$ , for the case

1 of node2vec, the maximum factor becomes  $\max\{\frac{1}{a}, \frac{1}{b}\}$ , then  $C_{uv}$  is still bounded by  $d_v$ . For autoregressive model,  $C_{uv} = 1$ . When  $\theta_{uv}$  is  $d_v - 1$ , the maximum factor of node2vec becomes  $\max\{\frac{1}{a}, 1\}$  and it degenerates to case 1 or case 2. The autoregressive model is not affected.

According to Theorem 1, the bound of  $C_{uv}$  is proportional to the node degree in the worst case, and the time complexity of the rejection method will be linear to the degree. In real-world graphs with standard parameter settings,  $C_{uv}$  is much smaller than the degree actually. Let us denote the average bounding constant of  $v$  as  $C_v$ , i.e.,  $C_v = \frac{1}{d_v} \sum_{u \in N(v)} C_{uv}$ . The brown line in Figure 4 shows the distribution of exact  $C_v$  on Flickr with standard parameter settings. For node2vec model, the most frequent values of  $C_v$  are below 10, and the autoregressive model has a similar trend. However, the maximal  $C_v$  of the autoregressive model is larger than the one of node2vec, this implies that the autoregressive model is more easily to be close to the bound defined in Theorem 1.

### 3.3 The Computation of Bounding Constants

As introduced in Section 2.2, the efficiency of rejection method is determined by the bounding constant. Therefore, we discuss how to compute (or estimate) the bounding constant in the context of second-order random walk. According to Equation 3, the exact  $C_v$  can be computed by enumeration which entails  $O(d_v^2)$  time complexity for node  $v$ . For the whole graph, the total time cost has the same complexity as the one of triangle counting [15], which is expensive. We resort to computing approximate ones for improving the efficiency of bounding constants computation.

On the basis of time complexity  $O(d_v^2)$ , we notice that nodes with higher degree incur larger time cost. Therefore, we estimate the bounding constants via sampling when the node degree exceeds a predefined threshold  $D_{th}$ . Formally, we estimate the  $C_{uv}$  as  $C_{uv} = \max_{z \in SN(v)} \left\{ \frac{w_{uz}}{w_{vz}} \frac{W_z}{W_v} \right\}$ , where  $SN(v)$  is a uniformly sampled neighborhood of  $v$  with size  $D_{th}$ . Then the estimation time cost is  $O(d_v D_{th})$ .

We empirically study the effectiveness of the above estimation. Figure 4 shows the distributions of estimated bounding constants on Flickr by setting  $D_{th} = [200, 400, 600, 800, 1000]$ . The results demonstrate that a relatively small threshold (e.g., 600) can achieve a good estimation. In addition, the above discussion assumes the sample size equals to the threshold. Theoretically, on the basis of the law of large numbers [8], the proposed method can achieve the exact estimation when the sample size is large enough.

## 4 COST ANALYSIS OF NODE SAMPLER

In this section, we analyze the time and memory cost of three node samplers and introduce a cost model in the context of second-order random walk.

Without loss of generality, we assume that a probability value is stored in  $b_f$  bytes and a node id is stored in  $b_i$  bytes. The unit of time cost is denoted by  $K$ .  $d_{max}$  is the maximum node degree in a graph. In addition, in second-order random walk models, the computation of biased weight is related to the existence of common neighbors, and we assume the cost of this operation is  $c$ , which is related to the node degree.

### 4.1 The Cost of Node Samplers

**Naive Node Sampler.** According to Equation 2, naive node sampler needs the cumulative distribution for sampling, and it builds the distribution on the basis of edge weights.

1) Memory cost: The sampler uses an array of length  $d_v$  to store probabilities for a node. Across the whole graph, we only need to allocate a single array of length  $d_{max}$ . In the view of a node, the average memory cost is  $\frac{b_f d_{max}}{|V|}$  bytes.

2) Time cost: The time cost consists of the distribution building cost and sampling cost. Assume linear search is used to locate the index, the sampling cost of node  $v$  is  $d_v K$ . To build the distribution, for each neighborhood, we need to check the existence of the edge between the previous node and next node, therefore, the cost is  $d_v c K$ . In total, the time cost is  $d_v(c + 1)K$ .

**Rejection Node Sampler.** Generally, the rejection node sampler follows Equations 3 and 4.

1) Memory cost: Considering that the start node of second-order random walk and successive sampled nodes for rejection are both sampled from the n2e distribution (proposal distribution  $Q$ ), we use alias method which entails  $(b_f + b_i) \times d_v$

Node Sampler	Memory Cost (M)	Time Cost (T)
Naive	$\frac{b_f d_{max}}{ V }$	$d_v(c + 1)K$
Rejection	$(2b_f + b_i)d_v$	$C_v c K$
Alias	$(b_f + b_i)(d_v^2 + d_v)$	$K$

**Table 1: Cost comparison of a node sampler by using three different sampling methods for second-order random walk. Notice that  $C_v$  is the average bounding constant of node  $v$ .  $b_f$  and  $b_i$  are the bytes of storing a non-integer and integer value respectively.**

bytes memory cost. Besides, to compute the acceptance ratio with regard to each edge  $(u, v)$ , we need to store a factor  $\min_{t \in N(v)} \left\{ \frac{w_{uz}}{w_{vz}} \right\}$ . There are total  $d_v$  different factors, the storage is  $b_f \times d_v$ . In total, for a node  $v$ , the memory cost is  $d_v \times (b_f + b_i) + d_v \times b_f$  bytes.

2) Time cost: The sampling time complexity of rejection node sampler is the average bounding constant  $O(C_v)$ , and then the cost is  $C_v K$ . In addition, for each sample, we need to compute  $w'$  to obtain the acceptance ratio, which relies on checking the existence of the edge between the previous node and next node, resulting total  $C_v$  numbers of edge existence checking. Therefore, the total time cost is  $C_v c K$ .

**Alias Node Sampler.** As mentioned before, the alias method requires two tables for a single distribution. One stores probabilities and the other is store alias-outcomes (i.e., node ids).

1) Memory cost: Given a node  $v$ , there are  $d_v$  e2e distributions, thus entailing  $d_v \times (b_f + b_i) \times d_v$  bytes memory cost. Furthermore, according to Algorithm 1, to sample the start node for a second-order random walk (i.e.,  $t = 0$ ), we need to sample it from the n2e distribution, then the alias method also incurs another two tables. Overall, the node  $v$  entails  $(d_v + 1) \times d_v \times (b_f + b_i)$  bytes memory cost.

2) Time cost: Since the sampling performance of the alias method is  $O(1)$ , the alias node sampler can generate a sample in constant time, which is  $K$ .

### 4.2 Cost Model

We summarize the above cost analysis as the cost model listed in Table 1 for second-order random walk. We can instantiate a cost model by determining the data types and the complexity of checking the existence of common neighbors. For example, assuming that the probability is stored in *float* data type, the node id is stored in *integer* data type, and the binary search is used for common neighbor checking, then  $b_f = 4$ ,  $b_i = 4$  and  $c = \log(d_v)$ .

From the table, we figure out that the more memory used the better time efficiency achieved. Concretely, we denote the memory costs of naive node sampler, rejection node sampler and alias node sampler by  $M_n$ ,  $M_r$  and  $M_a$  respectively. Similarly, the time costs are  $T_n$ ,  $T_r$  and  $T_a$ . According to Theorem 1,  $1 \leq C_v \leq d_v$  holds. Then we have the following

two orders:  $M_a > M_r > M_n$ , and  $T_a < T_r < T_n$ . Therefore, in the ideal scenario where memory is unlimited or large enough, we simply use alias node sampler to achieve the best efficiency. However, as introduced in the Introduction, this incurs the memory explosion problem in practice. To smartly utilize the memory, we turn to assign different samplers for each node, so that maximize the efficiency without causing memory explosion. Finally, we propose a memory-aware second-order random walk framework.

## 5 MEMORY-AWARE SECOND-ORDER RANDOM WALK FRAMEWORK

Figure 2 illustrates the overview of the memory-aware second-order random walk framework. The core of the framework is a cost-based optimizer. The optimizer guarantees the high-efficiency of second-order random walk on various memory budgets. It not only greedily finds an efficient node sampler assignment from scratch, but also is able to fast update the assignment when the memory budget changes online.

During the execution, the framework 1) first initializes the cost model and computes the bounding constants for the rejection node sampler. 2) Then it executes the cost-based optimizer to generate an efficient node sampler assignment without violating the memory constraint. 3) On the basis of the assignment, it initializes each node sampler across the graph. 4) Finally, the framework is ready to do the second-order random walk. In addition, the framework provides flexible programming interfaces for users to define new samplers and application-oriented random walk models. Thus, the framework can be a middleware in existing second-order random walk based applications to improve the sampling efficiency.

### 5.1 Cost-based Optimizer

The responsibility of the cost-based optimizer is to find a node sampler assignment in which the sampling efficiency is maximized while the memory footprint is constrained by the given budget. To formally analyze the node sampler assignment problem, we define it as follows:

**DEFINITION 1. (Node Sampler Assignment Problem).** Given a graph  $G = (V, E)$  and a set of node samplers  $NS = \{ns_j\}$ ,  $1 \leq j \leq S$ , the cost model is  $CM = \{(T_{ij}, M_{ij})\}$ , where  $T_{ij}$  is the time cost of node  $i$  using sampler  $ns_j$  and  $M_{ij}$  is the corresponding memory cost, the goal of the problem is to assign a sampler to each node in graph  $G$  such that the total time cost is minimized and the memory cost does not exceed a predefined memory budget  $M$ . The formal objective is

$$\text{minimize} \sum_{i=1}^{|V|} \sum_{j=1}^{j=S} T_{ij} x_{ij}, \quad (5)$$

$$\text{subject to} \sum_{i=1}^{|V|} \sum_{j=1}^{j=S} M_{ij} x_{ij} \leq M, \quad (6)$$

$$\sum_{j=1}^{j=S} x_{ij} = 1, i = 1, 2, \dots, |V|, \quad (7)$$

$$x_{ij} = \{0, 1\}, i = 1, 2, \dots, |V|; j = 1, 2, \dots, S. \quad (8)$$

The following theorem shows that the above problem is a 0-1 Multiple-Choice Knapsack Problem (MCKP).

**THEOREM 2.** *The node sampler assignment problem is a 0-1 Multiple-Choice Knapsack Problem [33].*

**PROOF.** Let  $M_{\max}$  be the maximum memory consumption among all  $M_{ij}$ , i.e.,  $M_{\max} = \max_{1 \leq i \leq |V|; 1 \leq j \leq S} \{M_{ij}\}$ .

We define  $M_{ij}^* = M_{\max} - M_{ij}$ , then Equation 6 becomes  $\sum_{i=1}^{|V|} \sum_{j=1}^{j=S} M_{ij}^* x_{ij} \geq |V| * M_{\max} - M$ .

According to the definition of 0-1 MCKP [33, 41], the assignment problem with the new memory constraint is an exact 0-1 MCKP.

In this paper, we have defined three node samplers, and they are alias node sampler, rejection node sampler, and naive node sampler. The cost model is listed in Table 1. Users can further extend the node sampler set by defining new samplers on the basis of our flexible programming interface which is introduced in Section 5.4.

### 5.2 Greedy Algorithms of Node Sampler Assignment

The original 0-1 MCKP is an NP-hard problem. It has pseudo-polynomial algorithm which is a dynamic programming solution [5], however, the dynamic programming method entails high time complexity  $O(|V|M)$  and cannot process large graphs and large memory budgets. Here we focus on greedy algorithms which are often efficient for the big data scenarios.

**5.2.1 LP Greedy Algorithm.** To solve the 0-1 MCKP, a classical solution is to transform the original problem into linear MCKP (LMCKP) by relaxing the constraint (8) and allowing  $x_{ij}$  to be real value. The optimal solution of LMCKP is the lower bound for the 0-1 MCKP. Then the original problem is solved to optimality through enumeration [6], or finds the approximation by rounding [33]. Here we use the rounding technique to achieve an efficient node sampler assignment and theoretically analyze its approximation bound.

**Properties of LMCKP.** We restate two properties of the optimal solution for LMCKP in the context of graph.

**PROPERTY 1. *P-Domination.*** For any node  $v_i \in V$ , if  $T_{ij} \geq T_{ik}$  and  $M_{ij} \geq M_{ik}$ , then  $x_{ij} = 0$  in the optimal solution.

---

**Algorithm 2** LP Greedy Algorithm

---

**Input:** Graph:  $G(V, E)$ , Cost Model:  $CM = \{(T_{ij}, M_{ij})\}$ , Memory Budget:  $M$   
**Output:** Assignment:  $assignDict$ , Trace: path

```

1: usedMem = 0
2: path = []
3: for each  $v_i$  in  $V$  do
4:   eliminate P-Domination and LP-Domination of node
   samplers of  $v_i$  according to Property 1 and 2.
5:   assign node sampler with smallest memory cost ( $NS_{i1}$ )
   to node  $v_i$ , i.e.,  $assignDict[v_i] = NS_{i1}$ 
6:   usedMem = usedMem +  $M_{i1}$ 
7: end for
8: compute the gradients  $\frac{T_{ij+1}-T_{ij}}{M_{ij+1}-M_{ij}}$  for each node.
9: sort the gradient array  $q$  in ascending order.
10: while  $q.\text{notEmpty}()$  do
11:    $NS_{ik} = (T_{ik}, M_{ik})$ , i.e., select the minimal  $\frac{T_{ij+1}-T_{ij}}{M_{ij+1}-M_{ij}}$ 
    from  $q$ .
12:    $(T_i, M_i) \leftarrow assignDict[v_i]$ 
13:   if  $usedMem - M_i + M_{ik} > M$  then
14:     break;
15:   end if
16:    $assignDict[v_i] = NS_{ik}$ 
17:   usedMem = usedMem -  $M_i + M_{ik}$ 
18:   path.append( $NS_{ik}$ ) //maintain the assignment trace for
    update.
19: end while

```

---

PROOF. Refer to the proof of Proposition 1 in the work [33].

**PROPERTY 2. LP-Domination.** For every node  $v_i \in V$ , if  $T_{ir} \geq T_{is} \geq T_{it}$ ,  $M_{ir} \leq M_{is} \leq M_{it}$ , and  $\frac{T_{is}-T_{ir}}{M_{is}-M_{ir}} > \frac{T_{it}-T_{is}}{M_{it}-M_{is}}$ , then  $x_{is} = 0$ .

PROOF. Refer to the proof of Proposition 2 in the work [33].

On the basis of the two properties, we assume the cost model  $CM = \{(T_{ij}, M_{ij})\}$  on every node are ordered as follows:  $T_{i1} \geq T_{i2} \dots \geq T_{iS}$ ,  $M_{i1} \leq M_{i2} \dots \leq M_{iS}$ , and there is no LP-Domination. For any cost pair violates the above condition, we can set the corresponding  $x$  to be zero. Fortunately, the cost model in Table 1 satisfies the condition. In general cases, a set of user-defined node samplers may not always hold the condition, i.e., there are P-Domination or LP-Domination samplers. Notice that the samplers which are not dominated are the lower convex boundary among all the user-defined samplers [26], therefore, for each node, we can find these undominated samplers by ordering them in increasing memory costs and successively testing the sampler according to Property 1 and 2. Next we give the theorem about the optimality of LMCKP.

**THEOREM 3.** An optimal solution of LMCKP has at most two fractional variables, which are from the same node, i.e.,

two variables are  $x_{ij}$  and  $x_{il}$ . Furthermore, the variables are adjacent, i.e.,  $l = j + 1$ .

PROOF. Refer to the proof of Proposition 3 in the work [33].

**Solution.** LP greedy algorithm uses LMCKP solver to get the optimal solution of LMCKP, then it rounds up the solution, i.e., assume  $x_{ij}$  and  $x_{ij+1}$  are two fractional variables, it sets  $x_{ij} = 1$  and  $x_{ij+1} = 0$  for approximation. Algorithm 2 lists the main procedure. First for each node, it eliminates the P-Domination and LP-Domination in the input cost model, and assigns the node the most time expensive node sampler, e.g.,  $NS_{i1}$ . Then it selects the minimal  $\frac{T_{ij+1}-T_{ij}}{M_{ij+1}-M_{ij}}$  to update the assignment, which indicates the most profitable changes currently. When the memory budget runs up, the algorithm finishes and does the approximation implicitly.

More concretely, we present an example of using LP greedy algorithm on a toy undirected graph, which has 4 nodes and 4 edges. The node sampler set consists of naive, rejection and alias node samplers and the memory budget is 188 Bytes. Figure 5 visualizes the key intermediate statistics of the example including the corresponding cost model and the sorted gradients for each node. First, when LP greedy algorithm finishes the initialization (Lines 3-7 in Algorithm 2), each node is assigned naive node sampler and the *usedMem* is 12. Then it selects the minimal gradient  $-0.114$ , changes node sampler of node 3 from naive to rejection, and the *usedMem* becomes  $12 - 3 + 24 = 33$ . This greedy procedure (Lines 10-19) does not stop until the memory budget is reached. In this example, when node 2 is determined to use alias method by the algorithm, the *usedMem* reaches 144 and the remained memory budget  $188 - 144 = 44$  cannot support any further node sampler update. Therefore, the final node sampler assignment is that nodes 0 and 1 use rejection node sampler and nodes 2 and 3 use alias sampler.

**Analysis of approximation bounds.** LP greedy algorithm does not guarantee to find the optimal solution of MCKP. But the following theorem shows that the LP greedy algorithm cannot be arbitrary worse with regard to the second-order random walk model.

**THEOREM 4.** Given a graph  $G$ , and the three node samplers are used, if the optimal solution of MCKP is denoted by  $OPT$ , the solution of LP greedy algorithm is  $A$ , then  $OPT \leq A \leq \max\{\frac{c+1}{c}, c\}d_{\max}OPT$ , where  $d_{\max}$  is the maximum degree of  $G$  and  $c$  is the cost of checking common neighbors of an edge which is degree dependent.

PROOF. First, it is obviously  $OPT \leq A$  because of the approximation of  $A$ .

Second, let  $OPT^{lp}$  be the optimal solution of LMCKP, since  $OPT^{lp} \leq OPT$ , we prove  $A \leq \max\{\frac{c+1}{c}, c\}d_{\max}OPT^{lp}$  instead of  $A \leq \max\{\frac{c+1}{c}, c\}d_{\max}OPT$ .

Cost Model Table									
vid	$d_v$	$C_v$	$NS_1$ (Naïve)	$NS_2$ (Rejection)	$NS_3$ (Alias)	$M_{11}$	$T_{11}$	$M_{12}$	$T_{12}$
			$M_{11}$	$T_{11}$	$M_{12}$	$T_{12}$	$M_{13}$	$T_{13}$	
0	3	2.41	3.0	6.0	36.0	2.41	96.0	1.0	
1	1	1.00	3.0	2.0	12.0	1.00	16.0	1.0	
2	2	1.60	3.0	4.0	24.0	1.60	48.0	1.0	
3	2	1.60	3.0	4.0	24.0	1.60	48.0	1.0	

Seq.	vid	Grad.	Sampler	Mem.	Seq.	vid	Grad.	Sampler	Mem.
0	3	-0.114	N->R	33	4	3	-0.025	R->A	120
1	2	-0.114	N->R	54	5	2	-0.025	R->A	144
2	1	-0.111	N->R	63	6	0	-0.024	R->A	204
3	0	-0.109	N->R	96	7	1	0	R->A	208

Sorted gradients and node sampler update sequence. N=Naïve, R=Rejection, A=Alias.

**Figure 5: A concrete example of applying LP greedy algorithm on a toy graph. The memory budget is 188, and the final node sampler assignment is {0:R, 1:R, 2:A, 3:A}. The top table describes the cost model while the bottom table lists the sorted gradients where the rows in red color are the update log by LP greedy algorithm within the memory budget. Notice that four nodes are initialized with naive node samplers, the total used memory is  $4 \times 3 = 12$  at the beginning.**

Assume the node  $v_k$  has two fractional variables  $x_{kj} = \lambda$  and  $x_{kj+1} = 1 - \lambda$  in  $OPT^{lp}$ , then  $OPT^{lp} = \sum_{i=1, i \neq k}^{|V|} \sum_{l=1}^S T_{il} + \lambda T_{kj} + (1 - \lambda)T_{kj+1}$ , and  $A = \sum_{i=1, i \neq k}^{|V|} \sum_{l=1}^S T_{il} + T_{kj}$ .

Let  $D = \sum_{i=1, i \neq k}^{|V|} \sum_{l=1}^S T_{il}$ , and  $T_{kj} \geq T_{kj+1}$ , we have

$$\begin{aligned} \frac{A}{OPT^{lp}} &= \frac{D + T_{kj}}{D + \lambda T_{kj} + (1 - \lambda)T_{kj+1}} \leq \frac{D + T_{kj}}{D + T_{kj+1}} \leq \frac{T_{kj}}{T_{kj+1}} \\ &\leq \max\left\{\frac{T_n}{T_r}, \frac{T_r}{T_a}\right\} = \max\left\{\frac{d_{v_k}(c+1)}{C_{v_k}c}, C_{v_k}c\right\}, \end{aligned}$$

where  $T_n, T_r, T_a$  are the time cost of naive, rejection and alias node samplers respectively.

Finally, based on the Theorem 1 and  $C_v \geq 1$ , it is easy to figure out  $\max\left\{\frac{d_{v_k}(c+1)}{C_{v_k}c}, C_{v_k}c\right\} \leq \max\left\{\frac{c+1}{c}, c\right\}d_{max}$ . Therefore, the above theorem holds.

The above theorem gives the upper bound  $\max\left\{\frac{c+1}{c}, c\right\}d_{max}$  of approximation between LP greedy algorithm and the optimal one. The upper bound is not only related to the graph, but also influenced by the algorithm for common neighbor checking. When binary search is used for checking,  $c$  is  $\log(d_v)$ , the upper bound is  $d_{max}\log(d_{max})$  ( $d_{max} \geq 4$ ); when hash-set is used for checking,  $c$  is one, the upper bound becomes  $2d_{max}$ . Although the upper bound is loose and graph-related, in practice, our empirical studies in Section 6 showcase the high-efficiency of the assignment generated by the LP greedy algorithm for real-world graphs. Another intuitive explanation is that existing empirical studies [33] demonstrated that the search of P-domination in the class where fractional variables exist would lower the cost, while the search among other classes will not lower the cost. For the second-order random walk with the aforementioned three node samplers,

there is actually no P-domination in the same class. Therefore, the output of the LP greedy algorithm is (close to) the optimal solution in high-probability.

**5.2.2 Degree-based Greedy Algorithm.** An intuition for solving the node sampler assignment problem is to use the alias node sampler as many as possible. From the cost model, we have two following observations: 1) The node with larger degree usually achieves more efficiency improvements when changing node sampler from rejection/naïve to alias. This motivates us to assign nodes with large degree the alias node sampler first. 2) The node with smaller degree occupies less memory resulting more nodes can use alias node sampler within a memory budget. This motivates us to assign nodes with small degree the alias node sampler first.

On the basis of the above observations, we introduce a basic degree-based greedy algorithm, which works as follows: first, the nodes in the graph are sorted by degree in increasing (decreasing) order; then for each node, within the memory budget, it tries to assign node samplers in alias, rejection and naïve order.

Compared with the LP greedy algorithm, the degree-based solution is much simpler. However, it does not consider the time cost explicitly and is hard to obtain an approximation bound. The experimental results will demonstrate that the degree-based algorithms only work well when the memory budget is large.

### 5.3 Adaptive Solution for Dynamic Memory Budget

In the real-world environment, e.g., cloud service, public clusters, the available memory is dynamic. This requires that the node sampler assignment can be adaptively adjusted when the memory budget changes. Due to the linear property of greedy algorithms, we can easily extend the proposed solution to be adaptive.

During the original greedy process, we maintain the trace of greedy options (Line 18 in Algorithm 2). When the memory budget changes, we execute the following two strategies to update the node sampler assignment.

**Memory budget increase.** When the available memory increases, the previous assignment is not affected by the new total budget, so we just restart the greedy algorithm from the last state in the trace.

**Memory budget decrease.** In this case, we need to revoke some node assignments to reduce the total memory. Since the original greedy algorithm does the greedy choice in increasing order in terms of the memory size, we execute the reverse order to reduce the used memory. Specially, we pop the greedy choices from the trace until the total memory satisfying the new budget.

```

1  class NodeSampler {
2      virtual int sample(int nid) = 0;
3      virtual float timeCost(int nid) = 0;
4      virtual float memCost(int nid) = 0;
5  };
6  class SecondRandomWalker {
7      virtual float biasedWeight(int u, int v, int z) = 0;
8 };

```

**Figure 6: Programming interfaces of the memory-aware second-order random walk framework.**

## 5.4 Framework Implementation

We carefully implemented the memory-aware framework in C++. The graph is organized as CSR format [29]. Both the cost-based optimizer and the random walk model run in a vertex-centric manner. We parallelize the computation at node level by using OpenMP library.

**Programming interface.** To enable users to easily benefit from the cost-based optimizer, we develop two programming interfaces for the framework – NODESAMPLER interface and SECONDRANDOMWALKER interface, which are illustrated in Figure 6. The NODESAMPLER defines the sampling procedure from a discrete distribution with corresponding time and memory cost. The SECONDRANDOMWALKER defines the logic of computing biased edge weight for an interested second-order random walk.

**Optimization of the cost-based optimizer.** During the initialization phase, the cost-based optimizer needs bounding constant  $C_v$  to compute the time cost of rejection method. Since the exact computation of  $C_v$  is expensive and the optimizer does not require the exact time cost, we use the bounding constant estimation (Section 3.3) to obtain approximate  $C_v$  for improving the efficiency of initialization.

## 6 EXPERIMENTS

### 6.1 Experimental Settings

Without clarification, the experiments are run on a cloud server, which equips with a 64-bit 24-core CPU, 96GB memory and 2TB hard disk. The operating system is Ubuntu 18.04 64bits. The default parallelism of the framework is set to 16.

**Datasets.** We use six public datasets: Blogcatalog, Flickr, Youtube, LiveJournal, Twitter, UK200705. All the graphs are processed into undirected. Table 2 lists the statistics of datasets. The memory size  $M_g$  of each graph is collected during the runtime from `/proc/<pid>/statm` file.

**Compared methods.** The proposed memory-aware second-order random walk framework has four variants with different greedy algorithms and optimizations. *LP-std* is the solution with LP greedy algorithm. *LP-est* is the solution using the optimization of bounding constant estimation, and the default threshold is set to 600. *Deg-inc* and *Deg-dec* are two variants using degree-based greedy algorithm in increase

G	V	E	$d_{avg}$	$M_g$
Blogcatalog	10.3K	668K	64.8	13MB
Flickr	80.5K	11.8M	146.6	185MB
Youtube	1.1M	6.0M	5.3	108MB
LiveJournal	4.8M	86.2M	17.8	1,375MB
Twitter	41.6M	2.4B	39.1	~10GB
UK200705	105.9M	6.6B	62.6	~26GB

**Table 2: Dataset statistics.**  $M_g$  is the memory size of the graph.

and decrease order respectively. In addition, the memory-unaware second-order random walk with three basic node samplers are compared, and they are called naive, rejection, and alias for simplicity.

**Second-order random walk models.** Node2vec and autoregressive models are used.

- (1) The node2vec model is controlled by two hyperparameters  $a$  and  $b$ , denoted by  $NV(a, b)$ . We follow the original work [9] and set  $a, b \in [0.25, 1, 4]$ .
- (2) The autoregressive model is controlled by one hyperparameter  $\alpha$ , denoted by  $Auto(\alpha)$ . We set the values of  $\alpha$  to be  $[0, 0.2, 0.4, 0.6, 0.8]$ .

**Benchmarks.** To evaluate the efficiency of the proposed solution, we choose two different benchmarks.

- (1) Node2vec Random Walk: Following the random walk sampling pattern in node2vec, every node in a graph samples a set of random walks with a fixed length. Here we use the same parameter settings from node2vec model [9], and sample 10 walks per node with walk length of 80.
- (2) Second-order PageRank query: Wu et al. [38] propose the second-order PageRank query. Given a query node  $v$ , it runs the second-order random walk with restart to estimate the pagerank. The decay factor is 0.85, the maximum length is 20, and the total sample size is  $4|V|$ . In addition, we randomly choose 100 query nodes for each dataset.

The node2vec random walk task is run over the node2vec model, while the second-order PageRank query is run over the autoregressive model.

**Evaluation Metrics.** We compare the elapsed time (seconds) and memory footprint among different methods. The elapsed time is further decomposed into initialization time  $T_{init}$  and sampling time  $T_s$ . For the memory-aware second-order random walk framework, the  $T_{init}$  includes the cost of computing the bounding constants  $T_{C_v}$  and the cost of initializing node sampler related data structure  $T_{NS}$ . In other words, the initialization time is

$$T_{init} = \begin{cases} T_{C_v} + T_{NS} & LP - std, LP - est, \\ T_{NS} & otherwise. \end{cases} \quad (9)$$

For the sampling cost  $T_s$ , it is the average query time per node in autoregressive model, while it is the time cost of sampling 10 walks per node with walk length of 80 in node2vec. Each experiment is run five times, and the average results are reported.

## 6.2 Effectiveness of Greedy Algorithms

We show the efficiency of node sampler assignments generated by different greedy algorithms on various memory budgets. Four datasets are used. For Blogcatalog, Flickr, Youtube, we set the maximum memory budget to 3GB, 70GB, 25GB respectively, that ensures the alias method can run in memory. For LiveJournal, the ideal maximum memory budget ( $\approx$ 109GB) exceeds the physical total memory (96GB), so we set the maximum budget to 90GB. Then we vary memory budgets by setting the ratios of the given memory budget to the maximum budget to [0.1, 0.3, 0.5, 0.7, 0.9, 1.0].

We only report the results of four models – NV(4, 0.25), NV(0.25, 4), Auto(0.2), Auto(0.8). One reason is that the selected models are already representative since they have different ranges of the bounding constants (see Figure 4). Another reason is that other configurations have similar results, and the result space of all configurations is huge. In addition, Figure 7 only visualizes the sampling cost and initialization cost on Youtube and LiveJournal because of the limited space. Actually, the results on the other two datasets are consistent with the following analysis.

**Sampling cost  $T_s$  analysis.** In Figure 7 (a)-(d) and (i)-(l), we see that all the greedy algorithms achieve better efficiency by increasing the memory budget. The LP-std and LP-est always outperform the Deg-inc and Deg-dec. Especially, when the ratio is low, the gap of  $T_s$  can be up to 46×. Taking Figure 7(a) as an example, when running NV(0.25,4) on Youtube with 2.5 GB memory budget (i.e., ratio=0.1), the LP-std takes 48.115 seconds while Deg-inc needs 2211.71 seconds. This is because the degree is not linearly proportional to the cost of sampling, so the degree-based algorithms cannot reduce the cost quickly simply according to the degree order. The LP greedy algorithm models the cost saving via gradients, and it can improve the performance effectively even when the memory budget is low. On the other hand, when the memory budget is large enough (e.g., ratio=1.0), all the methods achieve similar efficiency. In addition, LP-std and LP-est have similar efficiency across different settings, this demonstrates that bounding constant estimation works well in practice.

**Initialization cost  $T_{init}$  analysis.** In Figure 7 (e)-(h) and (m)-(p), unlike the sampling cost  $T_s$ , the initialization cost  $T_{init}$  (Equation 9) gradually increases when the memory budget increases. This is because more alias node samplers are used, and they entail  $O(d^2)$  initialization cost for each node. Another observation is that the initialization of LP-std

and LP-est is slower than the one of Deg-inc and Deg-dec. One reason is that LP-std and LP-est need the bounding constants, whose computation entails  $T_{C_v}$  (yellow bar in Figure 7). Therefore, we decompose  $T_{init}$  of LP-std and LP-est into  $T_{C_v}$  and  $T_{NS}$ . We see that when the memory budgets are large, the  $T_{NS}$  of LP-std and LP-est is comparable to the one of Deg-inc and Deg-dec. When the memory budgets are small, the degree-based algorithms are more efficient with respect to the initialization cost. This is because the node sampler assignments generated by Deg-inc and Deg-dec are much simpler than the ones created by LP-std and LP-est.

Considering the overhead  $T_{C_v}$  of LP greedy algorithms, LP-est is proposed to reduce the initialization cost. Table 3 lists the  $T_{C_v}$  of LP-std and LP-est on four datasets. Since the computation of  $C_v$  is irrelevant to the memory budget, we showcase the average  $T_{C_v}$  of different memory budgets on the same second-order random walk model. Overall LP-est can reduce 8.76%-92.07% of  $T_{C_v}$  caused by LP-std, while it still remains the similar sampling efficiency.

## 6.3 Efficiency of the Proposed Framework

We compare our proposed framework with three memory-unaware solutions on four datasets. Since our framework is memory-aware and the larger memory budget results in better performance, we choose memory budget ratios 0.1 and 1.0 as representative for comparison. Table 5 summarizes the sampling cost  $T_s$  and initialization cost  $T_{init}$ .

**Memory footprint comparison.** We present the memory consumption of memory-unaware solutions in Table 4. Consistent with our theoretical analysis in Section 4, the naive occupies the least memory, the rejection consumes the memory comparable to the graph size, and the alias incurs the most memory, which has the memory explosion problem.

**Sampling cost  $T_s$  comparison.** From Table 5, we see that when the memory budget is large enough for the dataset, alias method achieves the best performance, and LP-std (1.0) entails the similar time cost. The slight performance gap between alias method and LP-std (1.0) is caused by the following two reasons. One is that for nodes with degree one, LP-std uses the naive method on the basis of the cost model. The other is the overhead of the memory-aware second-order random walk framework itself. However, when the dataset becomes large (e.g., LiveJournal), the alias method suffers from Out-Of-Memory (OOM) error, but LP-std (1.0) still works well. This result demonstrates the robustness of our framework.

Another interesting observation is that LP-std (0.1) achieves comparable performance as well, and it is always faster than the rejection method. This implies even when the memory budget is small, our framework is able to find a good node sampler assignment for efficient random walk generation.

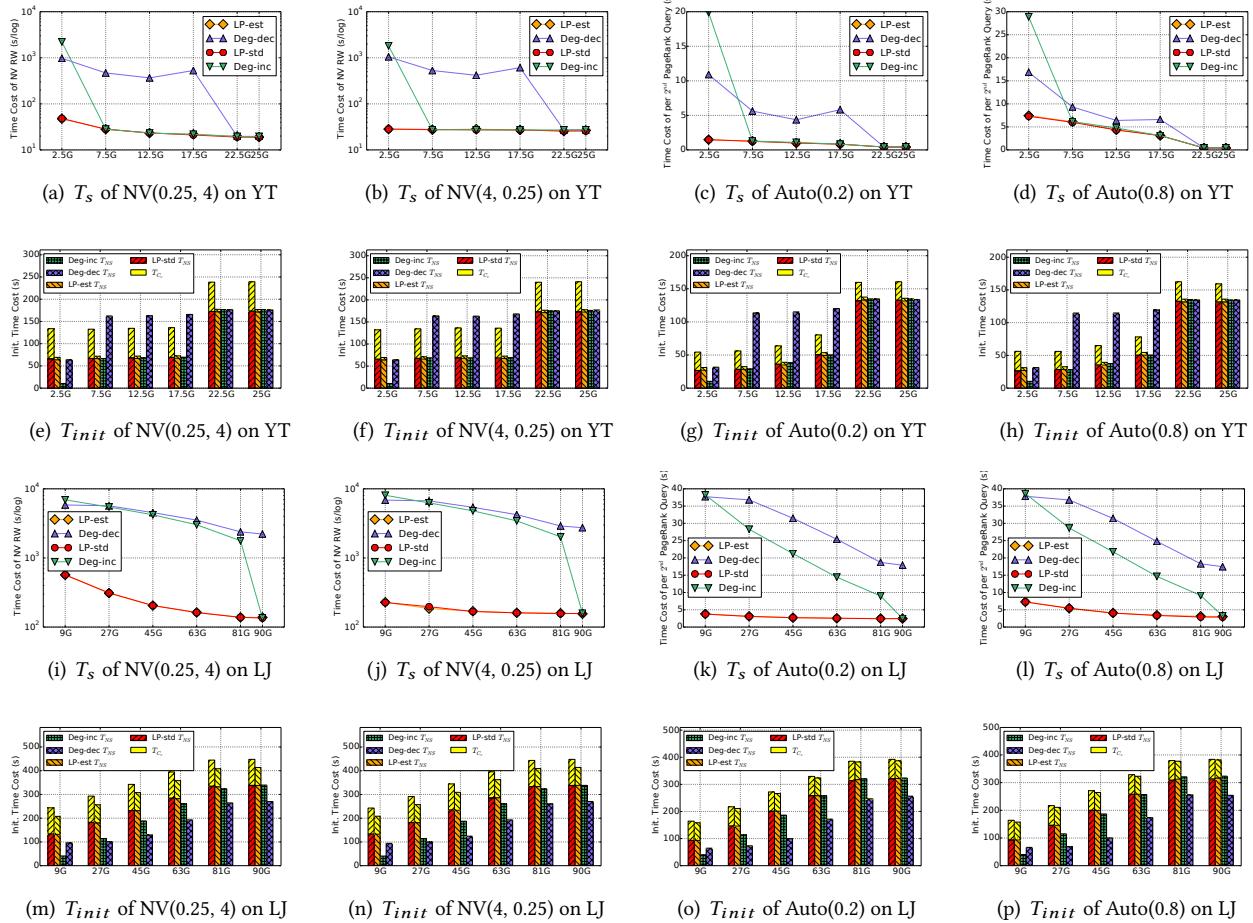


Figure 7: The comparison of  $T_s$  and  $T_{init}$  with respect to the greedy algorithms onYoutube (YT) and LiveJournal (LJ).

Dataset	NV(0.25, 4)			NV(4, 0.25)			Auto(0.2)			Auto(0.8)		
	LP-std $T_{C_v}$	LP-est $T_{C_v}$	Save cost									
Blogcatalog	2.830	1.670	40.99%	2.889	1.661	42.51%	1.867	1.470	21.26%	1.874	1.456	22.31%
Flickr	85.841	47.480	44.69%	85.981	47.477	44.78%	56.695	44.259	21.93%	56.734	44.322	21.88%
Youtube	66.453	5.333	91.97%	67.232	5.334	92.07%	28.128	4.139	85.29%	28.802	4.146	85.61%
LiveJournal	110.442	75.776	31.39%	109.949	75.697	31.15%	71.075	64.981	8.57%	71.040	64.816	8.76%

Table 3: The comparison of bounding constant computation cost  $T_{C_v}$  between LP-std and LP-est.

Graph	Naive	Rejection	Alias
Blogcatalog	0.3MB	8MB	2,848MB
Flickr	0.4MB	139MB	66,996MB
Youtube	6MB	174MB	22,949MB
LiveJournal	20MB	1,372MB	111,980MB*

Table 4: Memory footprint of memory-unaware solutions.

\* indicates the memory size is estimated.

**Initialization cost  $T_{init}$  comparison.** We compare the  $T_{init}$  of memory-unaware solutions and our proposed method. The naive method has the best initialization performance since it almost does not need to do any heavy preprocessing. The rejection method incurs the moderate initialization cost, and it only needs to compute the factor  $\min_{t \in N(v)} \frac{w_{vt}}{w'_{vt}}$  in Equation 4. The alias method entails the heaviest initialization cost among the three memory-unaware solutions.

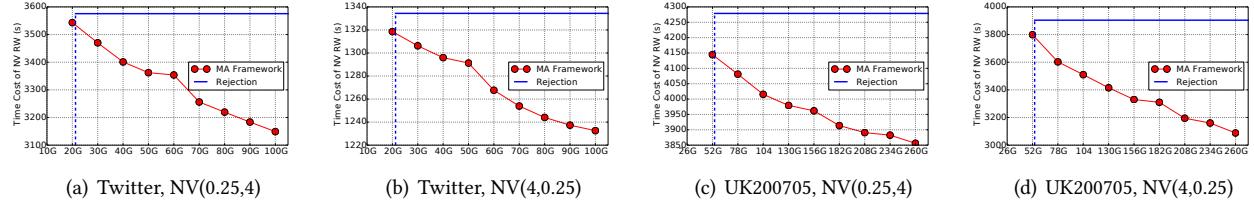
For the memory-aware solutions, the initialization cost consists of  $T_{C_v}$  and  $T_{NS}$ . When the memory budget is small, the initialization cost of our method is less than the one of alias method, since a few numbers of alias tables need to be created, but it is still greater than the rejection method because of  $T_{C_v}$ . When the memory budget is large, the initialization cost of the memory-aware framework is greater than the alias method due to  $T_{C_v}$  as well.

#### 6.4 Efficiency over Billion-edge Graphs

To further demonstrate the efficiency of the memory-aware (MA) framework over very large graphs, we run node2vec random walk task on two billion-edge graphs, Twitter and UK200705. Assuming the graph size is  $M_g$ , we set memory

Model	Cost Type	Blogcatalog					Flickr				
		Naive	Rejection	Alias	LP-std (0.1)	LP-std (1.0)	Naive	Rejection	Alias	LP-std (0.1)	LP-std (1.0)
NV(0.25,4)	$T_{init}$ $T_s$	<b>0</b> 40.894	<u>2.682</u> 1.247	9.451 <b>0.169</b>	6.020 0.881	12.048 0.173	<b>0</b> 723.564	<u>83.920</u> 22.230	229.846 <b>2.124</b>	181.263 15.052	315.415 2.205
NV(4,0.25)	$T_{init}$ $T_s$	<b>0</b> 28.462	<u>2.692</u> 0.241	9.329 <b>0.174</b>	6.151 0.204	12.176 <u>0.183</u>	<b>0</b> 588.911	<u>83.855</u> 4.121	229.846 <b>2.258</b>	182.058 3.401	315.709 2.324
Auto(0.2)	$T_{init}$ $T_s$	<b>0</b> 0.522	<u>1.852</u> 0.015	9.115 <b>0.004</b>	4.119 0.014	10.834 <b>0.004</b>	<b>0</b> 5.495	<u>55.433</u> 0.115	220.835 <b>0.040</b>	123.821 0.099	276.978 <u>0.047</u>
Auto(0.8)	$T_{init}$ $T_s$	<b>0</b> 0.757	<u>1.807</u> 0.050	9.207 <b>0.004</b>	4.156 0.047	11.058 <u>0.005</u>	<b>0</b> 6.378	<u>55.662</u> 0.379	220.735 <b>0.044</b>	123.596 0.312	277.582 <u>0.050</u>
Model	Cost Type	Youtube					LiveJournal				
		Naive	Rejection	Alias	LP-std (0.1)	LP-std (1.0)	Naive	Rejection	Alias	LP-std (0.1)	LP-std (1.0)
NV(0.25, 4)	$T_{init}$ $T_s$	<b>0.003</b> 2558.18	<u>64.357</u> 127.374	172.012 <b>18.188</b>	135.031 48.115	240.450 <u>18.980</u>	<b>0.013</b> 7472.88	<u>108.741</u> 803.293	OOM OOM	251.328 <u>569.428</u>	455.423 <b>135.041</b>
NV(4, 0.25)	$T_{init}$ $T_s$	<b>0.003</b> 2253.84	<u>66.410</u> 43.114	172.746 <b>24.403</b>	133.317 28.281	242.238 25.793	<b>0.013</b> 8815.27	<u>109.026</u> 289.735	OOM OOM	250.705 <u>226.582</u>	455.831 <b>157.496</b>
Auto(0.2)	$T_{init}$ $T_s$	<b>0.003</b> 24.202	<u>28.210</u> 1.780	132.701 <b>0.417</b>	55.747 1.486	161.893 <u>0.427</u>	<b>0.013</b> 44.077	<u>69.845</u> 4.411	OOM OOM	172.219 <u>3.743</u>	400.057 <b>2.448</b>
Auto(0.8)	$T_{init}$ $T_s$	<b>0.003</b> 33.631	<u>28.244</u> 8.282	132.422 <b>0.410</b>	57.520 7.318	160.562 <u>0.449</u>	<b>0.013</b> 44.118	<u>69.873</u> 8.828	OOM OOM	171.973 <u>7.265</u>	391.956 <b>2.877</b>

**Table 5: Efficiency comparison among memory-unaware approaches and the memory-aware framework. The bold style indicates the best performance and the underline style is the second best performance.**



**Figure 8: Sampling efficiency of the memory-aware framework on Twitter and UK200705.**

budgets by varying from  $M_g$  to  $10M_g$ . Table 2 gives the  $M_g$  of Twitter and UK200705. The experiments are conducted on a server equipping with 376GB memory. Among the three memory-unaware solutions, except the rejection method, naive method cannot finish the task in 4 hours, and alias method fails because of the Out-Of-Memory error. Therefore, we only compare the MA framework with the rejection method, which requires the minimal memory around 21G and 54G on Twitter and UK200705 respectively.

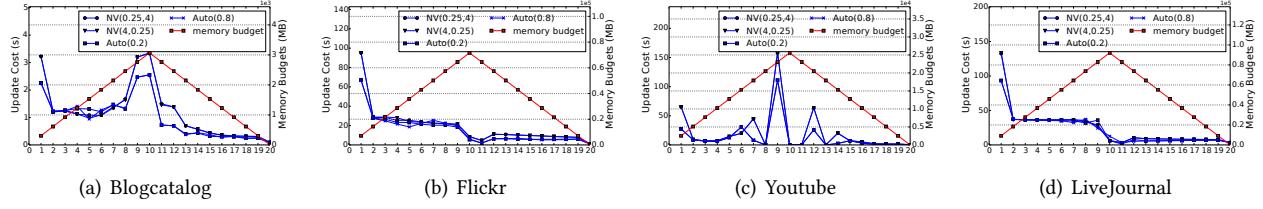
Figure 8 illustrates the experimental results. Notice that the performance of the MA framework with memory budget  $M_g$  is not listed because it cannot finish the execution in 4 hours. First, the MA framework can process the billion-edge graphs efficiently. On both Twitter and UK200705, it outperforms the rejection method even when the memory budget is smaller than the minimal memory requirement of the rejection method. This is because the framework assigns some nodes with small degree the naive method, thus saving memory for other nodes to use the alias method and improving the overall efficiency. Second, with the increase of memory budgets, the efficiency of the framework improves gradually. Taking NV(4, 0.25) on UK200705 as an example, the improvement is about 19%, i.e., from 3799.02s to 3088.17s, when memory budget increases from  $2M_g$  to  $10M_g$ . Compared with the results on previous median-size graphs, the

performance improvement of MA framework on billion-edge graphs is not that significant. The reason is that even when the memory budget is  $10M_g$ , it is still no more than 0.06% of the memory required by alias method (e.g., Twitter and UK200705 need total 1796TB and 379TB for alias method respectively), and the framework cannot allocate many alias node samplers, especially for the nodes with large degree.

## 6.5 Evaluation with Dynamic Memory Budgets

Here we evaluate the time cost of node sampler assignment update with dynamic memory budgets. For each dataset, we synthetically generate a memory budget update trace. Specifically, given a maximal memory budget  $M_{max}$ , first the memory budget linearly increases with the step  $\frac{M_{max}}{10}$ . Once  $M_{max}$  is reached, the memory budget decreases linearly with the same step. The red line in Figure 9 illustrates the trace of dynamic budget. The blue lines visualize the corresponding assignment update cost under the different second-order random walk models. Notice that  $T_{C_v}$  is excluded since it is only computed once during the memory budget update.

From the figures, except the first point which initializes the node sampler assignment from scratch, we see that the assignment update cost is small. Taking LiveJournal on Auto(0.2)



**Figure 9: The update cost of different models with dynamic memory budgets. x-axis is the timestamp of changing memory budget.**

as an example, during the increase of memory budget, the update cost is around 35 seconds, while at the beginning it needs about 93 seconds. When the memory budget decreases, it only needs about 7 seconds. According to Section 5.3, the logic of memory budget decrease is much simpler than the one of memory budget increase.

In addition, for Blogcatalog and YouTube, when memory budget increases, we find that there are burst update costs. For NV(0.25, 4) on YouTube, when memory budget increases from 20480MB to 23040MB, it takes about 158.047 seconds for the update. After profiling the data, this is because the framework chooses the node with the maximal degree (28,754) to use alias method, which entails the heavy initialization cost.

## 7 RELATED WORK

### 7.1 Random Walk on Graphs

Most existing works about random walk focus on designing new models to improve the accuracy of the applications. For example, Li et al. [16] introduced rejection-controlled Metropolis hastings algorithm and generalized maximum-degree random walk algorithm to improve the accuracy of graph property estimation. Sengupta et al. [31] uses random walk to estimate reachability of nodes in a large graph. Besides the first-order random walk, the second-order random walk is widely applied to model higher-order dependencies in various domains, including user trails on the Web [40], global shipping traffic [40], online social networks [7] and e-mail communications [23]. Boldi et al. [2] proposed a triangular random model to unveil arc-community structure in the social network, which assigns different weights for the triangular successors and the non-triangular successors. Wu et al. [38] studied the effectiveness of various second-order random walk models on clickstream data.

With regard to the efficiency of random walk-based applications, a general approach is to reduce the number of samples to reach the same accurate estimation. Nazi et al. [24] introduced a walk-estimate framework, which starts with a much shorter random walk, and then proactively estimates the sampling probability to fast approximate the target distribution. Another approach is to reduce the time cost of a sample generation. Zhou et al. [44] introduced a distributed second-order random walk algorithm on a Pregel-like graph

computation framework [21] to support large-scale node2vec. Our work also concentrates on efficiency optimization, but it emphasizes the performance of generating random walks in a single server. For the distributed solutions [44] that are built on Pregel-like graph computation frameworks, our proposed memory-aware framework can be applied to help improve the sampling efficiency for each worker. In addition, our framework can function as a middleware for other applications which use second-order random walk as a building block.

### 7.2 Cost-based Optimization

Cost-based optimization is a classical database technique for query optimization [4]. It looks at all of the possible ways in which a query can be executed and each way is assigned a “cost”, which indicates how efficiently that query can be run. Then, we pick the execution plan that has the least cost. It has been extensively used in the relational database, NoSQL, and various big data processing systems [12, 20, 34].

With regard to the graph systems, cost-based optimization is also used to improve the efficiency of graph query [11, 43]. In addition, distributed pattern matching uses cost-based optimization to speed up the external storage access [13]. The memory-aware second-order random walk framework uses the idea of cost-based optimization to find an efficient node sampler assignment so that it can generate random walks fast.

## 8 CONCLUSION

Second-order random walk becomes an important tool for modeling higher-order dependencies in data. We studied the problem of how to efficiently support the second-order random walk with various memory budgets in this work. We proposed a memory-aware framework by developing a cost-based optimizer. The optimizer assigns different node samplers for each node in the graph and ensures the efficiency is maximized on a memory budget. We designed an effective greedy algorithm to generate such high-efficiency assignment. Finally, we empirically evaluated the framework on four real-world large graphs, and the results clearly demonstrated the advantages of our memory-aware framework. In addition, the complex relationships between the efficiency

and graph structural properties in the context of second-order random walk is still an open question, and we treat it as our future work.

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