Efficient presolving methods for the influence maximization problem

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\textbf{Abstract}
We consider the influence maximization problem (IMP) which asks for identifying a limited number of key individuals to spread influence in a network such that the expected number of influenced individuals is maximized. The stochastic maximal covering location problem (SMCLP) formulation is a mixed integer programming formulation that effectively approximates the IMP by the Monte-Carlo sampling. For IMPs with a large-scale network or a large number of samplings, however, the SMCLP formulation cannot be efficiently solved by existing exact algorithms due to its large problem size. In this paper, we attempt to develop presolving methods to reduce the problem size and hence enhance the capability of employing exact algorithms in solving large-scale IMPs. In particular, we propose two effective presolving methods, called strongly connected nodes aggregation (SCNA) and isomorphic nodes aggregation (INA), respectively. The SCNA enables to build a new SMCLP formulation that is potentially much more compact than the existing one, and the INA further eliminates variables and constraints in the SMCLP formulation. A theoretical analysis on two special cases of the IMP is provided to demonstrate the strength of the SCNA and INA in reducing the problem size of the SMCLP formulation. We integrate the proposed presolving methods, SCNA and INA, into the Benders decomposition algorithm, which is recognized as one of the state-of-the-art exact algorithms for solving the IMP. We show that the proposed SCNA and INA provide the possibility to develop a much faster separation algorithm for the Benders cuts. Numerical results demonstrate that with the SCNA and INA, the Benders decomposition algorithm is much more effective in solving the IMP in terms of solution time.

\textbf{KEYWORDS}
Benders decomposition, influence maximization, integer programming, presolving methods, stochastic programming

1 \textbf{INTRODUCTION}

Nowadays, with the popularity of online social network sites such as Facebook, Instagram, and Twitter, propagation of influence in social networks has received more and more attention. Promotion of products, ideas, and specific behavior patterns can all be viewed as propagation of influence. In practice, influence spreads among individuals through the so-called word-of-mouth exchanges. Individuals with more social connections can be seen as more influential, which means they are more likely to exert influence on others. In this setting, one related optimization problem, called the \textit{influence maximization problem} (IMP), is to...
select a limited number of key individuals as a seed set, denoted as $S$, to trigger a spread process in the social network such that the expected number of influenced individuals is maximized after the spread. Mathematically, the IMP can be written as

$$\max_{S \subseteq \mathcal{V}, |S| \leq K} \sigma(S),$$

(1)

where $\mathcal{V}$ is the set of individuals in the social network, $S \subseteq \mathcal{V} (|S| \leq K \in \mathbb{Z}^+)$ is the seed set of key individuals that need to be identified, and $\sigma(S)$ is the influence function measuring the expected number of individuals in the social network that can be influenced by the individuals in the seed set $S$.

The IMP plays a crucial role in various social network applications, such as viral marketing [14, 21] and rumor control [9, 31]. For a comprehensive overview of this topic, we refer to [11]. Aside from the applications in social networks, similar concepts have been investigated in other areas, including the spread of epidemics [8, 22], network monitoring [41], habitat conservation [54], and analysis of protein-interaction networks [33]. Online network tools enable collection of a huge number of individuals and a huge amount of information about the network structures, and hence provide good opportunities to address these problems. However, they also lead to large-scale networks (with millions/billions of nodes and arcs), presenting new challenges in solving large-scale IMPs. Therefore, development of efficient algorithms to obtain a high-quality solution for large-scale IMPs is greatly needed.

### 1.1 Literature review

Kempe et al. [36] first proposed the discrete optimization problem formulation (1) for the IMP. Depending on different influence diffusion processes, they introduced two fundamental influence propagation models for the IMP: the independent cascade model (ICM) and the linear threshold model (LTM). They showed that under both the ICM and LTM, the IMP is NP-hard, indicating that achieving an optimal solution of the IMP is challenging for the large-scale cases. By proving that the influence function $\sigma(S)$ is monotone and submodular, they were able to design a greedy algorithm, which starts with $S = \emptyset$ and iteratively adds the individual with maximal marginal gain, with an approximation ratio of $(1 - 1/e)$ (here $e$ denotes the base of the natural logarithm). Unfortunately, as shown in [14, 15], given a fixed seed set $S$, it is #P-hard to compute $\sigma(S)$ exactly. Therefore, Kempe et al. [36] proposed Monte-Carlo sampling, which provides a subset of equiprobable scenarios, of a reasonable size, to estimate $\sigma(S)$ (each scenario is represented by a live-arc graph). Following [36], many researchers focused on the improvement of the greedy algorithm and the development of other heuristic algorithms for solving the IMP. Specifically, Leskovec et al. [41] utilized the submodularity of $\sigma(S)$ and presented an improved greedy algorithm called cost-effective lazy forward. According to their numerical results, their method is almost 700 times faster than the basic greedy algorithm of Kempe et al. [36]. Chen et al. [13] proposed another algorithmic enhancement for the greedy algorithm which reduces the graph searching time on computing the marginal gain (achieved by adding an individual into a seed set). Furthermore, they developed a much more efficient algorithm, called degree discount, for the IMP under the ICM that nearly matches the performance of the greedy algorithm. The two-phase influence maximization [56] and the influence maximization via martingales [57] heuristic algorithms also deserve special attention. They can not only guarantee an approximation ratio of $(1-1/e)$, but also enable to solve large-scale IMPs in nearly linear time. We refer to [14, 16, 25, 37] for more greedy or heuristic algorithms for solving the IMP and [44] for a detailed comparison among different heuristic algorithms.

Most heuristic algorithms find a suboptimal solution for the IMP with some worst-case guarantees. However, in some applications, it is crucial to identify an optimal solution instead of just a suboptimal one; see [27]. As a result, using exact algorithms to solve the IMP has attracted more and more attention recently. In most exact algorithms, a mixed integer programming (MIP) formulation is established. In particular, Wu and Küçükyavuz [59] transferred the IMP into the so-called two-stage stochastic submodular MIP model and proposed a delayed constraint generation algorithm to solve the problem to optimality. The computational results indicate that their algorithm is more efficient than the basic greedy algorithm in [36], especially when $K$ is large. Given a collection of scenarios $\Omega$, Güney [27], Güney et al. [28], and Li et al. [45] formulated the IMP as a stochastic maximal covering location problem (SMCLP) with $\Theta(|\mathcal{V}||\Omega|)$ variables and linear constraints. Following this line, Güney et al. [28] developed a reformulation of the SMCLP and proposed a Benders decomposition (BD) algorithm. Their experiment results show that the BD algorithm outperforms the one in [59] by several orders of magnitude in terms of solution time. We refer to [24, 29, 34, 35, 50] for employing exact algorithms in solving several variants of the IMP.

However, due to the NP-hardness of the IMP, the above exact algorithms are still inefficient, especially when the size of the network or the number of scenarios is large. Presolving [2] is an appealing strategy to address this issue. It removes redundant information and strengthens the model formulation with the aim of improving the performance of the subsequent solution procedure (e.g., the branch-and-cut or the BD approach). Indeed, presolving has been recognized as a standard routine of the state-of-the-art MIP solvers. For problems with specific structures, developing customized presolving methods is often much more effective; see [7, 18, 32, 46] for using customized presolving methods to solve various problems. In terms of the IMP, few articles are devoted to the development of customized presolving methods. To the best of our knowledge, only two simple
presolving methods have been developed in the literature [28, 34] and they have been proved to be beneficial to solving the IMP in certain cases. Consequently, it is crucial to develop more customized presolving methods to further enhance the capability of using exact algorithms to solve large-scale IMPs.

1.2 Contributions and outline

In this paper, we attempt to develop more presolving methods based on the SMCLP formulation to improve the solution efficiency for solving the IMP. The main contributions of this paper are summarized as follows.

- By exploiting the problem structure of the IMP, we propose two new presolving methods including (i) the strongly connected nodes aggregation (SCNA) which aggregates nodes in each strongly connected component (SCC), in a given live-arc graph, into a single virtual node; and (ii) the isomorphic nodes aggregation (INA), which extends the above idea to using isomorphic nodes among different live-arc graphs for aggregation (two nodes in different live-arc graphs are called isomorphic if the nodes that can influence them are identical in the corresponding live-arc graphs). We show that the proposed presolving methods can effectively reduce the problem size of the SMCLP formulation. In particular, after applying the SCNA, the IMP can be built on new live-arc graphs obtained by aggregating all SCCs in the original live-arc graphs, leading to a potentially much smaller SMCLP formulation.

- To demonstrate the strength of the proposed SCNA and INA in reducing the problem size of the SMCLP formulation, we provide a theoretical analysis on two special cases of the IMP: one is built on the one-way bipartite network under the LTM and the other one is built on the complete network under the ICM. For the first one, we give upper bounds, which are linear with the size of the network but independent of the number of scenarios, for the numbers of variables and constraints in the reduced SMCLP formulation (obtained by applying the proposed presolving methods). For the second one, we provide a lower bound for the probability that after applying the SCNA and INA, there are only |V| + 1 variables and two linear constraints in the reduced SMCLP formulation. We show that such a probability can tend to one under certain conditions.

- We integrate the SCNA and INA into the BD algorithm [28], which is recognized as one of the state-of-the-art exact algorithms to solve the IMP. We show that the proposed SCNA and INA provide the possibility to develop a much faster separation algorithm for the Benders cuts, as compared with the one in [28].

- Extensive numerical results on real-world networks demonstrate that (i) the proposed SCNA and INA are quite effective in reducing the problem size of the SMCLP formulation; (ii) when integrating them into the BD algorithm, they can effectively speed up the solution procedure of the IMP.

The remainder of the paper is organized as follows. Section 2 briefly reviews two fundamental influence propagation models (ICM and LTM) and the SMCLP formulation for the IMP. Section 3 presents the SCNA and INA and Section 4 further shows their theoretical strength in reducing the problem size of the SMCLP formulation. Section 5 describes the implementation of the INA and the integration of the SCNA and INA with the BD algorithm. Section 6 provides the computational results. Section 7 studies a generalization of the IMP and shows that the proposed SCNA and INA can also be applied under some realistic conditions. Finally, Section 8 gives some concluding remarks.

2 PROPAGATION MODELS AND PROBLEM FORMULATION

In this section, we briefly review the propagation models and the SMCLP formulation for the IMP [27, 28, 45]. We use a directed graph \( G = (V, A) \) to refer to a network, in which a node \( i \in V \) represents the individual involved in the influence spread, and an arc \( (i, j) \in A \) represents that individual \( i \) has the potential ability to influence (or activate) individual \( j \). The spread of influence in a given network \( G \) needs to obey certain propagation rules. Kempe et al. [36] provided the following two fundamental influence propagation models called ICM and LTM.

- In the ICM, each arc \( (i, j) \in A \) is assigned an activation probability \( \pi_{ij} \). The propagation process starts with a given seed set \( S \). If node \( i \) has been activated at the beginning of step \( t \), then during step \( t \), it has a single chance to activate its (inactive) neighbor node \( j \) with probability \( \pi_{ij} \) independently. If the activation is unsuccessful, node \( i \) has no chance to influence node \( j \) any more. Besides, for those nodes that are successfully activated during step \( t \), they will remain active and attempt to activate their inactive neighbor nodes during step \( t + 1 \). When no more inactive nodes are activated, the diffusion process is terminated.

- In the LTM, each arc \( (i, j) \in A \) is associated with a predefined weight \( b_{ij} \geq 0 \) satisfying \( \sum_{j : (i, j) \in A} b_{ij} \leq 1 \) for all \( i \in V \). In addition, each node \( j \in V \) selects a threshold value \( \theta_j \) randomly chosen from \([0, 1]\) before the propagation process.
Let $S_t$ denote the activated nodes set at the beginning of step $t$ \( (S_0 := S \text{ is the seed set}) \). Then an inactive node $j \in \mathcal{V}$ can be activated during step $t$ if and only if \( \sum_{i \in S} b_{ij} \geq \theta_j \), that is, the total contribution of its active neighbors’ influence weights exceeds its threshold value $\theta_j$. In analogy to the ICM, if a node is successfully activated during some step, it will remain active during the following steps, and the entire propagation process stops until no more nodes can be activated.

Given a seed set $S$, the results (i.e., the distributions of influenced nodes) returned by the ICM and LTM can be different [11]. For a comparison on the performance of the two models in different applications, we refer to [3, 43].

Although the influence spread under the ICM or LTM is a stochastic process, Kempe et al. [36] showed that it can be equivalently converted into a (discrete) deterministic process. More specifically, let $\Omega$ be the set of all possible scenarios of influence spread. Each scenario $\omega \in \Omega$ corresponds to a subgraph $G^\omega = (\mathcal{V}, A^\omega)$ of $G$ (called a live-arc graph) with a probability $p^\omega$ (satisfying $\sum_{\omega \in \Omega} p^\omega = 1$). Here arc \((i, j) \in A^\omega\) indicates that in scenario $\omega$, if node $i$ is activated during the influence spread, then node $j$ must be activated by it. Let $\sigma^\omega(S)$ represent the number of activated nodes in $G^\omega$. Then, $\sigma^\omega(S) = \{ i \in \mathcal{V} : \text{there exists a directed path in graph } G^\omega \text{ from a node in } S \text{ to node } i \}$. The influence function $\sigma(S)$ can equivalently be calculated by

\[
\sigma(S) = \sum_{\omega \in \Omega} p^\omega \sigma^\omega(S). \tag{2}
\]

We next discuss the computation of the probability $p^\omega$ of each scenario $\omega$ and the number of scenarios under the ICM or LTM. Under the ICM, to construct a live-arc graph $G^\omega$, each arc $(i, j) \in A$ is independently determined to be live with probability $\pi_{ij}$. Hence, the probability of $G^\omega$ is $p^\omega = \prod_{(i, j) \in A} \pi_{ij} \prod_{(i, j) \in A \setminus A^\omega} (1 - \pi_{ij})$ and the number of all possible live-arc graphs is $2^{\binom{|A|}{2}}$.

Under the LTM, for each node $j \in \mathcal{V}$, we select at most one incoming arc $(i, j)$ in $G$ to be live with probability $b_{ij}$, and do not select any arc with the probability $1 - \sum_{(i, j) \in A} b_{ij}$. As a result, (i) each live-arc graph $G^\omega$ has a probability $p^\omega = \prod_{j \in \mathcal{V}} \theta_j$, where $\theta_j := b_{ij}$ if $(i, j) \in A^\omega$; and $\theta_j := 1 - \sum_{(i, j) \in A} b_{ij}$, otherwise; and (ii) the number of all possible live-arc graphs is $\prod_{j \in \mathcal{V}} (\theta_j + 1)$, where $\theta_j$ denotes the number of incoming arcs of node $j$ in graph $G$. It is worthwhile remarking that the stochastic IMP under the ICM or LTM is equivalent to the IMP constructed via a finite number of live-arc graphs (in the sense that the distributions of nodes influenced by a given seed set are equivalent); see Kempe et al. [36].

Given a network $G$ and a set of scenarios $\Omega$, we next review the SMCLP formulation for the IMP [27, 28, 45]. First, for each scenario $\omega \in \Omega$ and node $i \in \mathcal{V}$, we denote $R(\mathcal{G}^\omega, i)$ as the reachability set of nodes that can activate node $i$ in live-arc graph $G^\omega$ (i.e., $R(\mathcal{G}^\omega, i) = \{ j \in \mathcal{V} : \text{there exists a directed path in graph } G^\omega \text{ from node } j \text{ to node } i \}$). Then, for each $\omega \in \Omega$ and $i \in \mathcal{V}$, we introduce binary variables $y_i$ and $z_{i\omega}$ to denote whether node $i$ is selected as a seed node and whether node $i$ can be activated in scenario $\omega$, respectively, that is,

\[
y_i = \begin{cases} 1, & \text{if node } i \in \mathcal{V} \text{ is chosen as a seed node;} \\ 0, & \text{otherwise;} \end{cases} \tag{3a}
\]

\[
z_{i\omega} = \begin{cases} 1, & \text{if node } i \in \mathcal{V} \text{ can be activated in scenario } \omega \in \Omega; \\ 0, & \text{otherwise.} \end{cases} \tag{3b}
\]

Using the above notations, the authors in [27, 28, 45] formulated the IMP as the following SMCLP:

\[
\max_{y, z} \sum_{\omega \in \Omega} p^\omega \sum_{i \in \mathcal{V}} z_{i\omega} \tag{3a}
\]

subject to

\[
\sum_{j \in R(\mathcal{G}^\omega, i)} y_j \geq z_{i\omega}, \quad \forall \omega \in \Omega, \quad \forall i \in \mathcal{V}, \tag{3b}
\]

\[
\sum_{j \in \mathcal{V}} y_j \leq K, \tag{3c}
\]

\[
y_j \in \{0, 1\}, \quad \forall j \in \mathcal{V}, \tag{3d}
\]

\[
z_{i\omega} \in \{0, 1\}, \quad \forall \omega \in \Omega, \quad \forall i \in \mathcal{V}. \tag{3e}
\]

In this formulation, the objective function \( (3a) \) maximizes the expected number of influenced nodes in the network $G$. Reachability constraints \((3b)\) indicate that if node $i$ in scenario $\omega$ can be activated, then at least one of the nodes in its reachability set $R(\mathcal{G}^\omega, i)$ is chosen as a seed node. Constraint \((3c)\) limits the cardinality of the set of seed nodes up to $K$. Finally, constraints \((3d)\) and \((3e)\) restrict variables $y$ and $z$ to be binary.

Unfortunately, formulation \((3)\) is computationally intractable for the network with realistic dimensions due to the huge number of scenarios (for the IMP under the ICM and LTM, the numbers of all possible scenarios are both exponential). Hence, the Monte-Carlo sampling approach is often used to approximate the influence diffusion process in which a reasonable size of equiprobable scenarios set $\Omega' \subset \Omega$ is generated, and the objective function in \((3a)\) is replaced by $\sum_{\omega \in \Omega'} p'_\omega \sum_{i \in \mathcal{V}} z_{i\omega}$ where $p'_\omega = 1/|\Omega'|$ and hence $\sum_{\omega \in \Omega'} p'_\omega = 1$ [27, 59]. The rationale behind this is that from the approximation result in [38], the probability
of obtaining an optimal solution of the IMP (3) by solving the sampling version of the IMP converges to one exponentially fast as \( |\Omega'| \to \infty \). In practice, however, the selection of the number of scenarios \( |\Omega'| \) is crucial for the approximation quality of the sampling version of the IMP. In general, the larger the \( |\Omega'| \), the smaller the approximation error is. We refer to Section 4.6 of [27] and Section 5.2 of [34] for the empirical studies of the effect of the number of scenarios on the approximation quality of the sampling version of the IMP and a variant of the IMP, respectively. Here we also want to highlight that the problem size of the sampling version of the IMP also grows linearly with the number of scenarios \( |\Omega'| \) (as both numbers of variables and constraints are \( \Theta(|\mathcal{V}| |\Omega'|) \)). This further makes it difficult to solve the problem by standard MIP solvers or the BD approach in [28], especially when the size of graph \( G \) is also large. In the next section, we shall resolve this difficulty by proposing two new presolving methods to reduce the problem size of the SMCLP formulation (3).

In the remaining of this paper, we will consider the sampling version of the IMP. For simplicity of notations, we continue to use \( \Omega \) and \( p^\omega \) to represent the set of sampling scenarios and the probability of occurrence of sampling scenario \( \omega \), respectively.

3  |  TWO PRESOLVING METHODS

In this section, by exploiting the problem structure of formulation (3), we propose two presolving methods to reduce the problem size of formulation (3). Specifically, Section 3.1 studies the SCNA which aggregates the nodes in each SCC, in a given live-arc graph, into a single node, and Section 3.2 investigates the INA which extends the idea of the SCNA to applying isomorphic nodes aggregations among different live-arc graphs.

3.1  |  Strongly connected nodes aggregation

In this subsection, we present a presolving method by considering the SCCs in a given live-arc graph. To begin with, we consider the example live-arc graph (corresponds to some scenario \( \omega \in \Omega \)) in Figure 1. In this graph, there exists a directed path (an arc) from node 1 to node 2, and as a result, if node 1 is activated by some seed node, node 2 can also be activated. Conversely, the fact that there exists a directed path (an arc) from node 2 to node 1 implies that if node 2 is activated by some seed node, node 1 can also be activated. This means that either (i) nodes 1 and 2 are simultaneously activated by some seed node; or (ii) neither of them can be activated. Consequently, \( z_1^\omega = z_2^\omega \) must hold in formulation (3). This reveals some redundancy in formulation (3) as it uses two variables \( z_1^\omega \) and \( z_2^\omega \) and two constraints in (3b) without considering \( z_1^\omega = z_2^\omega \). Indeed, to simplify the problem formulation, we can remove variable \( z_2^\omega \) and its corresponding reachability constraint in (3b) and add the objective coefficient of variable \( z_1^\omega \) into that of variable \( z_2^\omega \).

In general, for any given two strongly connected nodes \( i_1, i_2 \in \mathcal{V} \) in a live-arc graph \( G^\omega \) (i.e., there exists a directed path from node \( i_1 \) to node \( i_2 \) in \( G^\omega \) and vice versa), we can remove one of the two variables and the corresponding reachability constraint in (3b) from formulation (3). Notice that for a given SCC in a live-arc graph, as all of its nodes are strongly connected, we can recursively apply the above argument until there remains only a single variable and a single constraint in (3b) associated with this SCC. This provides us with the following presolving method.

**SCNA.** For each live-arc graph \( G^\omega \), let \( \{ SCC^\omega_i \}_{i=1}^{n_\omega}, s_\omega \in \mathbb{Z}_{++} \), be all its SCCs. For each SCC \( SCC^\omega_i \), variables \( z^\omega_i, i \in SCC^\omega_i \), are first substituted by a new variable \( z^\omega_{\text{SC}} \) with its objective coefficient being \( p^\omega |SC^\omega_i| \). Then, all but one of constraints in (3b) associated with variable \( z^\omega_{\text{SC}} \) are removed from the SMCLP formulation (3).

To implement the SCNA, we only need to identify all SCCs in all live-arc graphs \( G^\omega, \omega \in \Omega \). For each graph \( G^\omega \), this can be done in linear time \( O(|\mathcal{V}| + |A^\omega|) \) using, for example, the Kosaraju–Sharir’s algorithm [53]. Consequently, the overall complexity to implement the SCNA for formulation (3) is \( O(\sum_{\omega \in \Omega}(|\mathcal{V}| + |A^\omega|)) \).

After applying the SCNA, the IMP can be equivalently constructed based on a new set of live-arc graphs, which are potentially much more compact than the original live-arc graphs. To be more specific, by aggregating each SCC of \( G^\omega \) into a single node with the weight being the size of the SCC, we can get a directed acyclic graph, denoted as \( G'^\omega = (\mathcal{V}', \mathcal{A}') \). Each node

![Figure 1](image-url)
This reduction the numbers of variables \(y\) and the corresponding reachability constraints in the reduced formulation are equal to \(\sum_{\omega \in \Omega} |\overline{V}_\omega|\) (which is the number of the SCCs in the original live-arc graphs). This can be potentially much smaller than those in formulation (3), especially for the case where the numbers of SCCs are much smaller than the numbers of nodes in the live-arc graphs. As a result, it can be expected that solving formulation (6) is much more efficient than solving formulation (3). In addition, the fact that formulation (6) is built on the (potentially) compact and directed acyclic live-arc graphs plays an important role in improving the performance of the BD algorithm (see Section 5.2 further ahead).

### 3.2 Isomorphic nodes aggregation

The SCNA performs reductions on two nodes \(i_1\) and \(i_2\) in a given live-arc graph \(G^o\) where nodes \(i_1\) and \(i_2\) are strongly connected, or equivalently, the reachability sets of nodes \(i_1\) and \(i_2\) are identical, that is, \(R(G^o, i_1) = R(G^o, i_2)\). In this subsection, we concentrate on the extension of the result to the isomorphic nodes among different live-arc graphs. As it has been previously mentioned, two nodes \(i_1\) and \(i_2\) in two different live-arc graphs \(G^o\) and \(G^i\) are called isomorphic if their reachability sets are identical, that is,

\[
R(G^o, i_1) = R(G^i, i_2).
\]

We begin with the following observation stating that the values of variables \(z\) are determined by the values of variables \(y\) in formulation (3).

**Observation 3.1.** There must exist an optimal solution \((\overline{y}, \overline{z})\) of formulation (3) such that

\[
z^o_{\omega_i} = \min \left\{ 1, \sum_{j \in R(G^o, i)} y_j \right\}, \forall \omega \in \Omega, \forall i \in V.
\]

In particular, if node \(i \in V\) does not have any incoming arc in \(G^o\) for some \(\omega \in \Omega\), that is, \(R(G^o, i) = \{i\}\) is a singleton, the associated reachability constraint in (3b) reduces to \(y_i \geq z^o_{\omega_i}\). By Observation 3.1, we can get \(z^o_{\omega_i} = \min\{1, y_i\} = y_i\). As a result, we can perform a reduction on formulation (3) by aggregating \(z^o_{\omega_i} := y_i\) and removing the associated constraint in (3b). We call this reduction the *singleton node aggregation* (SNA). Indeed, this is exactly the “P2” presolving method proposed in Güney et al. [28].

We next use Observation 3.1 to derive the INA. Let \(i_1\) and \(i_2\) be two isomorphic nodes in two different live-arc graphs \(G^o\) and \(G^i\). By Observation 3.1, there must exist an optimal solution \((\overline{y}, \overline{z})\) such that
\[ \tau^0_{i_1} = \min \left\{ 1, \sum_{j \in R(G^0, i_1)} y_j \right\} \quad \text{and} \quad \tau^0_{i_2} = \min \left\{ 1, \sum_{j \in R(G^0, i_2)} y_j \right\}. \] (9)

By (7), we have \( \tau^0_{i_1} = \tau^0_{i_2} \). This implies that setting \( \tau^0_{i_1} : = \tau^0_{i_2} \) in formulation (3) does not change its optimal value. Consequently, we have the following presolving method.

**INA.** If nodes \( i_1 \) and \( i_2 \) in two different live-arc graphs \( G^0 \) and \( G^n \) are isomorphic, variable \( z^0_{i_1} \) can be replaced by variable \( z^0_{i_2} \) and constraint \( \sum_{j \in R(G^0, i_1)} y_j \geq z^0_{i_1} \) can be removed from formulation (3).

The SCNA can be regarded as a special case of the INA, which is restricted to aggregating the isomorphic (strongly connected) nodes inside each live-arc graph. However, as it has been discussed in Section 3.1, implementing the SCNA can be done in \( \mathcal{O}\left( \sum_{\omega \in \Omega} (|\mathcal{V}| + |A^0|) \right) \), which is much faster than that of implementing the INA. The latter requires to check whether or not condition (7) holds for all 4-tuples \((\alpha, \eta, i_1, i_2)\) with an overall complexity of \( \mathcal{O}(\|\Omega\|^2 |\mathcal{V}|^3) \). This shows that, to implement the INA, it is better to first implement the SCNA, and then detect isomorphic nodes among different scenarios based on the compact formulation (6) (in Section 5.1, we shall provide a fast heuristic algorithm for implementing the INA). After applying the INA on formulation (6), the formulation of the IMP can be presented as follows:

\[
\begin{align*}
\max_{y, z} & \quad \sum_{\omega \in \Omega} \sum_{u \in \bar{G}^0} f^\omega_{u, z^\omega}, \\
\text{s.t.} & \quad (3c), (3d), \\
& \quad \sum_{v \in R(G^0, \omega)} y(S\bar{C}^\omega_v) \geq z^\omega_{u}, \quad \forall \omega \in \Omega, \ \forall u \in \bar{V}^\omega, \\
& \quad z^\omega_{u} \in \{0, 1\}, \quad \forall \omega \in \Omega, \ \forall u \in \bar{V}^\omega, \quad (10)
\end{align*}
\]

where \( \bar{V}^\omega \) denotes the set of nodes in \( \bar{G}^0 \) that are not aggregated by some other nodes in other live-arc graphs and \( f^\omega_{u, z^\omega} \) denotes the objective coefficient of variable \( z^\omega_{u} \) after applying the INA. For simplicity, for two isomorphic nodes \( i_1 \) and \( i_2 \) in two different live-arc graphs \( G^0 \) and \( G^n \), we aggregated variable \( z^\omega_{i_1} \) by variable \( z^\omega_{i_2} \) if \( \omega > \eta \). It is worthwhile to highlight that, with the increasing number of scenarios \( |\Omega| \), a node in a live-arc graph \( G^0 \) is more likely to be aggregated by other isomorphic nodes (in other live-arc graphs), and consequently, \( |\bar{V}^\omega| \) in formulation (9) tends to be smaller.

**Remark 3.2.** The SNA can be used to further simplify formulation (9). In particular, if \( \bigcup_{\omega \in R(G^0, \omega)} S\bar{C}^\omega_v = \{j\} \), we can aggregate \( z^\omega_{u} : = y_j \) and remove the corresponding reachability constraint in (9).

### 4 | THEORETICAL ANALYSIS

In this section, we demonstrate the strength of the proposed SCNA and INA in reducing the problem size of the SMCLP formulation (3) by analyzing two special cases of the IMP. In particular, for the first case where the IMP is built upon a one-way bipartite network under the LTM, we provide upper bounds, which are linear with the size of the network but independent of the number of scenarios, for the numbers of variables and constraints in the reduced SMCLP formulation (obtained by applying the proposed presolving methods). For the second case where the IMP is built upon a complete network under the ICM, we provide a lower bound for the probability that after applying the SCNA and INA, there are only \(|\mathcal{V}| + 1\) variables and two linear constraints in the reduced SMCLP formulation (9). Such a probability can tend to one under certain conditions.

#### 4.1 | One-way bipartite network under the LTM

A one-way bipartite network is a bipartite graph in which all arcs are from one side (the source nodes) to another side (the target nodes). One-way bipartite networks also arise from several applications [4, 6, 23, 30, 55]. For example, the authors in [4, 30, 55] considered one of the major decisions in a marketing plan that deals with the allocation of a given budget among media channels (i.e., source nodes) in order to maximize the influence on a set of potential customers (i.e., target nodes), which can be characterized by a one-way bipartite network. Other applications on the one-way bipartite networks include, e.g., the human sexual contact network in [23] (where the nodes denote the groups of two different genders and the arcs denote the sexual connections between males and females) and the collaboration network in [6] (the source and target nodes denote the organizations and the projects, respectively, and an arc between organization \( i \) and project \( j \) denotes that organization \( i \) participates in project \( j \)). Due to the simple structure, some theoretical properties on the influence propagation in one-way...
bipartite networks have also been established. In particular, it is possible to compute the exact influence coverage \( \sigma(S) \) by a dynamic programming procedure; see [60, 61]. We remark that most existing works assume that the influence spreads from the source nodes to the target nodes. In this subsection, we consider the generalized case where the source nodes and target nodes can also exert influence to themselves. In the following, we consider the IMP built upon such a one-way bipartite network under the LTM.

Let \( G_B = (M \cup N, A) \) be a given one-way bipartite graph. All arcs in \( A \) are from the source nodes set \( M \) to the target nodes set \( N \). For each scenario \( \omega \in \Omega \), we denote its live-arc graph as \( G_B^\omega = (M \cup N, A^\omega) \). Notice that under the LTM, in each live-arc graph \( G_B^\omega \), each node \( i, \omega \in \Omega \) does not have any incoming arc and each node \( j, \omega \in \Omega \) has at most one incoming arc. Therefore, under the LTM, the reachability set of a node \( i \in M \) is \( R(G_B^\omega, i) = \{i\} \), and the reachability set of a node \( j \in N \) is

\[
R(G_B^\omega, j) = \begin{cases} \{j\}, & \text{if node } j \text{ does not have any incoming arc in } G_B^\omega; \\ \{i, j\}, & \text{if there exists some node } i \in M \text{ such that } (i, j) \in A^\omega. \end{cases}
\]

(11)

Next, for each node \( i \in M \cup N \), we define a set of scenarios

\[
\Omega(i) := \{ \omega \in \Omega : R(G_B^\omega, i) = \{i\} \},
\]

(12)

and for each arc \((i, j) \in A\), we define another set of scenarios

\[
\Omega(i, j) := \{ \omega \in \Omega : R(G_B^\omega, j) = \{i, j\} \}.
\]

(13)

By definition, it follows that

\[
\Omega = \begin{cases} \Omega(i), & \text{for } i \in M; \\ (\bigcup_{i \in M} \Omega(i, j)) \cup \Omega(j), & \text{for } j \in N. \end{cases}
\]

(14)

Then, we have the followings:

- by applying the SNA for each \( i \in M \cup N \), we can aggregate \( z_i^{\omega} := y_i \) for all \( \omega \in \Omega(i) \) and remove the corresponding constraints in (3b);
- by applying the INA for each \((i, j) \in A' := \{(i, j) \in A : \Omega(i, j) \neq \emptyset\}\), we can aggregate all variables \( z_j^{\omega}, \omega \in \Omega(i, j) \), into a single variable, denoted as \( z_{ij} \), and remove the redundant constraints in (3b).

As a result, the reduced formulation is given by

\[
\begin{align*}
\max_{j, \omega \in \Omega(i)} & \quad \sum_{i \in M \cup N} s_i y_i + \sum_{(i, j) \in A'} c_{ij} z_{ij} \\
\text{s.t.} & \quad y_i + y_j \geq z_{ij}, \quad \forall (i, j) \in A', \\
& \quad \sum_{i \in M \cup N} y_i \leq K, \quad \forall i \in M \cup N, \\
& \quad z_{ij} \in \{0, 1\}, \quad \forall i \in M \cup N, \\
& \quad z_{ij} \in \{0, 1\}, \quad \forall (i, j) \in A',
\end{align*}
\]

(15a, 15b, 15c, 15d, 15e)

where \( s_i := \sum_{\omega \in \Omega(i)} p^\omega \) for \( i \in M \cup N \) and \( c_{ij} := \sum_{\omega \in \Omega(i, j)} p^\omega \) for \((i, j) \in A'\), respectively. Since \(|A'| \leq |A|\), we have the following theorem providing upper bounds for the numbers of variables and constraints in the reduced SMCLP formulation (14).

**Theorem 4.1.** Consider the IMP on the one-way bipartite network \( G_B \) with a finite set of scenarios \( \Omega \) under the LTM. Applying the SNA and INA on formulation (3), the numbers of variables and constraints in the reduced SMCLP formulation (14) are at most \(|M| + |N| + |A|\) and \(|A| + 1\), respectively.

Finally, we provide more analysis results for formulation (14). To proceed, we note that using (13) and \( \sum_{\omega \in \Omega} p^\omega = 1 \), we have the following properties on the objective coefficients of formulation (14).

**Remark 4.2.** (i) \( s_i = 1 \) for all \( i \in M \); and (ii) \( s_j + \sum_{(i, j) \in A'} c_{ij} = 1 \) for all \( j \in N \).

**Proposition 4.3.** The linear programming (LP) relaxation of formulation (14) is tight. Moreover, formulation (14) can be solved in strongly polynomial time.

**Proof.** The proof is given in Appendix A.
4.2 Complete network under the ICM

In this subsection, we study another special case of the IMP where the considered network is a complete graph (denoted as $G_C = (V_C, A_C)$) and the influence propagation model is the ICM. Let $G_C^\omega = (V_C, A_C^\omega)$ be a live-arc graph and denote $n = |V_C|$. Recall that for the live-arc graph constructed under the ICM, each arc is determined to be live independently with probability $\pi_{ij}$. Consequently, $G_C^\omega$ can be seen as a directed Erdős-Rényi (ER) random graph [10, 20]. If the arc probabilities are homogeneous, that is, $\pi_{ij} = p$ for all $(i,j) \in A_C$ and some $p \in (0, 1]$, $G_C^\omega$ is a homogeneous directed ER random graph; otherwise it is an inhomogeneous directed ER random graph. Homogeneous directed ER random graph is shown to be strongly connected with a probability tending to one (as $n \to \infty$) under certain conditions; see, for example, [26]. The following lemma further provides a lower bound for the probability of the strong connectivity of $G_C^\omega$ with respect to the number of nodes $n$ and the arc probability $p$.

**Lemma 4.4.** Suppose that $\pi_{ij} = p \in (0, 1]$ for all $(i,j) \in A_C$ and
\[
\max \left\{ (n-1)(1-p^2)^{n+1}, 2(1-p^2)^{n-1} \right\} \leq 1.
\]
Then
\[
\mathbb{P}(\text{Graph } G_C^\omega \text{ is strongly connected}) \geq 1 - n(n-1)(1-p^2)^{n-1}.
\]

**Proof.** The proof can be found in Section 1 of [12].

**Theorem 4.5.** Consider the IMP on the complete network $G_C$ with a finite set of scenarios $\Omega$ under the ICM. Suppose that $\pi_{ij} \geq p \in (0, 1]$ for all $(i,j) \in A_C$ and (15) holds. Then, by applying the SCNA and INA, there are only $n + 1$ variables and two linear constraints in the reduced SMCLP formulation (9) with a probability at least $p^*$ where
\[
p^* = (1 - n(n-1)(1-p^2)^{n-1})^{\left|\Omega\right|}.
\]

**Proof.** Notice that the probability that $G_C^\omega$ is strongly connected with $\pi_{ij} \geq p$ for all $(i,j) \in A_C$ is larger than or equal to that with $\pi_{ij} = p$ for all $(i,j) \in A_C$. This, together with Lemma 4.4 and the fact that each live-arc graph $G_C^\omega$ is constructed independently, shows that the probability that all live-arc graphs $G_C^\omega$, $\omega \in \Omega$, are strongly connected is at least $p^*$ (defined in (17)). The strong connectivity of graph $G_C^\omega$ implies that the number of SCCs in $G_C^\omega$ is one and $R(G_C^\omega, i) = V_C$ for all $i \in V_C$. As a result, with a probability at least $p^*$, (i) the number of variables $z$ in the reduced formulation (6) (after applying the SCNA on (3)) is equal to $|\Omega|$; and (ii) the number of variables $z$ in the reduced formulation (9) (after applying the INA on (6)) is equal to one. This, together with the fact that there is a cardinality constraint (3c) in (9), completes the proof.

The value $p^*$ in Theorem 4.5 can tend to one. For instance, if $p \in (0, 1]$ is a constant, then $p^* \to 1$ as $n \to \infty$ (notice that in this case, condition (15) also holds). This shows that after applying the SCNA and INA, there are only $n + 1$ variables and two linear constraints in the reduced SMCLP formulation (9) with a probability tending to one.

It is worthwhile remarking that Theorem 4.5 also sheds a useful insight that for the IMP with a general large and well-connected network $G$ (not necessary to be complete) with high arc probabilities $\pi_{ij}$, the SCNA and INA can be expected to effectively reduce the sizes of the live-arc graphs and SMCLP formulation (3). Indeed, a well-connected network is likely to contain large complete subgraphs. By Theorem 4.5, with high arc probabilities, the nodes in these complete subgraphs are likely to be strongly connected in all live-arc graphs $G^\omega$, $\omega \in \Omega$, and as a result, more reductions are likely to be detected. This is consistent with the computational results in Section 6.2 where more reductions can be detected by the proposed SCNA and INA for large and well-connected networks with large arc probabilities (see Table 2 further ahead).

5 THE ALGORITHMS

In this section, we first discuss the implementation of the INA in Section 5.1. Then we present the BD algorithm with the proposed SCNA and INA for solving the IMP in Section 5.2.

5.1 An algorithm for identifying presolving reductions by the INA

A straightforward implementation of the INA requires to first precompute and store the reachability sets of all nodes in all live-arc graphs $G^\omega$, $\omega \in \Omega$, and then detect all 4-tuples $(\omega, \eta, i_1, i_2)$ that satisfy condition (7). However, this leads to a high
runtime complexity and a large memory consumption, which are \( \Theta(|\Omega|^2 |V|^2) \) and \( \Theta(|\Omega| |V|^2) \), respectively. In this subsection, we shall overcome this weakness by presenting a hashing-based heuristic algorithm.

We first discuss the computation of the reachability sets. Güney et al. [28] computed the reachability set \( R(G^o, i) \) of each node \( i \) in each live-arc graph \( G^o, o \in \Omega \), by applying a reverse breadth–first search (BFS) starting from node \( i \). The computational complexity is \( \Theta(|V|+|A^o|) \). Here we notice that it can be (possibly) much faster to compute \( R(G^o, i) \) based on the compact graph \( \overline{G}^o \). Indeed, as it has been mentioned in Section 3.1, the reachability sets of nodes inside a given SCC \( SC^o_u \) of \( G^o \) are identical, where \( u \) is the corresponding node in graph \( \overline{G}^o \). Hence, to compute the reachability sets of the nodes in SCC \( SC^o_u \), we only need to compute the reachability set \( R(G^o, SC^o_u) \). To compute the latter one, we can apply a reverse BFS in the compact graph \( \overline{G}^o \) and use relation (4). The related complexity is \( \Theta(|\Omega|+|\overline{A}^o|) \), which is potentially much smaller than \( \Theta(|V|+|A^o|) \), especially when \( |\overline{A}^o| \) is much smaller than \( |A^o| \). Furthermore, as \( \overline{G}^o \) is a directed acyclic graph (as each node forms an SCC), we can perform a topological ordering to further speed up the procedure of computing the reachability sets of all nodes in \( \overline{G}^o \). To be more specific, topological ordering for the directed acyclic graph \( \overline{G}^o \) is a linear ordering of nodes such that for each arc \((u_1, u_2) \in \overline{A}^o \), node \( u_1 \) comes before node \( u_2 \) in the ordering. In our implementation, we traverse all nodes in \( \overline{G}^o \) to compute their reachability sets according to the topological ordering. In other words, when computing the reachability set of node \( u \in \overline{V}^o \), the reachability sets of nodes \( u', u' \in \mathcal{N}_w(u) \), have been computed, where \( \mathcal{N}_w(u) := \{ u' : (u', u) \in \overline{A}^o \} \) is the set of node \( u \)'s incoming neighbor nodes. Therefore, to compute node \( u \)'s reachability set, we only need to traverse its incoming neighbor nodes and use relation \( R(\overline{G}^o, u) = \bigcup_{v \in \mathcal{N}_w(u)} R(\overline{G}^o, v) \cup \{ u \} \). This avoids performing a whole reverse BFS and generally accelerates the computation of \( R(\overline{G}^o, u) \).

Next, we discuss the storage of the reachability sets. First, we can implement SCNA to alleviate the memory consumption of storing the reachability sets (as for each SCC, only a single node’s reachability set needs to be stored). Second, to further avoid a large memory consumption, we only store those nodes’ reachability sets whose sizes are smaller than or equal to a predefined parameter \texttt{MaxReacSize}. This also means that only nodes satisfying this criterion will be used for detecting the reductions by the INA. The rationale behind this strategy is that for the nodes with smaller reachability sets, it is more likely to detect nodes that are isomorphic to them, as illustrated in our computational results (see Section 6.4 further ahead).

Finally, we apply the INA by detecting the pairs whose reachability sets are identical, that is,

\[
\bigcup_{v \in R(\overline{G}^o, u)} SC^o_v = \bigcup_{v \in R(\overline{G}^o, u)} SC^o_v, \quad (19)
\]

for some \( u \in \overline{V}^o \) and \( u_0 \in \overline{V}^o \). To do this, we follow [2] to use a hashing-based method. The basic idea of the hashing-based method is to simultaneously build a hashing table that remembers the information of the reachability sets, and test whether there exists a reachability set in the hashing table that is identical to the one we are currently looking at. Specifically, let \( H \) be the hashing table and for each \( o \in \Omega \) and \( u \in \overline{V}^o \), let \( \bigcup_{v \in R(\overline{G}^o, u)} SC^o_v \) be the key with 3-tuple \((o, u, R(\overline{G}^o, u))\) being the stored value in table \( H \). At first, table \( H \) is initialized to be \( \emptyset \). Then, in each iteration, for scenario \( o \in \Omega \) and node \( u \in \overline{V}^o \), if \(|\bigcup_{v \in R(\overline{G}^o, u)} SC^o_v| \leq \text{MaxReacSize} \), table \( H \) is queried for \( \bigcup_{v \in R(\overline{G}^o, u)} SC^o_v \). If condition (18) holds for some corresponding entry \((\eta, u_0, R(\overline{G}^o, u))\) in table \( H \), we apply the INA by removing variable \( z^o_\eta \), deleting the associated constraint in (5), and adding the objective coefficient of variable \( z^o_\eta \) into that of variable \( z^o_{\eta'} \); otherwise, tuple \((o, u, R(\overline{G}^o, u))\) will be added into table \( H \). The procedure is repeated until all considered reachability sets are tested.

In summary, we present the implementation of the INA in Algorithm 1 to obtain the reduced SMCLP formulation (9). Here, for simplicity of presentation, the improvement of topological ordering is omitted in Algorithm 1. In step 4, we perform a reverse BFS on node \( u \) in graph \( \overline{G}^o \) to compute \( R(\overline{G}^o, u) \). In steps 5–9, we use the hashing-based method to detect whether there exists some entry \((\eta, u_0, R(\overline{G}^o, u))\) in table \( H \) such that (18) holds. If yes, we apply the INA reductions; otherwise, we add the new entry \((o, u, R(\overline{G}^o, u))\) into table \( H \).

### 5.2 BD algorithm for solving the IMP

Güney et al. [28] have proposed the BD algorithm to solve the IMP based on formulation (3). The authors showed the effectiveness of integrating the presolving method SNA into the BD algorithm. In this subsection, to further enhance the capability of using the BD algorithm to solve the IMP, we attempt to integrate the proposed SCNA and INA into the BD algorithm, or equivalently, to design a BD algorithm that is based on the reduced formulation (9) and the compact graphs \( \overline{G}^o, o \in \Omega \).
5.2.1 Reformulation of (9)

We first briefly introduce the BD reformulation of (9) (more details can be found in [28]). To begin with, we note that replacing each binary variable \( z^\omega_u \) by a continuous variable taking value in \([0, 1]\) does not change the optimal value of formulation (9). For each \( \omega \in \Omega \), let \( \varphi^\omega \) represent the variable that captures the contribution of scenario \( \omega \) to the objective function. Then, we can project out variables \( z \) and equivalently reformulate (9) as

\[
\max_{y, \varphi} \left\{ \sum_{\omega \in \Omega} \varphi^\omega : (3c), (3d), \varphi^\omega \leq \Phi^\omega(y), \forall \omega \in \Omega \right\},
\]

where function \( \Phi^\omega(y) \) is defined as follows:

\[
\Phi^\omega(y) := \max_{z} \left\{ \sum_{u \in \tilde{V}^\omega} f^\omega_u : z^\omega_u \leq \sum_{v \in R(\tilde{G}^\omega, u)} y(S^\omega_{cv}), 0 \leq z^\omega_u \leq 1, \forall u \in \tilde{V}^\omega \right\}.
\]

For a fixed \( \bar{y} \in [0, 1]^{|\mathcal{F}|} \), to model the inequalities \( \varphi^\omega \leq \Phi^\omega(y) \) for all \( \omega \in \Omega \), we use the Benders optimality cuts which are derived as follows. First, the dual of formulation (20) when \( y = \bar{y} \) is

\[
\min_{\alpha^\omega, \beta^\omega} \left\{ \sum_{u \in \tilde{V}^\omega} \alpha^\omega_u + \sum_{v \in R(\tilde{G}^\omega, u)} \bar{y}(S^\omega_{cv}) + \beta^\omega_u : \alpha^\omega_u + \beta^\omega_u \geq f^\omega_u, \alpha^\omega_u, \beta^\omega_u \geq 0, \forall u \in \tilde{V}^\omega \right\},
\]

where \( \alpha^\omega_u \) and \( \beta^\omega_u \) are the dual variables of constraints \( z^\omega_u \leq \sum_{v \in R(\tilde{G}^\omega, u)} \bar{y}(S^\omega_{cv}) \) and \( z^\omega_u \leq 1 \), respectively. Clearly, formulation (21) has a closed form solution \((\tilde{\alpha}^\omega_u, \tilde{\beta}^\omega_u)\):

\[
(\tilde{\alpha}^\omega_u, \tilde{\beta}^\omega_u) = \begin{cases} (0, f^\omega_u), & \text{if } \sum_{v \in R(\tilde{G}^\omega, u)} \bar{y}(S^\omega_{cv}) \geq 1; \\ (f^\omega_u, 0), & \text{otherwise}, \end{cases} \forall u \in \tilde{V}^\omega.
\]

Then the Benders optimality cuts for formulation (19) are given by

\[
\varphi^\omega \leq \sum_{u \in \tilde{V}^\omega} \tilde{\alpha}^\omega_u + \sum_{v \in R(\tilde{G}^\omega, u)} \bar{y}(S^\omega_{cv}) + \tilde{\beta}^\omega_u, \forall \omega \in \Omega.
\]

From (22), \((\tilde{\alpha}^\omega_u, \tilde{\beta}^\omega_u)\) depends on \( \bar{y} \). Thus, inequality (23) depends on \( \bar{y} \) as well, but we omit this dependence for notational convenience.

To solve formulation (19), we use a branch-and-Benders-cut approach in which a branch-and-cut search tree is created and the Benders optimality cuts (23) are separated at each branch-and-cut node.
An efficient separation algorithm

The problem of (19) in which inequalities (23) with \( \vec{y} = \mathbf{0} \), that is,

\[
\varphi^\omega \leq \sum_{u \in V^\omega} \bar{f}^u \sum_{v \in R(G^\omega, u)} y(SC^\omega_v), \ \forall \ \omega \in \Omega,
\]

are added. Here \( \mathbf{0} \) is the \( |V| \)-dimensional zero vector.

For a given point \( \vec{y} \in [0, 1]^{|V|} \), it is interesting to ask whether or not applying the SCNA or INA changes the Benders optimality cuts (23), or equivalently, whether or not the Benders optimality cuts (25), (26), and (23) based on formulations (3), (6), and (9) are equivalent, where

\[
\varphi^\omega \leq \sum_{i \in V} \left( \hat{\alpha}_i^\omega \sum_{j \in R(G^\omega, i)} y_j + \hat{\beta}_i^\omega \right), \ \forall \ \omega \in \Omega,
\]

\[
\varphi^\omega \leq \sum_{u \in V^\omega} \left( \alpha_u^\omega \sum_{v \in R(G^\omega, u)} y(SC^\omega_v) + \overline{\beta}_u^\omega \right), \ \forall \ \omega \in \Omega.
\]

Here,

\[
\left( \hat{\alpha}_i^\omega, \hat{\beta}_i^\omega \right) = \begin{cases} 
(0, p^\omega), & \text{if } \sum_{j \in R(G^\omega, i)} y_j \geq 1; \\
(p^\omega, 0), & \text{otherwise,}
\end{cases} \quad \forall \ i \in V,
\]

\[
\left( \alpha_u^\omega, \overline{\beta}_u^\omega \right) = \begin{cases} 
(0, p^\omega | SC^\omega_u|), & \text{if } \sum_{v \in R(G^\omega, u)} y(SC^\omega_v) \geq 1; \\
(p^\omega | SC^\omega_u|, 0), & \text{otherwise},
\end{cases} \quad \forall \ u \in V^\omega.
\]

This question is addressed by the following proposition.

**Proposition 5.1.** Given a point \( \vec{y} \in [0, 1]^{|V|} \) and a scenario \( \omega \in \Omega \) (i) (25) and (26) are equivalent; (ii) (23) and (26) may be different.

**Proof.** The proof is given in Appendix B.

By Proposition 5.1 (i), we know that Benders optimality cuts before and after applying the SCNA are identical (if INA is not applied). As a result, the SCNA will not change the path of the search tree created by the branch-and-Benders-cut algorithm. However, the separation of the Benders optimality cuts (26) after applying the SCNA could be much more efficient than that of the Benders optimality cuts (25) before applying the SCNA. Indeed, given a scenario \( \omega \), the separation of (26) can be done in \( \Theta(|V| + \sum_{u \in V^\omega} |R(G^\omega, u)|) \approx \Theta(\sum_{u \in V^\omega} |R(G^\omega, u)|) \) (by computing \( \gamma(SC^\omega_v) \) for all \( v \in V^\omega \) in \( \Theta(|V|) \) and (22) in \( \Theta(\sum_{u \in V^\omega} |R(G^\omega, u)|) \)). This could be much smaller than the complexity of a direct implementation of the separation of Benders optimality cuts (25), which is \( \Theta(\sum_{i \in V} |R(G^\omega, i)|) \), especially when

- \( G^\omega \)'s number of SCCs is much smaller than its number of nodes, or
- the number of elements in the reachability sets of the nodes in \( G^\omega \) are much smaller than that in \( G^\omega \).

Different from that of applying the SCNA, applying the INA can generally change the Benders optimality cuts, as stated in Proposition 5.1 (ii). This means that the path of the search tree created by the branch-and-Benders-cut algorithm could also be different. Nevertheless, as \( |V^\omega| \) may be smaller than \( |V| \), applying the INA can further improve the separation of the Benders optimality cuts. We remark that as stated in [28], efficient separation of Benders optimality cuts is crucial for the BD algorithm to successfully solve large-scale IMPs. In the following, we shall present a more efficient separation algorithm for the Benders optimality cuts by integrating the proposed SCNA and INA into the one developed in [28].

5.2.2 An efficient separation algorithm

Having a solution \( (\vec{y}, \vec{\varphi}) \in [0, 1]^{|V|} \times \mathbb{R}_{+}^{[2]} \) of the LP relaxation of this relaxed master problem of (19), we now describe an efficient separation algorithm for Benders optimality cuts (23). As mentioned in Section 5.1, it is unrealistic to compute and store all reachability sets of all nodes of all scenarios a priori due to the large memory consumption. Hence, similar to [28], we introduce parameter \textbf{MemLimPerScen} to denote the maximally allowed memory consumption per scenario. In particular, for each scenario \( \omega \in \Omega \), we store the reachability sets of nodes according to their topological ordering in the compact live-arc graph \( G^\omega \) until the memory consumption reaches \textbf{MemLimPerScen}. As a result, when computing Benders optimality cut (23), if \( R(G^\omega, u) \) has been stored, we access it directly; otherwise, we perform a reverse BFS to compute \( R(G^\omega, u) \) on the fly. Moreover, to further improve the efficiency of computing Benders optimality cut (23), we can omit to compute \( R(G^\omega, u) \) if \( (\hat{\alpha}_u^\omega, \overline{\beta}_u^\omega) = (0, f_u^\omega) \)
Algorithm 2. Separation of Benders optimality cuts (23)

**Input:** the compact live-arc graphs \( \hat{G}^\omega = (\hat{\mathcal{V}}, \hat{A}^\omega) \), \( \omega \in \Omega \), formulation (9), and point \((\overline{y}, \overline{\varphi}) \in [0, 1]^{|\mathcal{V}|} \times \mathbb{R}_+^{|\mathcal{A}|} \).

**Output:** the set \( C \) of Benders optimality cuts which are violated by point \((\overline{y}, \overline{\varphi}) \).

1. Initialize \( C := \emptyset \)
2. for \( \omega \in \Omega \) do
   3. Compute \( S^\omega := \{ u \in \hat{\mathcal{V}}^\omega : \gamma(SC^\omega_u) \geq 1 \} \)
   4. Perform a BFS in \( \hat{G}^\omega \) to compute \( I(\hat{G}^\omega, S^\omega) \)
   5. Initialize \( C^\omega := \sum_{\omega \in I(\hat{G}^\omega, S^\omega)} f^\omega_u \) and \( c^\omega_j := 0 \), \( j \in \mathcal{V} \)
   6. for \( u \in \hat{\mathcal{V}}^\omega \backslash I(\hat{G}^\omega, S^\omega) \) do
      7. if \( R(\hat{G}^\omega, u) \) is not stored in the memory then
         8. Perform a reverse BFS in \( \hat{G}^\omega \) to compute \( R(\hat{G}^\omega, u) \)
      end
      9. if \( \sum_{\xi \in R(\hat{G}^\omega, u)} \gamma(SC^\omega_{\xi v}) \geq 1 \) then
         10. Set \( C^\omega := C^\omega + f^\omega_u \)
      else
         11. Set \( c^\omega_j := c^\omega_j + f^\omega_u \) for all \( j \in \bigcup_{\xi \in R(\hat{G}^\omega, u)} SC^\omega_{\xi v} \)
      end
   end
   12. if \( \gamma(\overline{y}) \sum_{j \in \mathcal{V}} c^\omega_j y_j + C^\omega \) then
      13. Set \( C := C \cup \left\{ \varphi^\omega \leq \sum_{j \in \mathcal{V}} c^\omega_j y_j + C^\omega \right\} \)
   end
end

It is worth emphasizing the computational efficiency of our separation algorithm for the Benders optimality cuts over the one in [28]. First, in Algorithm 2, to compute sets \( I(\hat{G}^\omega, S^\omega) \) and \( R(\hat{G}^\omega, u) \), we perform (reverse) BFSes in the compact live-arc graph \( \hat{G}^\omega \), which is potentially much faster than that in [28] where the (reverse) BFSes in the (original) live-arc graph \( G^\omega \) are performed. Second, compared with that in [28], fewer reverse BFSes will be performed in Algorithm 2 due to the following two reasons.

- \( |\hat{\mathcal{V}}^\omega \backslash I(\hat{G}^\omega, S^\omega)| \) in step 6 of Algorithm 2 can be potentially much smaller than that in [28], which is \( |\mathcal{V} \backslash I(G^\omega, \hat{S}^\omega)| \).

Here \( \hat{S}^\omega := \{ j \in \mathcal{V} : \gamma_j = 1 \} \), and \( I(G^\omega, \hat{S}^\omega) \) denotes the set of nodes in \( \mathcal{V} \) that can be influenced by some node in \( \hat{S}^\omega \) in graph \( G^\omega \). Indeed, a node \( u \in \hat{\mathcal{V}}^\omega \backslash I(\hat{G}^\omega, S^\omega) \) corresponds to an SCC in \( G^\omega \) whose nodes cannot be influenced by the nodes in \( \bigcup_{v \in S^\omega} SC^\omega_v \). As a result,

\[
|\hat{\mathcal{V}}^\omega \backslash I(\hat{G}^\omega, S^\omega)| \leq |\mathcal{V} \backslash I(G^\omega, \hat{S}^\omega)| \leq |\mathcal{V} \backslash I(G^\omega, S^\omega)|,
\]

where the last inequality follows from \( \hat{S}^\omega \subseteq \bigcup_{v \in S^\omega} SC^\omega_v \) and the fact that \( I(G^\omega, S_1) \subseteq I(G^\omega, S_2) \) for any \( S_1 \subseteq S_2 \subseteq \mathcal{V} \).
TABLE 1  Eight real-world social networks.

| Network          | |V|  | |A|  | |A'| | |A|  | ρ | Description                                      |
|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|
| MSG              | 1899             | 59 835           | 20 296           | 31.5             | Messaging network of UC-Irvine [49] |
| GNU              | 10 879           | 79 988           | 79 988           | 7.4              | Gnutella peer-to-peer file sharing network [51] |
| HEP              | 15 233           | 117 782          | 64 426           | 7.7              | High energy physics paper citation network [13] |
| ENRON            | 36 692           | 367 662          | 367 662          | 10.0             | Email communication network from Enron [40] |
| FACEBOOK         | 50 515           | 1 638 612        | 1 638 180        | 32.4             | Facebook page network in the category of artist [52] |
| DEEZER           | 54 573           | 996 404          | 996 404          | 18.3             | Deezer friendship network of users in Croatia [52] |
| TWITTER          | 81 306           | 1 768 149        | 1 768 135        | 21.7             | Social network from Twitter [39] |
| EPINIONS         | 131 828          | 841 372          | 840 799          | 6.4              | Who-trust-whom social network of Epinions [42] |

- As stated in Section 5.2.1, the sum of the sizes of the reachability sets \( R(G^ω, i), i \in V \), could be much smaller than that of the sizes of the reachability sets \( R(G^ω, i), i \in V \). This implies that for a fixed \( \text{MemLimPerScen} \), more reachability sets \( R(G^ω, u) \) are likely to be stored a priori, and thus the condition in step 7 of Algorithm 2 is less likely to occur.

6  | COMPUTATIONAL RESULTS

In this section, we present the computational results to show the effectiveness of the proposed SCNA and INA. We use the BD algorithm in Section 5.2, which was implemented in C++ linked with IBM ILOG CPLEX optimizer 20.1.0 [19]. The Benders optimality cuts were added using CALLABLE LIBRARIES under the default settings of the branch-and-cut framework of CPLEX. The time limit was set to 14400 seconds, and all the experiments were performed on a cluster of Intel(R) Xeon(R) Gold 6140 CPU @ 2.30GHz computers. Only a single core was used in our experiments. We note here that throughout this section, all averages are taken to be geometric means. Since the statistics can be zero, we use the shifted geometric mean with a shift of 1 (the shifted geometric mean of values \( x_1, x_2, \ldots, x_n \) with shift \( s \) is defined as \( \prod_{k=1}^n (x_k + s)^{1/n} - s; \) see [1]).

6.1  | Networks and settings

Our benchmark dataset consists of eight real-world social networks. Four of them have been used in [28, 59] (MSG, GNU, HEP, and ENRON) and the other four networks are from the SNAP database 1 (FACEBOOK, DEEZER, TWITTER, and EPINIONS). The latter ones are large-scale networks which are used to test the performance of the SCNA and INA in large-scale cases.

The selections of parameters of the IMP are also similar to the one in [28, 59]. More specifically, for the IMP under both the ICM and LTM, the cardinality restriction \( K \) in formulation (3) is selected in \( \{5, 10, 15, 25\} \). As stated at the end of Section 2, the number of scenarios \( |\Omega| \) is a key parameter to achieve the trade-off between the approximation quality and solution time for solving the sampling version of the IMP. In Section 2 of [12], we follow [34] to conduct experiments to compare the approximation gaps (defined in equation (4) of [34]) of the sampling version of the IMP with different number of scenarios. The approximation gap estimates the difference of the objective value of the sampling version of the IMP and the true objective value of the IMP. The results showed that \( |\Omega| = 1000 \) is a reasonable choice to achieve a solution with a small approximation gap for instances constructed by the considered networks. Therefore, in our experiments, to reflect different approximation levels of the sampling version of the IMP, the number of scenarios \( |\Omega| \) is selected in \( \{250, 500, 1000\} \). We next discuss the selections of the activation probability \( \pi_j \) (under the ICM) and the weight \( b_{ij} \) (under the LTM) for each arc \((i, j)\). Let \( n_{ij} \) denote the number of parallel arcs from node \( i \) to node \( j \). For the IMP under the ICM, each single arc is assigned the same activation probability \( p \) chosen in \( \{0.01, 0.05, 0.10\} \). For the IMP under the LTM, we set influence weight on arc \((i, j)\) as \( 1/n_{ij} \), where \( n_{ij} := \sum_{\{i,j\} \in \mathcal{A}} n_{ij} \) (i.e., the number of incoming arcs of node \( j \) in \( G \)) is a normalization factor to ensure that the sum of weights of the incoming arcs to \( j \) is at most 1. Each live-arc graph \( G^ω \) of a scenario \( \omega \in \Omega \) is randomly constructed as follows. For a live-arc graph \( G^ω \) under the ICM, arc \((i, j) \in \mathcal{A} \) is included in \( \mathcal{A}^ω \) with probability \( \pi_{ij} = 1 - (1 - p)^{n_{ij}} \) (representing that at least one of the parallel

\(^1\)https://snap.stanford.edu/data/.

\(^2\)\( \frac{|A|}{|V|} \) is equal to the arithmetic mean of nodes’ indegrees or outdegrees of the directed network.
arcs of \((i,j)\) appears in \(G^\omega\). For a live-arc graph \(G^\omega\) under the LTM, at most one of node \(j\)'s incoming arcs \((i,j)\) is included in \(A^\omega\) with probability \(b_{ij} = n_{ij}/n_j\) (also representing that at least one of the parallel arcs of \((i,j)\) appears in \(G^\omega\)). We remark that, with more parallel arcs from node \(i\) to node \(j\), the activation probability \(x_{ij}\) (under the ICM) or weight \(b_{ij}\) (under the LTM) is larger, and thus arc \((i,j)\) will be more likely to be included in graph \(G^\omega\). For the IMP with each combination of the above parameters, five instances are randomly generated. Therefore, for each network in Table 1, we have 180 and 60 instances for the IMP under the ICM and LTM, respectively.

In our experiments, we compare the performance of the following three settings:

- Default: solving the IMP based on the Benders reformulation of (3) with the SNA applied;
- SCNA: Default with the SCNA applied;
- SCNA+INA: Default with the SCNA and INA applied.

Following [28], the memory control parameter MemLimPerScen is set to \(8/|\Omega|\) GB. Unless otherwise stated, in the implementation of the INA (i.e., Algorithm 1), parameter MaxReacSize is set to 8 and 4 for the IMP under the ICM and LTM, respectively. Finally, to avoid generating too many Benders optimality cuts at fractional points, we follow [28] to stop the separation procedure if the dual bound improves by less than 0.001.

### 6.2 Results for the IMP under the ICM

In this subsection, we test the effectiveness of the proposed presolving methods SCNA and INA for the IMP under the ICM. Table 2 reports the reductions by applying the SNA, SCNA, and INA. For convenience, we only report the results for the case \(|\Omega| = 1000\) since the results for the other two cases (\(|\Omega| = 250, 500\)) are similar. For each setting under which the corresponding presolving method(s) are applied, we use \(\Delta Z\) and \(\Delta R\) to represent the reductions in percentage of the numbers of variables \(z\) and elements in the reachability sets of nodes in all live-arc graphs. In addition, we use \(\Delta Z_{\text{SCNA}}\) and \(\Delta R_{\text{SCNA}}\) to denote the reductions in percentage by applying the SCNA, and \(\Delta Z_{\text{INA}}\) and \(\Delta R_{\text{INA}}\) to denote the reductions in percentage by applying the INA. For the SCNA, we additionally list the average percentages of nodes and arcs reductions (\(\Delta V = \sum_{\omega \in \Omega}(|V^\omega| - |\overline{V}^\omega|)/(|\Omega||V^\omega|)\) and \(\Delta A = \sum_{\omega \in \Omega}(|A^\omega| - |\overline{A}^\omega|)/(|\Omega||A^\omega|)\)) to compare the sizes of the compact live-arc graphs \(\overline{G}^\omega = (\overline{V}^\omega, \overline{A}^\omega)\) and the original live-arc graphs \(G^\omega = (V^\omega, A^\omega)\). For the SCNA, the reductions on the numbers of nodes and variables \(z\) are equal, and hence we have \(\Delta Z_{\text{SCNA}} = \Delta V\) under setting SCNA. It is worthwhile remarking that \(\Delta Z\), \(\Delta R\), \(\Delta V\), and \(\Delta A\) can reflect the efficiency of the separation algorithm for the Benders optimality cuts over the one in which no presolving method is applied. The larger the \(\Delta Z\), \(\Delta R\), \(\Delta V\), and \(\Delta A\), the more efficient the separation algorithm is.

As it can be seen in Table 2, when the network has a small connectivity \(\rho\) or activation probability \(p\), the singleton nodes (nodes without any incoming arc) are more likely to appear in the live-arc graphs. As a result, the SNA can eliminate a considerably large numbers of variables and elements in the reachability sets. In contrast, the SCNA is more effective in eliminating the variables and elements in the reachability sets when the network has a relatively large value of \(\rho\) or \(p\). This is reasonable since as \(\rho\) or \(p\) increases, the original live-arc graphs contain more arcs, and as a result, more nodes are likely to be strongly connected with other nodes. As for the numbers of nodes and arcs in the compact live-arc graphs, we can observe that they are much smaller than those in the original live-arc graphs, and in general, the larger \(\rho\) or \(p\) is, the more reductions are detected by the SCNA. For the INA, we can observe a clear reduction on the number of variables (beyond the SCNA). However, in most cases, the reduction on the number of elements in the reachability sets is relatively small, which is due to the fact that most eliminated variables are associated with small reachability sets as we set MaxReacSize = 8 in Algorithm 1 (in Section 6.4, we will perform numerical experiments to confirm that setting MaxReacSize = 8 in Algorithm 1 is enough to identify almost all pairs of isomorphic nodes).

As discussed in the end of Section 3.2, with the increasing number of scenarios in the IMP, a node in a live-arc graph \(G^\omega\) is more likely to be aggregated by other isomorphic nodes (in other live-arc graphs). Table 3 further reports the reductions on the problem size of the SMCLP formulation (3) with different numbers of scenarios under setting SCNA+INA. For convenience, we only report results for the case \(\rho = 0.1\) as the results for the other two cases are similar. As observed from Table 3, \(\Delta Z_{\text{INA}}\) and \(\Delta R_{\text{INA}}\) tend to slightly increase with the increasing of \(|\Omega|\). This shows that for the IMP under the ICM, the reductions derived by the INA slightly increase with the increasing number of scenarios.

We now evaluate the performance improvement of the integration of the SCNA and INA with the BD algorithm. In Table 4, for each of the eight networks, we report the total number of instances that can be solved within the time limit (\#S), the average CPU time in seconds (\#T), the average number of branch-and-cut nodes (\#N), the average number of added Benders optimality cuts (\#Z), the average number of added Benders optimality cuts (\#Z).

---

3This setting can be seen as the implementation in [28]. Unfortunately, we could not access the code of [28] online. Therefore, the results reported in this section are based on our implementation. Notice that, however, due to the differences in hardware and randomness in sampling, it cannot be expected that the results of setting Default are the same as those in [28].
instances, the average relative gap reduces from 7 to 5%
the CPU time includes the presolving time spent on applying the SCNA/INA and the separation time. Notice that the CPU time includes the presolving time spent on applying the SCNA/INA and the separation time. For instances that cannot be solved by any of the three settings, we report the average relative gap (Gap) in percentage, defined as 100 \times \frac{UB - LB}{LB} \text{ where } LB and UB are the lower bounds and upper bounds returned by CPLEX. “0” in a row of Table 4 denotes that all instances of the corresponding network can be solved by at least one of the three settings. Detailed statistics of these results can be found in Tables 1a-8a of [12]. As it can be seen in Table 4, the performance of setting SCNA is much better than that of setting Default, especially for instances with large and well-connected networks. In total, setting SCNA can solve 1320 instances with a CPU time of 289.1 seconds, while setting Default can only solve 1122 instances with a CPU time of 675.1 seconds. For unsolved instances, the average relative gap reduces from 7.0% to 0.1%. The main improvement comes from efficiency of the separation for the Benders optimality cuts since by Proposition 5.1, we know that before and after applying the SCNA method, the Benders

| Network | Default | SCNA | SCNA+INA |
|---------|---------|------|----------|
| MSG     | 0.01    | 80.9%| 8.9%     |
| (ρ = 31.5) | 0.05    | 56.4%| 0.2%     |
| GNU     | 0.01    | 93.1%| 85.9%    |
| (ρ = 7.4) | 0.05    | 72.3%| 33.9%    |
| ENRON   | 0.01    | 93.3%| 84.0%    |
| (ρ = 10.0) | 0.05   | 76.8%| 1.6%     |
| FACEBOOK | 0.01   | 79.1%| 0.3%     |
| (ρ = 32.4) | 0.05   | 47.8%| <0.1%    |
| DEEZER  | 0.01   | 84.4%| 65.4%    |
| (ρ = 18.3) | 0.05 | 50.6%| <0.1%    |
| TWITTER | 0.01   | 83.8%| 0.4%     |
| (ρ = 21.7) | 0.05 | 57.9%| <0.1%    |
| EPINIONS | 0.01 | 95.9%| 10.8%   |
| (ρ = 6.4) | 0.05 | 87.7%| 0.1%    |
| 1000 | 81.2%| <0.1%| 87.6% (6.4%) |

| Network | Δz_{SCNA} | Δr_{SCNA} | Δz_{SCNA} | Δr_{SCNA} | Δz_{SCNA} | Δr_{SCNA} |
|---------|------------|------------|------------|------------|------------|------------|
| MSG     | 1.22%  | 0.01%  | 2.40%  | <0.01%  | 2.67%  | <0.01%  |
| (ρ = 31.5) | 1.32%| 0.01%| 2.88%| 0.01%| 5.28%| 0.01%|
| GNU     | 8.32%  | 0.05%  | 4.65%  | <0.01%  | 5.85%  | <0.01%  |
| (ρ = 7.4) | 9.58%| 0.06%| 5.81%| 0.01%| 6.38%| <0.01%|
| ENRON   | 14.73% | 0.12%  | 5.21%  | <0.01%  | 5.85%  | <0.01%  |
| (ρ = 10.0) | 16.28%| 0.13%| 6.43%| 0.01%| 4.12%| <0.01%|
optimal cuts are identical, and hence the path of the search tree must be identical. The latter is further confirmed by the results of networks MSG and EPINIONS where the numbers of added Benders optimality cuts and branch-and-cut nodes are identical under settings Default and SCNA. Notice that it is reasonable to observe that the numbers of added Benders optimality cuts and branch-and-cut nodes in other networks are different since some instances cannot be solved within the time limit.

As for setting SCNA+INA, we observe that it slightly outperforms setting SCNA. In total, setting SCNA+INA can solve 11 more instances than setting SCNA, with the CPU time decreasing from 289.1 to 249.5 seconds. This is consistent with the former results in Table 2 in which only isomorphic nodes with small reachability sets can be detected by the INA and hence its contribution to speed up the solution procedure is not very large.

From the above results, we can conclude that for the IMP under the ICM, (i) the SCNA can effectively reduce the numbers of variables $z$ and elements in the reachability sets and the sizes of networks and hence is beneficial to solving the IMP especially when the network is large and well-connected; and (ii) the INA can further remove a fairly large fraction of variables $z$ from the SMCLP formulation and slightly speed up the solution procedure.

### 6.3 Results for the IMP under the LTM

In this subsection, we present similar computational results for the IMP under the LTM in Tables 5–7. To begin with, we note that for node $j$ in graph $G$, if it is a singleton node (i.e., it does not have any incoming arc), then in each live-arc graph $G^w$, it is also a singleton node; otherwise, it has at most one incoming arc in each live-arc graph $G^w$ (indeed, in the tested instances, it has exactly one incoming arc as $\sum_{(i,j) \in A} b_{ij} = 1$ holds; see Section 6.1). As a result, for any pair of nodes $i_1$ and $i_2$ in live-arc graph $G^w$, there exists at most one directed path from node $i_1$ to node $i_2$. This implies that in graph $G^w$, the subgraph induced by the nodes in SCC $SCC^w_i$ with $|SCC^w_i| \geq 2$ must be a single circle (which is in sharp contrast to the IMP under the ICM in which the subgraph induced by the nodes in SCC $SCC^w_i$ with $|SCC^w_i| \geq 2$ can be a union of multiple circles). This property for the IMP under the LTM, however, implies that only a small portion of nodes are likely to be strongly connected with other nodes, and the reachability sets of the nodes are also likely to be small. Consequently, through applying the SCNA, we can only observe a mild reduction on the numbers of nodes and arcs and the numbers of variables and elements in the reachability sets in Table 5. However, due to the small sizes of the reachability sets, more nodes are likely to be isomorphic among different scenarios and hence the INA can detect more reductions, as compared to the IMP under the ICM. This is shown in Table 5 in which we observe a fairly large reduction on the number of variables by applying the INA. The reduction on the number of elements in the reachability sets is relatively small which can be explained by the fact that most eliminated variables by the INA are associated with small reachability sets as the INA are associated with small reachability sets as the sum of elements in the reachability sets is relatively small which can be explained by the fact that most eliminated variables by the INA.

Therefore, the reductions by applying the SNA for the IMP under the LTM totally depend on the number of singleton nodes in network $G$, which is very small in most cases. Indeed, only network EPINIONS contains a relatively large percentage of singleton nodes (35.9%); see column $\Delta \bar{Z}$ under setting Default in Table 5. For the IMP under the LTM, with the increasing number of scenarios, a relatively large increase on the reductions by the INA can be observed, as shown in Table 6.

We now present the overall performance improvement of integrating the SCNA and INA into the BD algorithm in Table 7. Detailed statistics of these results can be found in Tables 1b-8b of [12]. From Table 7, we can see that the performance of
settings SCNA and SCNA+INA is slightly better than setting Default. Indeed, we only observe a minor improvement on the average CPU time (T), the number of solved instances (#S), and the average relative gap of the unsolved instances (Gap) through applying the SCNA and INA. This can be explained by the reasons that (i) the reduction on the sizes of networks through applying the SCNA is small (as shown in Table 5); (ii) the time spent in implementing the SCNA and INA is relative large (as shown in column PT in Table 7); and (iii) only isomorphic nodes with small reachability sets can be detected by the INA (as shown in column ΔR^\text{INA} under setting SCNA+INA in Table 5). In addition, we note from Table 7 that the IMPs under the LTM are generally much easier than those under the ICM. In total, under the LTM, only 16 among 480 instances (3.3%) cannot be solved by setting Default within the given time limit while 318 among 1440 instances (22.1%) cannot be solved by the same setting under the ICM.
6.4 | Selection of parameter MaxReacSize

As it has been mentioned in Section 5.1, MaxReacSize is a parameter to achieve a trade-off between the effectiveness and efficiency of implementing the INA: the larger the parameter MaxReacSize, the more isomorphic nodes that might be identified and the higher the computational complexity. Therefore, in this subsection, we compare the performance of different selections of parameter MaxReacSize. Tables 8 and 9 report the computational results for the IMP under the ICM and LTM, respectively. For simplicity, we only report the results for the case |Q| = 1000 (for the ICM, we only report the results for the case p = 0.1).

In the two tables, we use $m_0$ to represent the average memory consumption (in GB) of storing all reachability sets after removing those detected by the SCNA, and $M$ to denote the average memory consumption (in GB) of only storing the reachability sets with the size restriction. Instead of storing the reachability sets to obtain the required memories $M$, we calculate the total number of elements of the stored reachability sets and convert it to the needed memory size. In our experiments, we set the size restriction $MaxReacSize = 2, 4, 8, 1000$. Table 8 indicates that due to the limited memory, we were not able to construct the whole hashing table to implement the INA.

For the ICM, Table 8 shows that it requires a prohibitively large memory $m_0$ to store all the reachability sets. However, when restricting the size of the considered reachability sets to a small value of $MaxReacSize$, the memory overhead significantly reduces; see column $M$ in Table 8. In addition, with the increasing value of $MaxReacSize$, the improvement on the percentage of the eliminated variables $\Delta Z_{ \text{IINA} }$ becomes smaller and smaller. Indeed, for networks MSG, GNU, HEP, and ENRON, the proposed algorithm with $MaxReacSize = 8$ can identify almost all isomorphic nodes as those with $MaxReacSize = |V|$. For the other four networks, setting $MaxReacSize = 8$ enables to identify almost the same amount of isomorphic nodes as those obtained by setting $MaxReacSize = 1000$.

We now discuss the results for the IMP under the LTM in Table 9. On one hand, for the IMP under the LTM, since there exists at most one incoming arc for each node in each live-arc graph, the sizes of the reachability sets are likely to be smaller than those for the IMP under the ICM. This leads to a smaller total memory consumption $m_0$ and a larger memory consumption $M$ when restricting $MaxReacSize$ to a small value, as compared to those for the IMP under the ICM. As a result, the computational overhead of implementing the INA is very high, even for a relatively small $MaxReacSize$ (e.g., $MaxReacSize = 8$). In addition,
in analogy to the IMP under the ICM, with a small value of parameter \( \text{MaxReacSize} \), the proposed algorithm can identify almost the same amount of isomorphic nodes as the case \( \text{MaxReacSize} = |\mathcal{V}| \). Therefore, for the IMP under the LTM, we choose \( \text{MaxReacSize} = 4 \) in the implementation of the INA to achieve a trade-off between the performance and the time complexity.

7 | EXTENSIONS

In this section, we investigate a generalization of the IMP (3), which arises from many existing applications including the IMP and its variants [47, 48, 60, 61], and discuss the extensions of the proposed SCNA and INA to this generalization.

The considered generalization is of the form

\[
\max_{y, z} f(y, z) \quad \text{s.t. (3b), (3d), (3e),}
\]

\[
y \in \mathcal{V}, z \in \mathcal{Z},
\]

where \( f: \{0, 1\}^{|\mathcal{V}|} \times \{0, 1\}^{|\mathcal{Z}|} \rightarrow \mathbb{R} \), \( \mathcal{Y} \subseteq \mathbb{R}^{|\mathcal{V}|} \), and \( \mathcal{Z} \subseteq \mathbb{R}^{|\mathcal{Z}|} \). Similar to the IMP (3), problem (31) is built upon a finite number of live-arc graphs \( \mathcal{G}^\omega = (\mathcal{V}, \mathcal{A}^\omega), \omega \in \Omega \). However, in contrast to the IMP (3), problem (31) can flexibly allow any objective function and any constraint in sets \( \mathcal{Y} \) and \( \mathcal{Z} \). Indeed, the IMP (3) can be seen as a special case of problem (31) where \( f(y, z) = \sum_{\omega \in \Omega} p^\omega \sum_{i \in \mathcal{V}} z_i^\omega, \mathcal{Y} = \left\{ y \in \mathbb{R}^{|\mathcal{V}|} : \sum_{j \in \mathcal{V}} y_j \leq K \right\}, \text{ and } \mathcal{Z} = \mathbb{R}^{|\mathcal{Z}|} \). Due to the flexibility, various variants of the IMP can also be seen as special cases of (31). For instance, by choosing \( \mathcal{Y} = \left\{ y \in \mathbb{R}^{|\mathcal{V}|} : \sum_{j \in \mathcal{V}} c_j y_j \leq B \right\} \) and the same \( \mathcal{Z} \) and \( f(y, z) \) as that of the IMP, problem (31) reduces to the budgeted influence maximization problem (BIMP) studied in [48]. Here \( c_j, j \in \mathcal{V} \), is the cost of choosing node \( j \) as a seed node and \( B \) is the total budget. We next present another two special cases of problem (31).

- The seed minimization problem (SMP) [47]. In this problem, \( f(y, z) = -\sum_{j \in \mathcal{V}} y_j \), \( \mathcal{Y} = \mathbb{R}^{|\mathcal{V}|} \), and \( \mathcal{Z} = \left\{ z \in \mathbb{R}^{|\mathcal{Z}|} : \sum_{\omega \in \Omega} p^\omega \sum_{i \in \mathcal{V}} z_i^\omega \geq \right\} (D \in \mathbb{R}^+). \) This problem can be seen as a dual form of the IMP (3), which minimizes the number of seed nodes with an expected influence coverage threshold \( \mathcal{D} \) in a network.

- The seed minimization problem with probabilistic influence coverage guarantee (SMPPICG) [60, 61]. In this problem, \( f(y, z) = -\sum_{j \in \mathcal{V}} y_j \), \( \mathcal{Y} = \mathbb{R}^{|\mathcal{V}|} \), \( \mathcal{Z} = \text{Proj}(\mathcal{W}) \) where \( \mathcal{W} = \left\{ (z, \xi) \in \mathbb{R}^{|\mathcal{V}|} \times \{0, 1\}^{|\mathcal{Z}|} : \sum_{i \in \mathcal{V}} z_i^\omega \geq D_i, \forall \omega \in \Omega, \sum_{\omega \in \Omega} \xi_i^\omega \geq 1 - \varepsilon \right\} \) (\( \varepsilon \in (0, 1) \) is the confidence level).

Instead of ensuring an expected influence coverage threshold \( \mathcal{D} \), the problem requires to influence at least \( \mathcal{D} \) nodes with a probability at least \( 1 - \varepsilon \).

We next discuss the extensions of the proposed SCNA and INA to problem (31). To proceed, we need the following two realistic assumptions.

(i) \( f(y, z) \) is nondecreasing with respect to variables \( z \), i.e., if \( z^1, z^2 \in \mathcal{Z} \) and \( z^1 \leq z^2 \), then \( f(y, z^1) \leq f(y, z^2) \).

(ii) \( \mathcal{Z} \) is up-monotone, that is, if \( z^1 \in \mathcal{Z} \) and \( z^1 \leq z^2 \), then \( z^2 \in \mathcal{Z} \) as well (such a set is also called a reverse normal set [58]).

The two assumptions imply that when node \( i \) is reachable in scenario \( \omega \) from some seed nodes (i.e., \( \sum_{j \in \mathcal{R}(G^\omega, i_j)} y_j \geq 1 \)), activating node \( i \) in scenario \( \omega \) (i.e., setting \( z_i^\omega := 1 \) provides a better solution for the decision maker while does not violate his/her requirement. It can be easily verified that for the IMP, BIMP, SMP, and SMPPICG, the two assumptions are satisfied.

**Proposition 7.1.** Suppose that problem (31), with assumptions (i) and (ii), has an optimal solution. Then there must exist an optimal solution \((\bar{y}, \bar{z})\) such that (8) holds.

**Proof.** Let \((\bar{y}, \bar{z})\) be an optimal solution of problem (31). Suppose that (8) does not hold for some \( \omega_0 \in \Omega \) and \( i_0 \in \mathcal{V} \). Then we must have \( z_i^\omega = 0 \) and \( \sum_{j \in \mathcal{R}(G^\omega, i_0, j_j)} y_j \geq 1 \). Setting \( z_i^\omega := 1 \), we obtain a new point \((\bar{y}, \bar{z}')\). By assumption (ii), \( \bar{z}' \in \mathcal{Z} \), and by \( \sum_{j \in \mathcal{R}(G^\omega, i_0, j_j)} y_j \geq 1 \), constraints (3b) hold at point \((\bar{y}, \bar{z}')\). This implies that \((\bar{y}, \bar{z}')\) is a feasible solution of problem (31). Moreover, by assumption (i), \( f(y, z') \geq f(y, z) \), indicating that \((\bar{y}, \bar{z}')\) must also be an optimal solution of problem (31). Recursively using the above argument, the statement follows.

By Proposition 7.1, if \( R(G^\omega, i_1) = R(G^\omega, i_2) \), we can set \( z_i^\omega := z_i^\omega \) in problem (31) (with the two realistic assumptions (i) and (ii)). As a result, the proposed SCNA and INA can also be applied to problem (31) to reduce the problem size and improve the solution efficiency.
8 | CONCLUDING REMARKS

In this paper, we proposed two new presolving methods, called the SCNA and INA, and integrated them into the BD algorithm to solve the IMP. The SCNA enables to build an SMCLP formulation for the considered problem based on the (potentially) much more compact live-arc graphs, which are obtained by aggregating strongly connected nodes in the original live-arc graphs. The INA further reduces the problem size of the SMCLP formulation by aggregating isomorphic nodes among different live-arc graphs. We provided a theoretical analysis on two special cases of the IMP to show the strength of the proposed SCNA and INA in reducing the problem size of the SMCLP formulation. Furthermore, with the SCNA and INA, a (potentially) much faster separation procedure for the Benders optimality cuts is developed, which plays a crucial role in speeding up the BD algorithm. We have performed extensive experiments to analyze the performance impact of the proposed SCNA and INA on solving the IMP with real-world networks. Computational results show that the proposed SCNA and INA can effectively reduce the problem size, speed up the separation of Benders optimality cuts, and (hence) improve the overall performance of using the BD algorithm to solve the IMP. We also studied a generalization of the IMP and demonstrated that the proposed SCNA and INA are applicable to this generalization under some realistic assumptions.

There still exist some instances where the proposed SCNA and INA cannot effectively reduce the problem size of the SMCLP formulation. Indeed, in Section 4 of [12], we have provided a worst-case example showing that the percentage of the eliminated variables in the SMCLP formulation of the IMP tends to zero with a probability tending to one. Consequently, it is interesting to develop more powerful presolving methods for solving the IMP. In addition, it also deserves to investigate whether the proposed presolving methods are computationally effective in solving other variants of the IMP.

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DATA AVAILABILITY STATEMENT

The datasets generated or analyzed during this study are available from the corresponding author on reasonable request.

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APPENDIX A: PROOF OF PROPOSITION 4.3

Proof. Clearly, if \( K \geq |\mathcal{M}| + |\mathcal{N}| \), point \( (y, z) = (e, e) \) is optimal for formulation (14) and its LP relaxation, where \( e \) is an all-ones vector with appropriate dimension. As a result, the statement follows. Therefore, in the following, we consider the case \( K < |\mathcal{M}| + |\mathcal{N}| \).

Let \((\bar{y}, \bar{z})\) be an optimal solution of the LP relaxation of formulation (14). If there exists some \( i_0 \in \mathcal{M} \) and \( j_0 \in \mathcal{N} \) such that \( y_{i_0} < 1 \) and \( z_{j_0} > 0 \), then we can construct a new point \((\hat{y}, \hat{z})\) as follows:

- \( \hat{y}_{i_0} := y_{i_0} + \varepsilon, \hat{y}_i := y_i - \varepsilon, \) and \( \hat{z}_j := y_j \) for \( i \in \mathcal{M} \cup \mathcal{N} \setminus \{i_0, j_0\} \);
- \( \hat{z}_{j_0} := \max\{z_{j_0} - \varepsilon, 0\} \) for \( (i, j) \in \mathcal{A}' \) and \( \hat{z}_j := z_j \) for \( (i, j) \in \mathcal{A}' \) with \( j \neq j_0 \),

where \( \varepsilon > 0 \) is a sufficiently small value. It is easy to see that point \((\hat{y}, \hat{z})\) is also feasible for the LP relaxation of formulation (14). Moreover, point \((\hat{y}, \hat{z})\) must be optimal since

\[
\sum_{i \in \mathcal{M} \cup \mathcal{N}} s_i \hat{y}_i + \sum_{(i, j) \in \mathcal{A}'} c_{ij} \hat{z}_{ij} - \left( \sum_{i \in \mathcal{M} \cup \mathcal{N}} s_i \bar{y}_i + \sum_{(i, j) \in \mathcal{A}'} c_{ij} \bar{z}_{ij} \right) = s_i \bar{y}_i + \varepsilon + s_j (\bar{y}_j - \varepsilon) + \sum_{i : (i, j) \in \mathcal{A}'} c_{ij} \max\{z_{j_0} - \varepsilon, 0\} - s_i \bar{y}_i - s_j \bar{y}_j - \sum_{i : (i, j) \in \mathcal{A}'} c_{ij} \bar{z}_{ij} \leq s_i \varepsilon - s_j \varepsilon + \sum_{i : (i, j) \in \mathcal{A}'} c_{ij} \max\{-\varepsilon, -\bar{z}_{ij}\} \geq s_i \varepsilon - s_j \varepsilon - \sum_{i : (i, j) \in \mathcal{A}'} c_{ij} \varepsilon = \varepsilon \left( s_i - s_j - \sum_{i : (i, j) \in \mathcal{A}'} c_{ij} \right) = 0,
\]

where the last equality follows from Remark 4.2. Recursively applying the above argument, we will obtain an optimal solution \((\bar{y}, \bar{z})\) of the LP relaxation of formulation (14) fulfilling:

\[(\star) \text{ if } \bar{y}_{i_0} > 0 \text{ for some } j_0 \in \mathcal{N}, \text{ then } \bar{y}_i = 1 \text{ for all } i \in \mathcal{M} \text{ must hold.}\]

Furthermore, we can, without loss of generality, assume the followings on point \((\bar{y}, \bar{z})\).

1. \( \sum_{i \in \mathcal{M} \cup \mathcal{N}} \bar{y}_i = K. \) Otherwise, we can increase some \( \bar{y}_i \), with \( \bar{y}_i < 1 \), without decreasing the objective value (as \( \sum_{i \in \mathcal{M} \cup \mathcal{N}} \bar{y}_i < K < |\mathcal{M}| + |\mathcal{N}| \));
2. \( \bar{z}_{ij} = \min\{\bar{y}_i + \bar{y}_j, 1\} \) for all \((i, j) \in \mathcal{A}'\). Otherwise, we can increase \( \bar{z}_{ij} \), with \( \bar{z}_{ij} < \min\{\bar{y}_i + \bar{y}_j, 1\} \), without decreasing the objective value.

Together with \((\star)\) and 1), point \((\bar{y}, \bar{z})\) must satisfy the followings.
(i) if $K \geq |\mathcal{M}|$, then $\bar{y}_i = 1$ for all $i \in \mathcal{M}$; and
(ii) if $K < |\mathcal{M}|$, then $\bar{y}_j = 0$ for all $j \in \mathcal{N}$.

We next prove the statement in the proposition by treating cases (i) and (ii) separately.

(i) In this case, $\bar{y}_i = 1$ for all $i \in \mathcal{M}$ and by 2), $\overline{z}_{ij} = 1$ for all $(i,j) \in \mathcal{A}'$. Setting $y_i : = 1$ for all $i \in \mathcal{M}$ and $z_{ij} : = 1$ for all $(i,j) \in \mathcal{A}'$. the LP relaxation of formulation (14) reduces to

$$\max_y \left\{ \sum_{j \in \mathcal{N}} s_j y_j + |\mathcal{M}| + \sum_{(i,j) \in \mathcal{A}'} c_{ij} : \sum_{j \in \mathcal{N}} y_j = K - |\mathcal{M}|, \ y_j \in [0, 1], \ \forall j \in \mathcal{N} \right\}. \tag{A1}$$

Then point $\{\bar{y}_j\}_{j \in \mathcal{N}}$ must be an optimal solution of formulation (A1). On the other hand, suppose that $s_j \geq \cdots \geq s_{j_{|N|}}$ where $\{j_1, \ldots, j_{|N|}\} = \mathcal{N}$. It is easy to show that point $\{|y'_j|\}_{j \in \mathcal{N}}$ is also an optimal solution of formulation (A1) where $y'_{j_i} = 1$ for $i = 1, \ldots, K - |\mathcal{M}|$ and $y'_j = 0$ otherwise. We next extend point $\{|y'_j|\}_{j \in \mathcal{N}}$ to a higher dimensional point $(\bar{y}', \bar{z}') \in \{0, 1\}^{|\mathcal{M}| + |\mathcal{N}|} \times \{0, 1\}^{|\mathcal{A}'|}$ by additionally setting $y'_j : = 1$ for all $i \in \mathcal{M}$ and $z'_{ij} : = 1$ for all $(i,j) \in \mathcal{A}'$. The 0-1 point $(\bar{y}', \bar{z}')$ must be optimal to formulation (14) and its LP relaxation. This implies that the LP relaxation of formulation (14) is tight and formulation (14) is strongly polynomial time solvable for this case.

(ii) In this case, $\bar{y}_j = 0$ for all $j \in \mathcal{N}$ and by 2), $\overline{z}_{ij} = \bar{y}_i$ for all $(i,j) \in \mathcal{A}'$. Setting $y_i : = 0$ for all $i \in \mathcal{N}$ and $z_{ij} : = \bar{y}_i$ for all $(i,j) \in \mathcal{A}'$, the LP relaxation of formulation (14) reduces to

$$\max_y \left\{ \sum_{i \in \mathcal{M}} \left( 1 + \sum_{j : (i,j) \in \mathcal{A}'} c_{ij} \right) y_i : \sum_{i \in \mathcal{M}} y_i = K, \ y_i \in [0, 1], \ \forall i \in \mathcal{M} \right\}. \tag{A2}$$

Then point $\{\overline{y}_i\}_{i \in \mathcal{M}}$ must be an optimal solution of formulation (A2). Using a similar argument in case (i), we can also prove the statement in this case.

**APPENDIX B: PROOF OF PROPOSITION 5.1**

**Proof.** We first prove case (i). Note that for all nodes in a given SCC $SC_{\omega}^w$ of $\mathcal{G}^\omega$ ($u \in \overline{V}^\omega$), their reachability sets are identical and equal to $R(\mathcal{G}^\omega, SC_{\omega}^w)$. As a result, $(\bar{\alpha}_u^\omega, \bar{\beta}_u^\omega)$ for all $i \in SC_{\omega}^w$ must also be identical, denoted by $(\bar{\alpha}_u^\omega, \bar{\beta}_u^\omega)$. Hence, inequality (25) can be rewritten as

$$q_u^\omega \leq \sum_{u \in \overline{V}^\omega} \left| SC_{\omega}^w \right| \bar{\alpha}_u^\omega \sum_{j \in R(\mathcal{G}^\omega, SC_{\omega}^w)} y_j + \left| SC_{\omega}^w \right| \bar{\beta}_u^\omega$$

where the equality follows from (4). By (4) and the definitions of $(\bar{\alpha}_u^\omega, \bar{\beta}_u^\omega)$ and $(\bar{\alpha}_u^\eta, \bar{\beta}_u^\eta)$ in (27) and (28), we must have $(\bar{\alpha}_u^\omega, \bar{\beta}_u^\omega) = (\left| SC_{\omega}^w \right| \bar{\alpha}_u^\omega, \left| SC_{\omega}^w \right| \bar{\beta}_u^\omega)$ for all $u \in \overline{V}^\omega$. Thus, the two Benders optimality cuts (26) and (B1) are equivalent.

We next prove case (ii). Indeed, for two different scenarios $\omega$ and $\eta$, the corresponding Benders optimality cuts based on formulation (6) are given by

$$q_u^\omega \leq \sum_{u \in \overline{V}^\omega} \left( \bar{\alpha}_u^\omega \sum_{v \in R(\mathcal{G}^\omega, u)} y(SC_{\omega}^v) + \bar{\beta}_u^\omega \right)$$

$$q_u^\eta \leq \sum_{u \in \overline{V}^\eta} \left( \bar{\alpha}_u^\eta \sum_{v \in R(\mathcal{G}^\eta, u)} y(SC_{\omega}^v) + \bar{\beta}_u^\eta \right).$$

Suppose that node $u_1$ in graph $\mathcal{G}^\omega$ is isomorphic to node $u_2$ in graph $\mathcal{G}^\eta$. By applying the INA, we can remove, for example, variable $z_{u_1}^w$ and the corresponding reachability constraint in (5) from formulation (6). The new objective coefficient $f_{u_1}^\omega$ is set to the sum of the old objective coefficients of variables $z_{u_1}^w$ and $z_{u_1}^\eta$. As a result, the term $\bar{\alpha}_u^\omega \sum_{v \in R(\mathcal{G}^\omega, u)} y(SC_{\omega}^v) + \bar{\beta}_u^\omega$ in Benders optimality cut (B2) will be incorporated into Benders optimality cut (B3),
leading to two new Benders optimality cuts

\[ \varphi^\omega \leq \sum_{u \in \overline{V} \setminus \{u_1\}} \left( \alpha_u^\omega \sum_{v \in \hat{\mathcal{R}}_u} y(SC_v^\omega) + \overline{\beta}_u^\omega \right) \quad \text{and} \quad \tag{B4} \]

\[ \varphi^\eta \leq \sum_{u \in \overline{V}} \left( \alpha_u^\eta \sum_{v \in \hat{\mathcal{R}}_u} y(SC_v^\eta) + \overline{\beta}_u^\eta \right) + \left( \overline{\alpha}_{u_1}^\omega \sum_{v \in \hat{\mathcal{R}}_{u_1}} y(SC_v^\omega) + \overline{\beta}_{u_1}^\omega \right). \quad \tag{B5} \]

Obviously, the new Benders optimality cuts (B4) and (B5) may be different from the old ones (B2) and (B3), respectively.