

ASAP: Towards Accurate, Stable and Accelerative Penetrating-Rank Estimation on Large Graphs

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Abstract. Pervasive web applications increasingly require a measure of similarity among objects. Penetrating-Rank (P-Rank) has been one of the promising link-based similarity metrics as it provides a comprehensive way of jointly encoding both incoming and outgoing links into computation for emerging applications. In this paper, we investigate P-Rank efficiency problem that encompasses its accuracy, stability and computational time. (1) We provide an accuracy estimate for iteratively computing P-Rank. A symmetric problem is to find the iteration number K needed for achieving a given accuracy ϵ . (2) We also analyze the stability of P-Rank, by showing that small choices of the damping factors would make P-Rank more stable and well-conditioned. (3) For undirected graphs, we also explicitly characterize the P-Rank solution in terms of matrices. This results in a novel non-iterative algorithm, termed ASAP, for efficiently computing P-Rank, which improves the CPU time from $O(n^4)$ to $O(n^3)$. Using real and synthetic data, we empirically verify the effectiveness and efficiency of our approaches.

1 Introduction

The study of quantifying structural similarity between vertices in ubiquitous networks has attracted considerable attention over the past decade. Typical structural similarity metrics include Google PageRank, Hyperlink-Induced Topic Search (HITS), Co-citation, Bibliographic Coupling, SimRank and SimFusion (*e.g.*, [1,2,3,4,5,6]).

P-Rank (Penetrating-Rank) is a new similarity measure of this kind, which was initially proposed by Zhao *et al.* [7]. The similarity scores flowing from in-link neighbors of entities are penetrated through their out-link neighbors. Emerging real-world applications of P-Rank include biological networks, collaborative filtering, graph classification, web document ranking, and outlier detection (*e.g.*, [8,9,4,10,5,6]).

In contrast to other similarity measures, P-Rank has become one of the important metrics owing to the following two reasons. (i) P-Rank provides a comprehensive way to jointly explore both in- and out-link relationships with semantic completeness. In comparison, other similarity measures, say SimRank, have the “limited information problem” [5], in which only in-link relationships can be partially exploited. (ii) P-Rank transcends other similarity measures in its most general form. Other measures such as SimRank [5], Amsler [1] can be regarded as just special cases of P-Rank [7].

Unfortunately, previous work on P-Rank suffers from the following limitations. (i) P-Rank solution is known to converge [7], but a precise accuracy estimation of P-Rank is not given. (ii) No prior work has well studied *the P-Rank condition number*, which indeed has played a paramount role in measuring how much the web graph can change in proportion to small changes in the P-Rank scoring results. (iii) The P-Rank time complexity in [7] retains *quartic* in the worst case for both digraphs and undirected graphs. To the best of our knowledge, there is no efficient P-Rank algorithm specially designed for *undirected graphs*. These practical needs call for an in-depth investigation of P-Rank and its efficiency.

Contributions. This paper studies the P-Rank problems regarding its *accuracy*, *stability* and *computational efficiency*. Our main results are summarized below.

1. We provide an accuracy estimation for P-Rank iteration (Section 3). We find that the number of iterations $K = \lceil \log \epsilon / \log (\lambda \cdot C_{\text{in}} + (1 - \lambda) \cdot C_{\text{out}}) \rceil$ suffices to acquire a desired accuracy of ϵ , where λ is a weighting factor, and C_{in} and C_{out} are in- and out-link damping factors, respectively.
2. We introduce the notion of P-Rank *condition number* κ_{∞} to investigate the stability issue of P-Rank (Section 4). We show that P-Rank is *well-conditioned* for small choices of the damping factors, by providing a tight *stability bound* for κ_{∞} .
3. We propose a novel *non-iterative* $O(n^3)$ -time algorithm (**ASAP**) for efficiently computing similarities over undirected graphs (Section 5). We explicitly couch the P-Rank solution *w.r.t.* matrix products in connection with its eigen-problem.
4. We experimentally verify the efficiency of our methods on real and synthetic data (Section 6). We find that P-Rank exponentially converges *w.r.t.* the iteration number. The stability of P-Rank is determined by the damping factors and weighting factors. The **ASAP** algorithm outperforms baseline algorithms on undirected graphs.

2 Preliminaries

In this section, we first give a brief overview of the P-Rank model. We then present notations and formulations of the P-Rank similarity.

2.1 An Overview of P-Rank

The basic essence of the P-Rank similarity model is distilled from the standard concept of SimRank metric [5]. Concretely, the theme of P-Rank involves three facets below:

1. Two distinct objects are similar if they are referenced by similar objects. (in-link recursion)
2. Two distinct objects are similar if they reference similar objects. (out-link recursion)
3. Every object is maximally similar to itself. (a base case)

Based on these facets, P-Rank scores, as described in its name, flowing from the incoming neighbors of objects are able to penetrate the outgoing neighbors. The concise elegance of the P-Rank intuition makes it one of the useful and cutting-edge structural metrics in the link-based analysis of information networks.

2.2 Notations

symbol	designation	symbol	designation
\mathcal{G}	information network	$s(a, b)$	P-Rank score between vertices a and b
$\mathcal{I}(a)$	in-neighbors of vertex a	C_{in} / C_{out}	in-link / out-link damping factor
$\mathcal{O}(a)$	out-neighbors of vertex a	λ	weighting factor
n	number of vertices in \mathcal{G}	\mathbf{A}	adjacency matrix of \mathcal{G}
m	number of edges in \mathcal{G}	\mathbf{S}	P-Rank similarity matrix of \mathcal{G}
K	number of iterations	\mathbf{I}	identity matrix

2.3 Formulation of P-Rank Model

We first define the network graph, and then introduce the P-Rank formulation.

Network Graph. A *network* is a labeled graph $\mathcal{G} = (\mathcal{V}, \mathcal{E}, \mathcal{A}; l)$, in which (i) \mathcal{V} is a finite set of vertices. A partition of \mathcal{V} is formed by dividing \mathcal{V} into N disjointed domain-specific parts \mathcal{V}_i ($i = 1, \dots, N$) s.t. $\mathcal{V} = \bigcup_{i=1}^N \mathcal{V}_i$ and $\mathcal{V}_i \cap \mathcal{V}_j = \emptyset$ if $i \neq j$; (ii) $\mathcal{E} \subseteq \mathcal{V} \times \mathcal{V}$ is an edge set, and each vertex-pair $(u, v) \in \mathcal{E}$ denotes an edge from vertex u to v ; (iii) \mathcal{A} is a finite alphabet set, with a labeling function $l : \mathcal{V} \rightarrow \mathcal{A}$.

Based on the network graph, P-Rank model can be formulated as follows:

P-Rank Similarity. Given a network \mathcal{G} , for any two distinct vertices $u, v \in \mathcal{V}$, we define a scoring function $s(u, v) \in [0, 1]$ as follows:

$$\begin{aligned}
 & s(u, u) = 1; \tag{1} \\
 & s(u, v) = \underbrace{\frac{\lambda \cdot C_{in}}{|\mathcal{I}(u)| |\mathcal{I}(v)|} \sum_{i=1}^{|\mathcal{I}(u)|} \sum_{j=1}^{|\mathcal{I}(v)|} s(\mathcal{I}_i(u), \mathcal{I}_j(v))}_{\text{in-link part}} + \underbrace{\frac{(1 - \lambda) \cdot C_{out}}{|\mathcal{O}(u)| |\mathcal{O}(v)|} \sum_{i=1}^{|\mathcal{O}(u)|} \sum_{j=1}^{|\mathcal{O}(v)|} s(\mathcal{O}_i(u), \mathcal{O}_j(v))}_{\text{out-link part}}
 \end{aligned}$$

where (i) $\lambda \in [0, 1]$ balances the contributions of in- and out-links, (ii) C_{in} and $C_{out} \in (0, 1)$ are the damping factors for in- and out-link directions, respectively, (iii) $\mathcal{I}_i(u)$ and $\mathcal{O}_i(u)$ are the individual members of $\mathcal{I}(u)$ and $\mathcal{O}(u)$, respectively, and (iv) $|\mathcal{I}(u)|$ and $|\mathcal{O}(u)|$ are the cardinalities of $\mathcal{I}(u)$ and $\mathcal{O}(u)$, respectively.

It is worth noting that to avoid $s(u, v) = \infty$ when $|\mathcal{I}(\cdot)|$ or $|\mathcal{O}(\cdot)| = 0$, we define: (a) the in-link part of Eq.(2) = 0 if $\mathcal{I}(u) \cup \mathcal{I}(v) = \emptyset$; (b) the out-link part of Eq.(2) = 0 if $\mathcal{O}(u) \cup \mathcal{O}(v) = \emptyset$; (c) $s(u, v) = 0$, if $(\mathcal{I}(u) \cup \mathcal{I}(v)) \cap (\mathcal{O}(u) \cup \mathcal{O}(v)) = \emptyset$.

P-Rank Iterative Paradigm. To compute the P-Rank similarity $s(u, v)$ of Eq.(2), the conventional approach is to construct an iterative paradigm as follows:

$$s^{(0)}(u, v) = \begin{cases} 0, & \text{if } u \neq v; \\ 1, & \text{if } u = v. \end{cases} \tag{3}$$

For each $k = 0, 1, 2, \dots$, the $(k + 1)$ -th iterate $s^{(k+1)}(\cdot, \cdot)$ can be computed as

$$\begin{aligned}
 s^{(k+1)}(u, u) &= 1. \\
 s^{(k+1)}(u, v) &= \frac{\lambda \cdot C_{in}}{|\mathcal{I}(u)||\mathcal{I}(v)|} \sum_{i=1}^{|\mathcal{I}(u)|} \sum_{j=1}^{|\mathcal{I}(v)|} s^{(k)}(\mathcal{I}_i(u), \mathcal{I}_j(v)) \\
 &\quad + \frac{(1-\lambda) \cdot C_{out}}{|\mathcal{O}(u)||\mathcal{O}(v)|} \sum_{i=1}^{|\mathcal{O}(u)|} \sum_{j=1}^{|\mathcal{O}(v)|} s^{(k)}(\mathcal{O}_i(u), \mathcal{O}_j(v)). \tag{4} \\
 s^{(k+1)}(u, v) &= \text{the in-link part of Eq.(4), if } \mathcal{O}(u) \cup \mathcal{O}(v) = \emptyset, \\
 s^{(k+1)}(u, v) &= \text{the out-link part of Eq.(4), if } \mathcal{I}(u) \cup \mathcal{I}(v) = \emptyset.
 \end{aligned}$$

where $s^{(k)}(u, v)$ is the k -th iterative P-Rank score between vertices u and v . It has been proved in [7] that the exact similarity $s(u, v)$ is the supremum of $\{s^{(k)}(u, v)\}_{k=0}^\infty$, denoted by $\sup_k \{s^{(k)}(u, v)\}$, i.e.,

$$\lim_{k \rightarrow \infty} s^{(k)}(u, v) = \sup_{k \geq 0} \{s^{(k)}(u, v)\} = s(u, v) \quad (\forall u, v \in \mathcal{V}). \tag{5}$$

3 Accuracy Estimate on P-Rank Iteration

Based on Eq.(4), the iterative P-Rank similarity sequence $\{s^{(k)}(\cdot, \cdot)\}_{k=0}^\infty$ was known to converge monotonically [7]. However, the gap between the k -th iterative similarity $s^{(k)}(\cdot, \cdot)$ and the exact similarity $s(\cdot, \cdot)$ still remains unknown. Practically, quantifying this gap is very important in estimating the accuracy of iteratively computing P-Rank. For instance, given a tolerated error $\epsilon > 0$, one may naturally ask ‘‘What is the number of P-Rank iterations needed for achieving such an accuracy?’’.

This motivates us to study the *P-Rank accuracy estimate problem*. Given a network \mathcal{G} , for any iteration number $k = 1, 2, \dots$, it is to find an upper bound ϵ_k of the gap between the k -th iterative similarity $s^{(k)}(\cdot, \cdot)$ and the exact $s(\cdot, \cdot)$, s.t. $|s^{(k)}(u, v) - s(u, v)| \leq \epsilon_k$ for any vertices u and v in \mathcal{G} .

The main result of this section is the following.

Theorem 1. *The P-Rank accuracy estimate problem has a tight upper bound*

$$\epsilon_k = (\lambda C_{in} + (1 - \lambda)C_{out})^{k+1}$$

such that

$$|s^{(k)}(u, v) - s(u, v)| \leq \epsilon_k. \quad (\forall k = 0, 1, \dots, \forall u, v \in \mathcal{V}) \tag{6}$$

A detailed proof of Theorem 1 can be found in [11] due to space limitations.

Theorem 1 provides an a-priori estimate for the gap between iterative and exact P-Rank similarity scores, which solely hinges on the weighing factor λ , the in- and out-link damping factors C_{in} and C_{out} , and the number k of iterations. To ensure more accurate P-Rank estimation results, it can be discerned from $\epsilon_k = (\lambda C_{in} + (1 - \lambda)C_{out})^{k+1} = (\lambda(C_{in} - C_{out}) + C_{out})^{k+1}$ that the smaller choices of C_{in} and C_{out} (i) with a smaller λ if $C_{in} > C_{out}$, or (ii) with a larger λ if $C_{in} < C_{out}$, are more preferable.

Example 1. Setting $C_{in} = 0.6, C_{out} = 0.4, \lambda = 0.3, k = 5$ will produce the following high accuracy :

$$\epsilon_k = (0.3 \times 0.6 + (1 - 0.3) \times 0.4)^{5+1} = 0.0095. \quad \square$$

Notice that the upper bound in Eq.(6) is attainable. Consider the network \mathcal{G}_0 depicted in Figure 1. It is apparent that $s^{(0)}(u, v) = 0$. For $k = 1, 2, \dots$, it can be easily obtained that $s^{(k)}(u, v) = \lambda C_{in} + (1 - \lambda)C_{out}$, which implies that $s(u, v) = \lambda C_{in} + (1 - \lambda)C_{out}$. Hence, in the case of $k = 0$, $|s(u, v) - s^{(k)}(u, v)| = (\lambda C_{in} + (1 - \lambda)C_{out})^{0+1}$ gives the precise upper bound in Eq.(6).

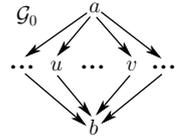


Fig. 1. “=” of Eq.(6) is attainable

Moreover, in the special case of $\lambda = 1$, Eq.(6) can be simplified to the SimRank accuracy estimate problem given in [8]. From this perspective, P-Rank accuracy estimate problem is the extension of Proposition 1 in [8] by jointly considering both in- and out-links of similarity computation.

Conversely, the exponential P-Rank convergence rate in Theorem 1 also implies that the number k of P-Rank iterations needed for attaining a desired accuracy ϵ amounts to

$$k = \lceil \log \epsilon / \log (\lambda \cdot C_{in} + (1 - \lambda) \cdot C_{out}) \rceil .$$

4 Stability Analysis of P-Rank Model

In this section, we investigate the important issues of P-Rank sensitivity and stability. We first reformulate the P-Rank formulae in matrix notations (Subsect. 4.1). Using this representation, we then give a rigorous bound of the P-Rank conditional number for its stability analysis (Subsect. 4.2).

4.1 P-Rank Matrix Representation

We start by introducing the following symbols used throughout this section.

For a network \mathcal{G} with n vertices, we denote by (i) $\mathbf{A} = (a_{i,j}) \in \mathbb{R}^{n \times n}$ the *adjacency matrix* of \mathcal{G} whose entry $a_{i,j}$ equals 1, if there exists an edge from vertex i to j ; or 0, otherwise, (ii) $\mathbf{S} = (s_{i,j}) \in \mathbb{R}^{n \times n}$ the *P-Rank similarity matrix* whose entry $s_{i,j}$ is equal to P-Rank score $s(i, j)$, and (iii) $\mathbf{Q} = (q_{i,j}) \in \mathbb{R}^{n \times n}$ and $\mathbf{P} = (p_{i,j}) \in \mathbb{R}^{n \times n}$ the *one-step backward and forward transition probability matrix* of \mathcal{G} , respectively, whose entries defined as follows:

$$q_{i,j} \triangleq \begin{cases} a_{j,i} / \sum_{j=1}^n a_{j,i}, & \text{if } \mathcal{I}(i) \neq \emptyset; \\ 0, & \text{if } \mathcal{I}(i) = \emptyset. \end{cases} \quad p_{i,j} \triangleq \begin{cases} a_{i,j} / \sum_{j=1}^n a_{i,j}, & \text{if } \mathcal{O}(i) \neq \emptyset; \\ 0, & \text{if } \mathcal{O}(i) = \emptyset. \end{cases} \tag{7}$$

With the above notations, the P-Rank formulae (1) and (2) can be rewritten as¹

$$\mathbf{S} = \lambda C_{in} \cdot \mathbf{Q} \cdot \mathbf{S} \cdot \mathbf{Q}^T + (1 - \lambda)C_{out} \cdot \mathbf{P} \cdot \mathbf{S} \cdot \mathbf{P}^T + (1 - \lambda C_{in} - (1 - \lambda)C_{out}) \cdot \mathbf{I}_n, \tag{8}$$

Also, dividing both sides of Eq.(8) by $(1 - \lambda C_{in} - (1 - \lambda)C_{out})$ results in

$$\mathbf{S}' = \lambda C_{in} \cdot \mathbf{Q} \cdot \mathbf{S}' \cdot \mathbf{Q}^T + (1 - \lambda)C_{out} \cdot \mathbf{P} \cdot \mathbf{S}' \cdot \mathbf{P}^T + \mathbf{I}_n, \text{ and} \tag{9}$$

¹ Although in this case the diagonal entries of \mathbf{S} are not equal to 1, \mathbf{S} still remains diagonally dominant, which ensures that “every vertex is maximally similar to itself”. Li *et al.* [12] has showed that this revision for SimRank matrix representation is reasonable, which can be applied in the similar way to P-Rank.

$$\mathbf{S} = (1 - \lambda C_{in} - (1 - \lambda)C_{out}) \cdot \mathbf{S}'.$$

Comparing Eq.(8) with Eq.(9), we see that the coefficient $(1 - \lambda C_{in} - (1 - \lambda)C_{out})$ of \mathbf{I}_n in Eq.(8) merely contributes an overall multiplicative factor to P-Rank similarity. Hence, setting the coefficient to 1 in Eq.(8) still preserves the *relative* magnitude of the P-Rank score though the diagonal entries of \mathbf{S} in this scenario might not equal 1^2 .

4.2 Conditional Number of P-Rank

Using the P-Rank matrix model of Eq.(9), we next theoretically analyze the stability of P-Rank, by determining its *conditional number* $\kappa_\infty(\cdot)$ in the infinity norm sense, which has important implications for the sensitivity of P-Rank computation. One complicated factor in P-Rank stability analyses is to bound the conditional number $\kappa_\infty(\cdot)$.

Before giving a formal definition of $\kappa_\infty(\cdot)$, we first introduce the following notations:

(i) We denote by $\text{vec}(\mathbf{X}) \in \mathbb{R}^{n^2}$ the *vectorization* of the matrix $\mathbf{X} \in \mathbb{R}^{n \times n}$ formed by stacking the columns of \mathbf{X} into a single column vector, and (ii) the operator \otimes the *Kronecker product* of two matrices.

Taking $\text{vec}(\cdot)$ on both sides of Eq.(9) and applying the Kronecker property of $(\mathbf{B}^T \otimes \mathbf{A})\text{vec}(\mathbf{X}) = \text{vec}(\mathbf{A}\mathbf{X}\mathbf{B})$ [13], we have

$$\text{vec}(\mathbf{S}) = \lambda C_{in}(\mathbf{Q} \otimes \mathbf{Q})\text{vec}(\mathbf{S}) + (1 - \lambda)C_{out}(\mathbf{P} \otimes \mathbf{P})\text{vec}(\mathbf{S}) + \text{vec}(\mathbf{I}_n).$$

Rearranging terms in the above equation produces

$$\underbrace{(\mathbf{I}_{n^2} - \lambda C_{in}(\mathbf{Q} \otimes \mathbf{Q}) - (1 - \lambda)C_{out}(\mathbf{P} \otimes \mathbf{P}))}_{\triangleq \mathbf{M}} \cdot \underbrace{\text{vec}(\mathbf{S})}_{\triangleq \mathbf{s}} = \underbrace{\text{vec}(\mathbf{I}_n)}_{\triangleq \mathbf{b}}. \tag{10}$$

Notice that Eq.(10) is a *linear* system of n^2 equations in terms of $s_{i,j}$ in nature as it can be reduced in the form of $\mathbf{M} \cdot \mathbf{s} = \mathbf{b}$, as illustrated in Eq.(10).

Based on Eq.(10), we now give a formal definition of *P-Rank conditional number*.

Definition 1 (P-Rank conditional number). For a network \mathcal{G} , let \mathbf{Q} and \mathbf{P} be the one-step backward and forward transition probability matrix of \mathcal{G} , respectively, defined by Eq.(7), and let

$$\mathbf{M} \triangleq \mathbf{I}_{n^2} - \lambda C_{in}(\mathbf{Q} \otimes \mathbf{Q}) - (1 - \lambda)C_{out}(\mathbf{P} \otimes \mathbf{P}). \tag{11}$$

The P-Rank conditional number of \mathcal{G} , denoted by $\kappa_\infty(\mathcal{G})$, is defined as

$$\kappa_\infty(\mathcal{G}) \triangleq \|\mathbf{M}\|_\infty \cdot \|\mathbf{M}^{-1}\|_\infty, \tag{12}$$

where $\|\cdot\|_\infty$ is the ∞ -norm that returns the maximum absolute row sum of the matrix.

The conditional number is introduced for being applied to P-Rank stability analysis.

Theorem 2. Given the network \mathcal{G} , for any weighting factor $\lambda \in [0, 1]$ and in- and out-link damping factors $C_{in}, C_{out} \in (0, 1)$, the P-Rank problem has the following tight bound of conditional number

$$\kappa_\infty(\mathcal{G}) \leq \frac{1 + \lambda \cdot C_{in} + (1 - \lambda) \cdot C_{out}}{1 - \lambda \cdot C_{in} - (1 - \lambda) \cdot C_{out}}. \tag{13}$$

² In what follows, we shall base our techniques on the P-Rank matrix form of Eq.(9).

For the interest of space, please refer to [11] for a detailed proof of Theorem 2.

Theorem 2 provides a tight upper bound of the P-Rank conditional number $\kappa_\infty(\mathcal{G})$. Intuitively, the value of $\kappa_\infty(\mathcal{G})$ measures how stable the P-Rank similarity score is to the changes either in the link structure of the network \mathcal{G} (by inserting or deleting vertices or edges), or in the damping and weighting factors of C_{in} , C_{out} and λ .

To get a feel for how $\kappa_\infty(\mathcal{G})$ affects P-Rank sensitivity, we denote by $\Delta\mathbf{A}$ the updates in \mathcal{G} to the adjacency matrix \mathbf{A} , $\Delta\mathbf{M}$ (*resp.* $\Delta\mathbf{s}$) the changes (caused by $\Delta\mathbf{A}$) to \mathbf{M} (*resp.* \mathbf{s}) defined in Eq.(10). As Eq.(10) is a linear equation, it is known that

$$\frac{\|\Delta\mathbf{s}\|_\infty}{\|\mathbf{s}\|_\infty} \leq \kappa_\infty(\mathcal{G}) \cdot \frac{\|\Delta\mathbf{M}\|_\infty}{\|\mathbf{M}\|_\infty} \leq \frac{1 + \lambda \cdot C_{in} + (1 - \lambda) \cdot C_{out}}{1 - \lambda \cdot C_{in} - (1 - \lambda) \cdot C_{out}} \cdot \frac{\|\Delta\mathbf{M}\|_\infty}{\|\mathbf{M}\|_\infty}. \quad (14)$$

This tells that the small $\kappa_\infty(\mathcal{G})$ (*i.e.*, small choices for C_{in} and C_{out}) would make P-Rank stable, implying that a small change $\Delta\mathbf{M}$ in the link structure to \mathbf{M} may not cause a large change $\Delta\mathbf{s}$ in P-Rank scores. Conversely, the large value of $\kappa_\infty(\mathcal{G})$ would make P-Rank *ill-conditioned*.

To see how the weighting factor λ affects $\kappa_\infty(\mathcal{G})$, we compute the partial derivatives of the bound for $\kappa_\infty(\mathcal{G})$ *w.r.t.* λ to get

$$\frac{\partial}{\partial \lambda} \left(\frac{1 + \lambda \cdot C_{in} + (1 - \lambda) \cdot C_{out}}{1 - \lambda \cdot C_{in} - (1 - \lambda) \cdot C_{out}} \right) = \frac{2(C_{in} - C_{out})}{(1 - \lambda \cdot C_{in} - (1 - \lambda) \cdot C_{out})^2}, \quad (15)$$

which implies that when $C_{in} > C_{out}$ (*resp.* $C_{in} < C_{out}$), for the increasing λ , a small change in \mathcal{G} may result in a large (*resp.* small) change in P-Rank, which makes P-Rank an *ill-conditioned* (*resp.* a *well-conditioned*) problem; when $C_{in} = C_{out}$, the value of $\kappa_\infty(\mathcal{G})$ is independent of λ .

5 An Efficient Algorithm for P-Rank Estimating on Undirected Graphs

In this section, we study the P-Rank optimization problem over undirected networks.

The key idea in our optimization is to maximally use the adjacency matrix \mathbf{A} to characterize the P-Rank similarity \mathbf{S} as a power series in a function form like $\mathbf{S} = \sum_{k=0}^{+\infty} f(\mathbf{A}^k)$. The rationale behind this is that $\mathbf{A} = \mathbf{A}^T$ for undirected networks, implying that there exists a diagonal matrix \mathbf{D} such that $\mathbf{Q} = \mathbf{P} = \mathbf{D} \cdot \mathbf{A}$ in Eq.(8). Hence, computing \mathbf{S} boils down to solving $\sum_{k=0}^{+\infty} f(\mathbf{A}^k)$. For computing $f(\mathbf{A}^k)$, it is far less costly to first diagonalize \mathbf{A} into $\mathbf{\Lambda}$ via eigen-decomposition than to compute \mathbf{A}^k in a brute-force way. Then calculating $f(\mathbf{A}^k)$ reduces to computing the function on each eigenvalue for \mathbf{A} .

More specifically, the main result of this section is the following.

Theorem 3. *For undirected networks, the P-Rank similarity problem of Eq.(8) can be solvable in $O(n^3)$ worst-case time.*

It is worth noting that SimRank over undirected graphs is a special case of the more general P-Rank when $\lambda = 1$ and $C_{in} = C_{out}$. In contrast to the $O(n^3 + Kn^2)$ -time of

Algorithm 1. ASAP ($\mathcal{G}, \lambda, C_{in}, C_{out}$)

-
- Input** : a labeled undirected network $\mathcal{G} = (\mathcal{V}, \mathcal{E}, \mathcal{A}; l)$, the weighting factor λ , and in- and out-link damping factors C_{in} and C_{out} .
- Output**: similarity matrix $\mathbf{S} = (s_{i,j})_{n \times n}$ with $s_{i,j}$ denoting P-Rank score between vertices i and j .
- 1 initialize the adjacency matrix \mathbf{A} of \mathcal{G} ;
 - 2 compute the diagonal matrix $\mathbf{D} = \text{diag}(d_{1,1}, d_{2,2}, \dots, d_{n,n})$ with its entry $d_{i,i} = (\sum_{j=1}^n a_{i,j})^{-1}$, if $\sum_{j=1}^n a_{i,j} \neq 0$; and $d_{i,i} = 0$, otherwise;
 - 3 compute the auxiliary matrix $\mathbf{T} = \mathbf{D}^{1/2} \cdot \mathbf{A} \cdot \mathbf{D}^{1/2}$
 - 4 decompose \mathbf{T} into the diagonal matrix $\mathbf{\Lambda} = \text{diag}(\Lambda_{1,1}, \Lambda_{2,2}, \dots, \Lambda_{n,n})$ and the orthogonal \mathbf{U} via QR factorization *s.t.* $\mathbf{T} = \mathbf{U} \cdot \mathbf{\Lambda} \cdot \mathbf{U}^T$;
 - 5 compute the auxiliary matrix $\mathbf{\Gamma} = (\Gamma_{i,j})_{n \times n} = \mathbf{U}^T \cdot \mathbf{D}^{-1} \cdot \mathbf{U}$ and $\mathbf{V} = \mathbf{D}^{1/2} \cdot \mathbf{U}$ and the constant $C = \lambda C_{in} + (1 - \lambda) C_{out}$;
 - 6 compute the matrix $\mathbf{\Psi} = (\psi_{i,j})_{n \times n}$ whose entry $\psi_{i,j} = \Gamma_{i,j} / (1 - C \cdot \Lambda_{i,i} \cdot \Lambda_{j,j})$;
 - 7 compute the P-Rank similarity matrix $\mathbf{S} = (1 - C) \cdot \mathbf{V} \cdot \mathbf{\Psi} \cdot \mathbf{V}^T$;
 - 8 **return** \mathbf{S} ;
-

SimRank optimization over undirected graphs in our early work [14], this work further optimizes P-Rank time in $O(n^3)$ with no need for iteration.

To show Theorem 3, we first characterize P-Rank elegantly on undirected graphs.

Theorem 4. For the undirected network \mathcal{G} with n vertices, let (i) $\mathbf{A} = (a_{i,j}) \in \mathbb{R}^{n \times n}$ be the adjacency matrix of \mathcal{G} , (ii) $\mathbf{D} \in \mathbb{R}^{n \times n}$ is a diagonal matrix³.

$$\mathbf{D} = \text{diag}\left(\left(\sum_{j=1}^n a_{1,j}\right)^{-1}, \dots, \left(\sum_{j=1}^n a_{n,j}\right)^{-1}\right), \quad (16)$$

and (iii) \mathbf{U} and $\mathbf{\Lambda}$ the eigen-decomposition results of the matrix $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$, in which $\mathbf{U} \in \mathbb{R}^{n \times n}$ is an orthogonal matrix whose columns are all eigenvectors of $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$, and $\mathbf{\Lambda} = (\Lambda_{i,j}) \in \mathbb{R}^{n \times n}$ is an diagonal matrix with its diagonal entries being all eigenvalues of $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$.

Then the P-Rank similarity matrix \mathbf{S}' of Eq.(9) can be explicitly computed as⁴.

$$\mathbf{S}' = \mathbf{D}^{1/2} \mathbf{U} \cdot \mathbf{\Psi} \cdot \mathbf{U}^T \mathbf{D}^{1/2}, \quad (17)$$

where

$$\mathbf{\Psi} = (\Psi_{i,j})_{n \times n} = \left(\frac{[\mathbf{U}^T \mathbf{D}^{-1} \mathbf{U}]_{i,j}}{1 - (\lambda \cdot C_{in} + (1 - \lambda) \cdot C_{out}) \Lambda_{i,i} \Lambda_{j,j}} \right)_{n \times n}, \text{ and} \quad (18)$$

$[\mathbf{U}^T \mathbf{D}^{-1} \mathbf{U}]_{i,j}$ denotes the (i, j) -entry of the matrix $\mathbf{U}^T \mathbf{D}^{-1} \mathbf{U}$.

Due to space limitations, a detailed proof of this theorem can be found in [11].

We next prove Theorem 3 by providing an algorithm for P-Rank computation over undirected networks.

³ We define $\frac{1}{0} \triangleq 0$, thereby avoiding division by zero when the column/row sum of \mathbf{A} equals 0.

⁴ $\mathbf{D}^{1/2}$ is a diagonal matrix whose diagonal entries are the principal square root of those of \mathbf{D} .

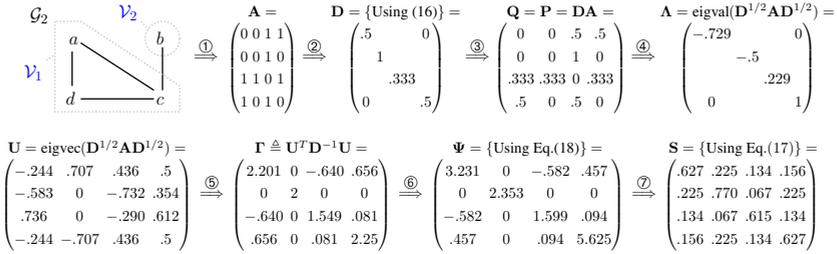


Fig. 2. An example of computing P-Rank over undirected network \mathcal{G}_2 via non-iterative method

Algorithm. The algorithm, referred to as **ASAP**, is shown in Algorithm 1. It takes as input a labeled undirected network \mathcal{G} , a weighting factor λ , and in- and out-link damping factors C_{in} and C_{out} ; and it returns the P-Rank similarity matrix $\mathbf{S} = (s_{i,j})_{n \times n}$ of all vertex-pairs in \mathcal{G} .

The algorithm **ASAP** works as follows. It first initializes the adjacency matrix \mathbf{A} of the network \mathcal{G} (line 1). Using \mathbf{A} , it then compute the auxiliary diagonal matrix \mathbf{D} whose (i, i) -entry equals the reciprocal of the i -th column sum of \mathbf{A} , if this reciprocal exists; or 0, otherwise (line 1). **ASAP** then uses QR eigen-decomposition [13] to factorize $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$ as $\mathbf{U} \mathbf{\Lambda} \mathbf{U}^T$, in which all columns of \mathbf{U} are the eigenvectors of $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$, and all diagonal entries of $\mathbf{\Lambda}$ are the corresponding eigenvalues of $\mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$ (lines 1-1). Utilizing \mathbf{U} and $\mathbf{\Lambda}$, it calculates $\mathbf{\Psi}$ (lines 1-1) to obtain the similarity matrix \mathbf{S} (lines 1-1), which can be justified by Eqs.(17) and (18).

We now give a running example to show how **ASAP** computes the P-Rank similarity matrix \mathbf{S} for a given network.

Example 2. Consider a labeled undirected network \mathcal{G}_2 with 4 vertices $\mathcal{V} = \mathcal{V}_1 \cup \mathcal{V}_2 = \{a, c, d\} \cup \{b\}$ and 4 edges $\mathcal{E} = \{(a, c), (a, d), (c, d), (b, c)\}$. Figure 2 depicts the detailed process of computing \mathbf{S} step by step with no need for any iteration. **ASAP** returns \mathbf{S} as the P-Rank matrix, which is exactly the solution to the P-Rank formula of Eq.(8).

To complete the proof of Theorem 3, we next show that algorithm **ASAP** has cubic-time complexity bound in the number of vertices.

Complexity. (i) In lines 1-1, computing the diagonal \mathbf{D} and $\mathbf{T} = \mathbf{D}^{1/2} \mathbf{A} \mathbf{D}^{1/2}$ takes $O(m)$ and $O(n^2)$ time, respectively, with n and m being the number of vertices and edges in \mathcal{G} respectively. (ii) In line 1, QR factorization of \mathbf{T} into the orthogonal \mathbf{U} and the diagonal $\mathbf{\Lambda}$ requires $O(n^3)$ worst-case time. (iii) In lines 1-1, computing the auxiliary matrices $\mathbf{\Gamma}$, \mathbf{V} , $\mathbf{\Psi}$ and the similarity matrix \mathbf{S} yields $O(n^3)$, $O(n^2)$, $O(n^2)$ and $O(n^3)$, respectively, which can be bounded by $O(n^3)$. Combining (i), (ii) and (iii), the total time of **ASAP** is in $O(n^3)$.

6 Experimental Evaluation

We next present an experimental study of our P-Rank estimating methods. Using real-life and synthetic data, we conducted two sets of experiments to evaluate the accuracy, stability, computational efficiency of our approaches for similarity estimation *v.s.* (a)

the conventional pruning P-Rank iterative paradigm [7] and (b) the memoization-based P-Rank algorithm [8,7].

Experimental setting. We used real-life data and synthetic data.

(1) *Real-life data.* The real-life data was taken from DBLP⁵. We extracted the 10-year (from 1998 to 2007) author-paper information from the entire DBLP dataset. We picked up papers published on 6 major conferences (“ICDE”, “VLDB”, “SIGMOD”, “WWW”, “SIGIR” and “KDD”). Every two years made a time step. For each time step, we built a co-authorship network incrementally from the one of previous time step. We chose the relationship that there is an edge between authors if one author wrote a paper with another. The sizes of these DBLP networks are as follows:

DBLP Data	1998-1999	1998-2001	1998-2003	1998-2005	1998-2007
n	1,525	3,208	5,307	7,984	10,682
m	5,929	13,441	24,762	39,399	54,844

(2) *Synthetic data.* We also used a C++ boost generator to produce graphs, controlled by two parameters: the number n of vertices, and the number m of edges. We then produced a set of 5 networks (undirected RAND data) by increasing the vertex size n from 100K to 1M with edges randomly chosen.

(3) *Algorithms.* We have implemented the following algorithms in C++: (a) our algorithm **ASAP**; (b) the conventional P-Rank iterative algorithm **Iter** [7] with the radius-based pruning technique; (c) the memoization-based algorithm **Memo** [8] applied on P-Rank; (d) SimRank optimized algorithm **AUG** [14] over undirected graphs.

The experiments were run on a machine with a Pentium(R) Dual-Core (2.0GHz) CPU and 4GB RAM, using Windows Vista. Each experiment was repeated 5 times and the average is reported here.

For ease and fairness of comparison, the following parameters were used as default values (unless otherwise specified). We set the in-link damping factor $C_{in} = 0.8$, the out-link damping factor $C_{out} = 0.6$, the weighting factor $\lambda = 0.5$, the total iteration number $k = 10$, the desired accuracy $\epsilon = 0.001$.

Experimental Results. We now present our findings.

Exp-1: Accuracy. In the first set of experiments, we evaluated the accuracy of P-Rank iteration in Eq.(6) using synthetic and real data. We also investigated the impact of in- and out-link damping factors on P-Rank accuracy, using synthetic data.

We considered various weighting factors λ from 0 to 1 for 5 RAND networks. For each RAND data with vertex size ranged from 0.2M to 1M, fixing the damping factors $C_{in} = 0.8$ and $C_{out} = 0.6$, we varied the number k of P-Rank iterations in Eq.(4) ranged from 2 to 20 for each vertex-pair. Due to space limitations, Figure 3(a) only reports the results over the RAND network (with 1M vertices), which visualizes the P-Rank accuracy *w.r.t.* the number of iterations performed. Here, the accuracy is measured

⁵ <http://www.informatik.uni-trier.de/~ley/db/>

by the absolute difference between the iterative P-Rank and the exact solution⁶. Note that the logarithmic scale is chosen across the y -axis in Figure 3(a) to provide a more illustrative look for the asymptotic rate of P-Rank convergence. For each fixed λ , the downward lines for P-Rank iterations reveal an exponential accuracy as k increases, as expected in Theorem 1. We also observe that the larger λ may dramatically increase the slope of a line, which tells that increasing the weighting factor decreases the rate of convergence for P-Rank iteration.

Using the same RAND, we fixed the desired accuracy $\epsilon = 0.001$ and varied damping factors by increasing C_{in} and C_{out} from 0.1 to 0.9. Fixing $C_{out} = 0.6$, the result of varying C_{in} is reported in Figure 3(b) (for space constraints, a similar result of varying C_{out} is omitted), in which the x -axis represents the value of in-link damping factor, and y -axis gives the number of iterations needed for attaining the given accuracy ϵ . When $\lambda = 0$, the curve in Figure (b) visually approaches a horizontal line. This is because in this case, P-Rank boils down to an iterative form of the reversed SimRank with no in-link similarity considered, which makes C_{in} insensitive to the final P-Rank score. When $0 < \lambda \leq 1$, k shows a general increased tendency as C_{in} is growing. This tells us that small choices of damping factors may reduce the number of iterations required for a fixed accuracy, and hence, improves the efficiency of P-Rank, as expected.

To evaluate the impact of both C_{in} and C_{out} w.r.t. the accuracy, we used the real DBLP data. We only report the result on DBLP 1998-2007 data in Figure 3(c), which shows a 3D shaded surface from the average of accuracy value for all vertex-pairs on z -axis when we fixed $k = 10$ and $\lambda = 0.5$, and varied C_{in} and C_{out} in x -axis and y -axis, respectively. It can be seen that the residual becomes huge only when C_{in} and C_{out} are both increasing to 1; and the iterative P-Rank is accurate when C_{in} and $C_{out} < 0.6$. This explains why small choices of damping factors are suggested in P-Rank iteration.

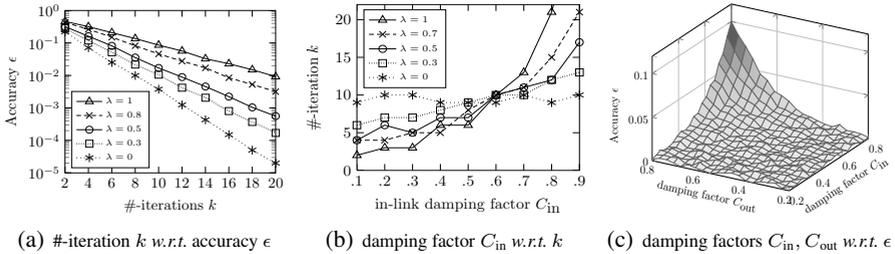


Fig. 3. Experimental Results on P-Rank Accuracy

Exp-2: Stability. We evaluated P-Rank stability using synthetic data and DBLP.

Fixing the value of out-link damping factor $C_{out} = 0.6$, we varied the values of C_{in} from 0.2 to 0.8. The result over RAND 1M data is reported in Figure 4(a), in which x -axis indicates various weighting factors λ with their sizes ranged from 0 to 1. Accordingly, varying λ from 0 and 1, Figure 4(b) visualizes the impact of in-link damping factor C_{in} on P-Rank stability when $C_{out} = 0.6$ is fixed.

⁶ To select the P-Rank “exact” solution $s(\cdot, \cdot)$, we used the Cauchy’s criterion for convergence and regarded the 100th iterative $s^{(100)}(\cdot, \cdot)$ score as the “exact” one s.t. $|s^{(100)}(\cdot, \cdot) - s^{(101)}(\cdot, \cdot)| \ll 1 \times 10^{-10}$.

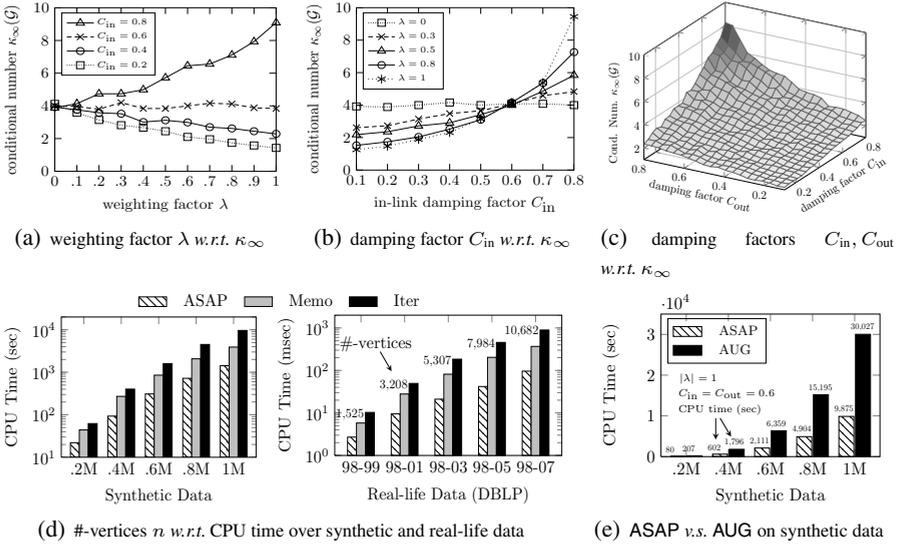


Fig. 4. Experimental Results on P-Rank Stability and Time Efficiency

The results in Figures 4(a) and 4(b) both show that increasing weighting factor λ induces a large P-Rank conditional number when $C_{in} > 0.6$. Notice that for different C_{in} , there is one common point $(\lambda, \kappa_\infty) = (0, 4)$ of intersection of all curves in Figure 4(a); correspondingly, in the extreme case of $\lambda = 0$, the curve in Figure 4(b) approaches to a horizontal line. These indicate that varying C_{in} when $\lambda = 0$ has no effect on the stability κ_∞ of P-Rank, for in this case only the contribution of out-links is considered for computing P-Rank similarity. When $C_{in} < 0.6$, however, $\kappa_\infty(\mathcal{G})$ is decreased as λ grows. This tells that small quantities of weighting factor and damping factors yield small P-Rank conditional numbers, and thus make the P-Rank well-conditioned, as expected in Theorem 2.

For real-life datasets, Figure 4(c) comprehensively depicts in 3D view the impacts of both C_{in} and C_{out} on P-Rank stability over DBLP 1M data, in which x - and y -axis represent in- and out-link damping factors respectively, and z -axis stands for the P-Rank conditional number. The result demonstrates that P-Rank is comparatively stable when both C_{in} and C_{out} are small (less than 0.6). However, when C_{in} and C_{out} are approaching to 1, P-Rank is ill-conditioned and not very useful since small perturbations in similarity computation may cause P-Rank scores drastically altered, which carries the risk of producing nonsensical similarity results. In light of this, small choices of damping factors are preferable.

Exp-3: Time Efficiency over Undirected Networks. In the third set of experiments, we used synthetic RAND and the real DBLP data to evaluate the benefits of the time-efficient algorithm ASAP. Figure 4(d) compares the performance of ASAP with those of Memo and Iter on both datasets. We use the logarithmic scale on the CPU time (y -axis). The iteration number for Iter and Memo is set to 10. Note that different time unit is chosen across the vertical axis in the two plots of Fig.4(d) to provide a clear

look for each bar shape. (i) Varying the number of vertices from 200K to 1M, the result on RAND indicates that **ASAP** outperformed **Memo** and **Iter**; the computational time of **ASAP** has almost one order of magnitude faster than **Iter**, *i.e.*, computing P-Rank similarity from the explicit characterization of its solution is efficient. In most cases, there are a considerable amount of repeated iterations for **Iter** and **Memo** to reach a fixed-point of P-Rank scores, and these impede their time efficiency in P-Rank computation. (ii) The result on DBLP also shows the running time with respect to the number of nodes for P-Rank estimation when the sizes of DBLP data increased from 1.5K to 10K. In all cases, **ASAP** performed the best, by taking advantage of its non-iterative paradigm.

To compare the performances of **ASAP** and **AUG**, we applied them to compute Sim-Rank similarities over synthetic RAND data, by setting $\lambda = 1$ for **ASAP** (a special case of P-Rank without out-links consideration). Figure 4(e) reports the result over synthetic RAND data. It can be seen that **ASAP** runs approx. 3 times faster than **AUG** though the CPU time of the **ASAP** and **AUG** are of the same order of magnitude. The reason is that after eigen-decomposition, **AUG** still requires extra iterations to be performed in the small eigen-subspace, which takes a significant amount of time, whereas **ASAP** can straightforwardly compute similarities in terms of eigenvectors with no need for iterations.

7 Related Work

There has been a surge of studies (*e.g.*, [2,15,3,9,4,5,16,6]) on link-based analysis over information networks in recent years. PageRank became popular since the famous result of Page *et al.* [6] was used by the Google search engine for ranking web sites based on link structures. Since then, a host of new ranking algorithms for web pages have been developed. The famous results include the HITS algorithm [2] proposed by Jon *et al.* (now used www.ask.com), SimRank [17,10,5], SimFusion [15,4] and P-Rank algorithm [7].

SimRank [5] is a recursive structural similarity measure based on the intuition that “*two objects are similar if they are related to similar objects*”, which extends the Bibliometric Coupling and Co-citation [3,1] beyond the local neighborhood so that the information of the entire network can be exploited globally. A naive iterative approach was initially proposed in [5] to compute the SimRank score with a pruning mechanism, which is in quartic worst-case time. Several optimization problems were investigated for SimRank estimation, including the pair-wise iterative methods [5,8], matrix-based approaches [8,12,17,14] and probabilistic algorithms [18].

More recently, Zhao *et al.* [7] presented a new P-Rank model when noticing the limited information problem of SimRank —the similarity scores are only determined by their in-link relationships. P-Rank measure refines SimRank by jointly considering both in- and out-links of entity pairs. The conventional algorithm for computing P-Rank is based on iterative techniques, which still requires quartic time complexity. To optimize its computational time, a similar memoization approach in [8] can be applied to P-Rank, which reduces its time to be cubic in the worst case. In comparison, our work focuses on the problems of P-Rank accuracy, stability and computational time over undirected graphs.

Closer to this work are [14,8]. A time-efficient algorithm **AUG** for SimRank computation was proposed on undirected graphs in [14], which is in $O(n^3 + kn^2)$ time. In contrast, we further improve [14] (i) by providing a non-iterative $O(n^3)$ -time algorithm that explicitly characterizes the similarity solution, and (ii) by extending SimRank to the general P-Rank measure. An accuracy estimation for SimRank was addressed in [8], it differs from this work in that our focus is on P-Rank estimation, in which the accuracy depends on both in- and out-link damping factors rather than the in-link damping factor alone.

8 Conclusions

In this study, several P-Rank problems were investigated. First, we have proposed an accuracy estimate for the P-Rank iteration, by finding out the exact number of iterations needed to attain a given accuracy. Second, we have introduced the notion of P-Rank conditional number based on the matrix representation of P-Rank. A tight bound of P-Rank conditional number has been provided to show how the weighting factor and the damping factors affect the stability of P-Rank. Finally, we have also devised an $O(n^3)$ -time algorithm to deal with the P-Rank optimization problem over undirected networks. Our empirical results have verified the accuracy, stability and computational efficiency of our approaches.

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