

# Scaling Biclique Percolation for Community Detection in Large Bipartite Networks

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## Abstract

Community detection, which uncovers closely connected vertex groups in networks, is vital for applications in social networks, recommendation systems, and beyond. Real-world networks often have bipartite structures (vertices in two disjoint sets with inter-set connections), creating unique challenges on specialized community detection methods. Biclique percolation community (BCPC) is widely used to detect cohesive structures in bipartite graphs. A biclique is a complete bipartite subgraph, and a BCPC forms when maximal bicliques connect via adjacency (sharing an  $(\alpha, \beta)$ -biclique). Yet, existing methods for BCPC detection suffer from high time complexity due to the potentially massive maximal biclique adjacency graph (MBAG). To tackle this, we propose a novel partial-BCPC based solution, whose key idea is to use partial-BCPC to reduce the size of the MBAG. A partial-BCPC is a subset of BCPC. Maximal bicliques belonging to the same partial-BCPC must also belong to the same BCPC. Therefore, these maximal bicliques can be grouped as a single vertex in the MBAG, significantly reducing the size of the MBAG. Furthermore, we move beyond the limitations of MBAG and propose a novel BCPC detection approach based on  $(\alpha, \beta)$ -biclique enumeration. This approach detects BCPC by enumerating all  $(\alpha, \beta)$ -bicliques and connecting maximal bicliques sharing the same  $(\alpha, \beta)$ -biclique, which is the condition for maximal bicliques to be adjacent. It also leverages partial-BCPC to significantly prune the enumeration space of  $(\alpha, \beta)$ -biclique. Experiments show that our methods outperform existing methods by nearly three orders of magnitude.

## 1 Introduction

Community detection aims to uncover groups of closely connected vertices in a network and is a core task in network analysis [11]. It plays a key role in applications such as social networks [26], biological systems [25] and recommendation systems [8]. In many real-world scenarios, networks are naturally modeled as bipartite graphs, represented by  $G = (U, V, E)$ , where vertices are divided into two disjoint sets  $U$  and  $V$ , and connections only occur between  $U$  and  $V$ . Representative examples include user-item networks in recommendation systems [23] and author-paper networks in bibliometrics [5]. This structure introduces new challenges for community detection, motivating the development of specialized methods for bipartite graphs.

The biclique percolation community (BCPC) is a widely studied approach for detecting cohesive structures in bipartite graphs [7, 12–16, 33]. A biclique is a complete bipartite subgraph, and

a biclique percolation community is defined as a maximal set of bicliques that are connected through *adjacency*. In the  $(\alpha, \beta)$ -biclique percolation community, two bicliques are considered *adjacent* if they share at least  $\alpha$  vertices on one side and  $\beta$  on the other (share an  $(\alpha, \beta)$ -biclique). Biclique percolation community (BCPC) has diverse applications across bipartite network scenarios. In social networks (users-entities as vertices, interaction as edges), BCPC uncovers interest-based user groups to support interest or user recommendation [12]. In Wikipedia networks (articles-editors as vertices, editorial activities as edges), BCPC identifies clusters, revealing topic-driven editor aggregation and coordinated content creation [14]. For internet resource access (resources-visitors as vertices), BCPC is used to analyze student-course resource relationships and their temporal evolution [13, 33]. In enterprise heterogeneous graphs (hosts-users as vertices, interactions as edges), BCPC is used to detect clusters to aid abnormal user behavior identification and malicious activity tracing for network security [16]. BCPC allows overlaps between clusters, and this property is crucial because real-world users or entities are likely to belong to multiple communities. This property has also been extensively studied [12–14, 16].

Despite its widespread practical utility, the BCPC method faces significant computational challenges due to its high time complexity. To the best of our knowledge, the existing BCPC methods are all based on [15]. Its framework was later implemented in [27] and used to build indexes for personalized search problems [7]. The basic idea is based on maximal biclique adjacency graph (MBAG). This method first enumerates all maximal bicliques and then uses them as vertices to construct a maximal biclique adjacency graph. In this graph, there is an edge between two vertices if and only if the overlapping part of the corresponding two maximal bicliques contains an  $(\alpha, \beta)$ -biclique. The connected components in the adjacency graph correspond one-to-one with the  $(\alpha, \beta)$ -BCPC. However, real-world bipartite graphs may contain a large number of maximal bicliques (up to  $2^{n/2}$  theoretically [10]), and the adjacency graph formed by these maximal bicliques could contain a significant number of edges (up to  $10^{11}$ , see Table 1). Traversing such a large-scale adjacency graph is highly time-consuming.

To overcome the limitations of existing methods, we first propose a partial-BCPC based solution, which focuses on reducing the size of the MBAG. The key idea is to leverage the prefixes of the maximal biclique enumeration tree during the enumeration process. Specifically, we introduce a novel concept of partial-BCPC, which is computed on-the-fly while enumerating maximal bicliques. During the enumeration, if a prefix of the enumeration tree contains at least  $\alpha$  vertices from the vertex set  $U$  and  $\beta$  vertices from the vertex set  $V$ , we can determine that all maximal bicliques under this branch belong to the same BCPC. These bicliques collectively form an incomplete BCPC, referred to as a partial-BCPC. Maximal bicliques in the same partial-BCPC can be regarded as a single vertex in the MBAG, thus reducing the size of the MBAG. Experiments show that our approach achieves a substantial reduction in the number of vertices in the MBAG —by two orders of magnitude (see

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Figure 5 (c) of Exp-3)—thereby enabling more efficient traversal and computation of BCPCs.

While the partial-BCPC solution operates on a reduced MBAG, it still requires a full traversal of this structure, which can remain prohibitively large. To overcome this limitation, we propose a novel approach based on  $(\alpha, \beta)$ -biclique enumeration. The basic idea is to enumerate all  $(\alpha, \beta)$ -biclques in the bipartite graph, and for each enumerated  $(\alpha, \beta)$ -biclque  $B$ , connect all maximal biclques containing  $B$  using union-find set. The reason this idea works is that two maximal biclques are adjacent only if they share an  $(\alpha, \beta)$ -biclque. Since the number of  $(\alpha, \beta)$ -biclques increases exponentially with respect to  $\alpha, \beta$  [30, 31], we leverage partial-BCPC once again to prune the  $(\alpha, \beta)$ -biclque enumeration tree. During the enumeration process, for each intermediate small biclque generated, we can maintain the set of maximal biclques containing it. If this set is empty or all the maximal biclques in the set already belong to the same partial-BCPC, then the current enumeration branch can be safely pruned. Experiments show that using partial-BCPC can reduce the number of nodes in the  $(\alpha, \beta)$ -biclque enumeration tree by more than five orders of magnitude (Exp-4). Consequently, it achieves a speedup of nearly three orders of magnitude compared to the state-of-the-art (SOTA) algorithm (Exp-1).

We summarize our contributions as follows.

**Novel partial-BCPC based solution.** We propose a novel partial-BCPC based solution, characterized by introducing a relaxed definition of BCPC, namely partial-BCPC. By leveraging partial-BCPC, we can significantly reduce the size of the MBAG used in existing methods, thereby directly accelerating the traversal process of the MBAG. We show that leveraging partial-BCPC can reduce the number of vertices in the MBAG by up to two orders of magnitude (i.e., eliminating up to 99% of the vertices in MBAG, see Exp-3).

**Novel  $(\alpha, \beta)$ -biclque based solution.** Unlike traditional approaches that rely on traversing the MBAG, we propose a novel framework that directly enumerates  $(\alpha, \beta)$ -biclques and connects the maximal biclques that share them utilizing union-find sets. To reduce the cost of enumerating  $(\alpha, \beta)$ -biclques, we again leverage partial-BCPC, which reduces the number of nodes in the enumeration tree by up to five orders of magnitude (Exp-4).

**Extensive experiments.** We conduct experiments on 10 real-world datasets, and the experimental results demonstrate the effectiveness and efficiency of our proposed algorithms. Specifically, we test the performance of our algorithms under a wide range of parameter settings. For example, on dataset Youtube, when  $\alpha = 2, \beta = 2$ , the SOTA method (MBAG) takes 13,376 seconds, while the partial-BCPC based solution (PBCPC+) takes 197 seconds, achieving an improvement of nearly two orders of magnitude. In contrast, the  $(\alpha, \beta)$ -biclque based solution (BiclqueP) takes 34 seconds, with an improvement of nearly three orders of magnitude. Additionally, we conduct a case study of BCPC in community detection. The results show that compared with other traditional community models (e.g., biclique, bitruss, bicore), BCPC can find more practically meaningful communities, which demonstrates the effectiveness of BCPC.

**Reproducibility.** The source code of this paper is released on <https://anonymous.4open.science/r/bcpc> for reproducibility purposes.

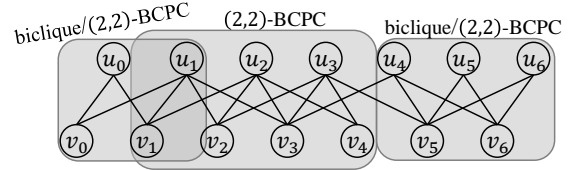


Figure 1: A bipartite graph  $G$  and  $(\alpha, \beta)$ -BCPC ( $\alpha = 2, \beta = 2$ )

## 2 Preliminaries

Let  $G = (U, V, E)$  be an undirected and unweighted bipartite graph, where  $U, V$  are two disjoint sets of vertices and  $E$  is a set of edges in form of  $(u, v), u \in U, v \in V$ . For a vertex  $u \in U$ ,  $Nei_G(u)$  is the set of neighbors of  $u$ , i.e.,  $Nei_G(u) = \{v | (u, v) \in E\}$ . A bipartite graph  $G' = (U', V', E')$  is a subgraph of  $G$  if  $U' \subseteq U, V' \subseteq V, E' \subseteq E$ .

**DEFINITION 1 (BICLIQUE).** Given a bipartite graph  $G = (U, V, E)$ , a biclique is a pair of vertex sets  $B = (X, Y)$  where  $X \subseteq U, Y \subseteq V$  and  $\forall u \in X, v \in Y, (u, v) \in E$ .

For simplicity, we let  $B.X = X, B.Y = Y$  if  $B = (X, Y)$ . For two biclques  $B, B'$ ,  $B \subseteq B'$  indicates that  $B.X \subseteq B'.X, B.Y \subseteq B'.Y$ . A maximal biclique  $B$  is a biclique that there is no other biclique  $B'$  satisfying  $B \subseteq B'$ .

**DEFINITION 2 ( $(\alpha, \beta)$ -BICLIQUE ADJACENCY).** Given two integer  $\alpha, \beta$  and two biclques  $B_1, B_2$ ,  $B_1$  and  $B_2$  are  $(\alpha, \beta)$ -adjacent if  $|B_1.X \cap B_2.X| \geq \alpha, |B_1.Y \cap B_2.Y| \geq \beta$ .

**DEFINITION 3 ( $(\alpha, \beta)$ -BICLIQUE CONNECTIVITY).** Given two integer  $\alpha, \beta$  and two biclques  $B_1, B_n$ ,  $B_1$  and  $B_n$  are  $(\alpha, \beta)$ -connected if there is a sequence of biclques  $B_1, B_2, \dots, B_n$  that  $\forall i = 1, \dots, n-1, B_i$  and  $B_{i+1}$  are  $(\alpha, \beta)$ -adjacent.

**DEFINITION 4 ( $(\alpha, \beta)$ -BICLIQUE PERCOLATION COMMUNITY (( $\alpha, \beta$ )-BCPC)).** Given a bipartite graph  $G$  and two integer  $\alpha, \beta$ , an  $(\alpha, \beta)$ -biclque percolation community is a maximal set of maximal biclques in  $G$  that any two maximal biclques in the set are  $(\alpha, \beta)$ -connected.

**EXAMPLE 1.** Let  $\alpha = 2, \beta = 2$ , Figure 1 illustrates a bipartite graph  $G$  and some of the  $(2, 2)$ -BCPCs. These BCPCs locates on the left, in the middle, and on the right, respectively. The  $(2, 2)$ -BCPC in the middle is composed of three maximal biclques:  $B_1 = (\{u_1, u_2\}, \{v_1, v_2, v_3\})$ ,  $B_2 = (\{u_1, u_2, u_3\}, \{v_2, v_3\})$ , and  $B_3 = (\{u_2, u_3\}, \{v_2, v_3, v_4\})$ . Among these biclques,  $B_1$  and  $B_2$  are adjacent because their intersection,  $(\{u_1, u_2\}, \{v_2, v_3\})$ , satisfies the condition in Definition 2. For the same reason,  $B_2$  and  $B_3$  are also adjacent. Therefore,  $B_1, B_2$ , and  $B_3$  can together form a valid  $(2, 2)$ -BCPC. On the left and right sides of Figure 1 are two maximal biclques. These maximal biclques are not adjacent (Definition 2) to any other maximal biclques, and thus can be regarded as two special BCPCs.

**Problem statement.** Given a bipartite graph  $G$  and two integer  $\alpha, \beta$ , the goal of our work is to detect all  $(\alpha, \beta)$ -BCPCs (BCPC detection).

### 2.1 Maximal Biclque Enumeration

The existing state-of-the-art maximal biclque enumeration methods [1, 6, 10] are all based on the set-enumeration tree structure, where vertices are iteratively traversed to build nodes of the enumeration tree, with each node representing a distinct biclque. In this paper, we adopts the algorithmic framework in [10] to

The existing state-of-the-art  $(\alpha, \beta)$ -biclique enumeration method is proposed in [30]. Given a bipartite graph  $G = (U, V, E)$ , the core idea is to enumerate combinations of vertices in the  $U$  set while maintaining their common neighbors in the  $V$  set. It also constructs an ordinary directed graph, called the *2-hop graph*, on the vertices of the  $U$  set to accelerate the enumeration process.

**Algorithm 2:** The  $(\alpha, \beta)$ -biclique enumeration framework in [30]

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**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All  $(\alpha, \beta)$ -bicliques in  $G$

```

1 Construct 2-hop graph  $\tilde{H} = (U, \tilde{E}_H)$  on vertex set  $U$  [30];  $\tilde{H} = (U, \tilde{E}_H)$ : a direct
  graph,  $(u, v) \in \tilde{E}_H$  indicates that  $u < v$ ,  $u, v$  have common neighbors in  $V$  */
2 BicliqueListing( $\tilde{H}, \emptyset, V$ );
3 Procedure BicliqueListing( $\tilde{H}, R_U, C_V$ )
4 if  $|R_U| = \alpha$  then
5   foreach  $R_V \subseteq C_V, |R_V| = \beta$  do Output  $(R_U, R_V)$  as an  $(\alpha, \beta)$ -biclique;
6   return;
7 foreach  $u \in \tilde{H}$  do
8    $C'_V \leftarrow C_V \cap \text{Nei}_{\tilde{H}}(u)$ ;
9   if  $|C'_V| < \beta$  or  $\text{Nei}_{\tilde{H}}(u) < \alpha - |R_U| - 1$  then continue;
10  Let  $\tilde{H}'$  be induced subgraph of  $\tilde{H}$  on  $\text{Nei}_{\tilde{H}}(u)$ ;
11  BicliqueListing( $\tilde{H}', R_U \cup \{u\}, C'_V$ );

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Algorithm 2 presents the basic framework. It first constructs the 2-hop graph  $\tilde{H}$  with  $U$  as its vertex set (Line 1), where an edge  $(u, v)$  exists if  $u < v$  and  $u, v$  share at least one common neighbor in set  $V$  of the bipartite graph. In Line 4, if  $R_U$  already contains  $\alpha$  vertices, it is sufficient to enumerate all combinations of  $\beta$  vertices from  $C_V$  to obtain the  $(\alpha, \beta)$ -bicliques. In Line 7, the algorithm enumerates vertices  $u$  in  $\tilde{H}$ , then in Line 8 updates the candidate set  $C'_V$  in  $V$ , and proceeds with the next-level enumeration in Line 11.

### 2.3 Union-Find Set [9]

The Union-Find Set, also referred to as the Disjoint-Set data structure, serves as an efficient method for managing a collection of disjoint sets. Each set is represented as a tree, where the nodes represent the elements of the set. Within this tree, every node points to its parent, with the root of the tree acting as the representative of the set. This tree structure allows for fast **union** and **find** operations, making the Union-Find Set well-suited for tasks such as graph connectivity problems. **Find**( $i$ ) is used to get the representative of the unique set containing  $i$ , **union**( $i, j$ ) is used to merge the two sets containing  $i$  and  $j$  into their union. Specifically, within the corresponding tree structure, the representative node of one set is assigned as the parent of the representative node of another set.

In all practical scenarios, the **union** and **find** operations of the union-find set achieve constant amortized time complexity. The space complexity of this structure is  $O(n)$ , with  $n$  denoting the total number of elements it manages [9].

### 3 Existing Solution

The existing BCPC detection method was proposed and implemented in [7, 15, 27]. To the best of our knowledge, no other work has primarily focused on the algorithmic implementation and improvement of BCPC detection. The basic idea is based on maximal\_biclique\_adjacency\_graph (MBAG). This method first enumerates all maximal bicliques and then uses them as vertices to construct a maximal biclique adjacency graph. In this graph, there is an edge between two vertices if and only if the overlapping part of the corresponding two maximal bicliques contains an  $(\alpha, \beta)$ -biclique (satisfying Definition 2). The connected components in the adjacency graph correspond one-to-one with the  $(\alpha, \beta)$ -BCPC. In this method, the  $(\alpha, \beta)$ -biclique connectivity of maximal bicliques in  $(\alpha, \beta)$ -BCPC are maintained by the classic union-find set [9].

**Algorithm 3:** Existing MBAG-based solution

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**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All BCPC in  $G$

```

1  $\mathcal{B} \leftarrow$  set of all maximal bicliques;
2  $\mathcal{B} \leftarrow \{B | B \in \mathcal{B}, |B.X| \geq \alpha, |B.Y| \geq \beta\}$ ;
3  $\text{Nei}(B) \leftarrow$  set of maximal bicliques that share vertices with  $B$ ;
4  $UF \leftarrow$  an initial union-find set, i.e.,  $UF.find(B) = B$  for any input  $B$ ;
5 for  $B \in \mathcal{B}$  do
6   for  $B' \in \text{Nei}(B)$  do
7     if  $|B.X \cap B'.X| \geq \alpha$  and  $|B.Y \cap B'.Y| \geq \beta$  then  $UF.union(B, B')$ ;
8 return  $UF$ ;

```

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Algorithm 3 is the existing MBAG-based algorithm identifying all  $(\alpha, \beta)$ -BCPCs in a bipartite graph  $G = (U, V, E)$ . It starts by enumerating all maximal bicliques in  $G$  and filters them based on size constraints (Lines 1-2). The reason is that for a maximal biclique  $B$ , if  $|B.X| < \alpha$  or  $|B.Y| < \beta$ , then  $B$  can not form a BCPC with other maximal bicliques. A union-find set  $UF$  is initialized in Line 4. The algorithm then iterates over each maximal biclique  $B$  and its neighbors  $B' \in \text{Nei}(B)$ , checking if  $|B.X \cap B'.X| \geq \alpha$ ,  $|B.Y \cap B'.Y| \geq \beta$  (Lines 5-7). If the conditions are satisfied,  $B$  and  $B'$  are merged into the same BCPC by  $UF.union(*)$  (Line 7). After processing all maximal bicliques, the union-find set  $UF$  represents the final BCPCs in the bipartite graph.

The major drawback of this approach is that the size of the MBAG can be extremely large (up to  $10^{11}$ , see Table 1). Consequently, traversing such a massive MBAG to derive all BCPCs is clearly impractical.

### 4 Novel Partial-BCPC Based Solution

As we discussed in Section 3, current approach relying on MBAG (Algorithm 3) faces significant challenges due to the vast scale of MBAG. This immense size leads to inefficient traversal operations. Consequently, it becomes evident that optimizing the performance of MBAG-based method hinges on effectively reducing the size of the MBAG.

**Key idea.** We reduce the size of the MBAG by merging vertices within it. The key is to leverage the prefixes of the maximal biclique enumeration tree during the enumeration process. During the enumeration, if a prefix of the enumeration tree contains at least  $\alpha$  vertices in  $U$  and  $\beta$  vertices in  $V$ , we can determine that all maximal bicliques under this branch belong to the same BCPC. Connections between these maximal bicliques are unnecessary to be traversed. These bicliques collectively form an incomplete BCPC, referred to as a partial-BCPC.

**EXAMPLE 3.** Let  $\alpha = \beta = 1$ . Consider Figure 2, where the subtree rooted at node  $nd(B_1, h_2)$  contains three maximal biclique branches:  $B_1$ ,  $B_2$ , and  $B_3$ . All three maximal bicliques include both  $nd(B_1, h_2).R_U$  and  $nd(B_1, h_2).R_V$ . Therefore, we can regard  $B_1$ ,  $B_2$ , and  $B_3$  as forming an incomplete BCPC, i.e., a partial-BCPC.

Next, we will formally define partial-BCPC, the maximal biclique enumeration tree and the special prefixes used to compute partial-BCPCs.

**DEFINITION 5 (PARTIAL-BCPC).** Given a bipartite graph  $G$  and two integer  $\alpha, \beta$ , a partial-BCPC  $\mathcal{P}$  is a set of maximal bicliques satisfying  $\mathcal{P} \subseteq \mathcal{B}$ , where  $\mathcal{B}$  is an  $(\alpha, \beta)$ -BCPC in  $G$ .

The distinction from Definition 4 lies in the fact that Definition 5 does not require the set of maximal bicliques in the partial-BCPC to be maximal. This reflects the characteristic of partial-BCPC as an

incomplete BCPC. In practice, partial-BCPCs are also maintained by union-find set.

**DEFINITION 6 (MAXIMAL BICLIQUE ENUMERATION TREE (MBE TREE)).** Given a bipartite graph  $G = (U, V, E)$ . The MBE tree  $T = (N, \vec{E})$  represents the search space of the maximal biclique enumeration algorithm on  $G$ . Here,  $N$  denotes the set of nodes in the tree, where each node  $nd \in N$  contains a six-tuple  $(R_U, R_V, C_U, C_V, X_U, X_V)$ , as defined in the basic framework of MBE in Section 2.1. The set  $\vec{E}$  consists of directed edges of the form  $(nd_i, nd_j, u)$ , where  $u \in U$  or  $u \in V$ . An edge  $(nd_i, nd_j, u) \in \vec{E}$  indicates that node  $nd_j$  is a child of node  $nd_i$ , generated by selecting vertex  $u$  from  $nd_i.C_U$  or  $nd_i.C_V$ . A node  $nd \in N$  is a leaf node if  $nd.C_U \cup nd.C_V = \emptyset$ . For a leaf node  $nd \in N$ , if  $nd.X_U \cup nd.X_V = \emptyset$ , then  $(nd.R_U, nd.R_V)$  is a maximal biclique.

For simplicity, we use  $nd.RC_X_U$  and  $nd.RC_X_V$  to denote  $nd.R_U \cup nd.C_U \cup nd.X_U$  and  $nd.R_V \cup nd.C_V \cup nd.X_V$ , respectively.

**DEFINITION 7 (( $\alpha, \beta$ )-PREFIX AND ( $\alpha, \beta$ )-NODE).** Given a bipartite graph  $G = (U, V, E)$  and its MBE tree  $T = (N, \vec{E})$ , let  $nd_1 \in N$  be the root node of  $T$ ,  $PF(nd_1) = (nd_1, nd_2, \dots, nd_l)$  is the path from  $nd_1$  to  $nd_l$ , i.e.,  $\forall i = 1, 2, \dots, l-1, (nd_i, nd_{i+1}, *) \in \vec{E}$ .  $PF(nd_l)$  is an  $(\alpha, \beta)$ -prefix if and only if  $|nd_l.R_U| = \alpha, |nd_{l-1}.R_U| < \alpha, |nd_l.R_V| \geq \beta$  or  $|nd_l.R_V| = \beta, |nd_{l-1}.R_V| < \beta, |nd_l.R_U| \geq \alpha$ . In this case,  $nd_l$  is an  $(\alpha, \beta)$ -node.

In short, an  $(\alpha, \beta)$ -node  $nd$  is the first node that satisfies  $|nd.R_U| \geq \alpha$  and  $|nd.R_V| \geq \beta$  in all branches where  $nd$  resides. For all its ancestor nodes  $nd_i$ , it is not possible to simultaneously satisfy  $|nd_i.R_U| \geq \alpha$  and  $|nd_i.R_V| \geq \beta$ .

**THEOREM 1.** For any two  $(\alpha, \beta)$ -nodes  $nd_1, nd_2$ ,  $nd_1$  can not be an ancestor or a descendant of  $nd_2$ .

**PROOF.** Based on Definition 7,  $nd_1$  is the first node that satisfies  $|nd_1.R_U| \geq \alpha$  and  $|nd_1.R_V| \geq \beta$  in all branches where  $nd_1$  resides. The same applies to  $nd_2$ . Thus,  $nd_1, nd_2$  are in two different branches, and  $nd_1$  can not be an ancestor or a descendant of  $nd_2$ .  $\square$

## 4.1 The Basic Framework

This section presents the basic framework for our partial-BCPC based solution. The framework consists of two fundamental steps: (1) Computing partial-BCPCs using the MBE tree; (2) Traversing the reduced MBAG based on the partial-BCPCs to derive the final BCPCs. When computing partial-BCPCs, the framework connects the maximal bicliques within the subtree rooted at specific nodes in the MBE tree using a union-find set. Specifically, we focus on  $(\alpha, \beta)$ -nodes in MBE tree. Ancestors of an  $(\alpha, \beta)$ -node are ignored as they fail to satisfy  $|R_U| \geq \alpha$  and  $|R_V| \geq \beta$ . Descendants of an  $(\alpha, \beta)$ -node, while satisfying  $|R_U| \geq \alpha$  and  $|R_V| \geq \beta$ , are redundant since any maximal biclique branches sharing the prefix corresponding to the descendants must also share the  $(\alpha, \beta)$ -prefix corresponding to the  $(\alpha, \beta)$ -node. Thus, focusing solely on  $(\alpha, \beta)$ -nodes enables efficient computation of partial-BCPCs.

**Implementation.** Algorithm 4 presents the basic framework. It begins by maximal biclique enumeration and extracting MBE tree  $T = (N, \vec{E})$  (Line 1). It then initializes the neighbor list of maximal bicliques and a union-find set  $UF$  to manage the partial-BCPCs and BCPCs (Lines 2-3). In Lines 5-8, the algorithm iterates over each  $(\alpha, \beta)$ -node  $nd$  in  $T$  (Line 5), invoking the ProcessNode function to gather all maximal bicliques under subtree rooted at  $nd$  into  $\mathcal{P}$  (Line 6). Subsequently, it merges all maximal bicliques in  $\mathcal{P}$  into the

### Algorithm 4: Partial-BCPC based solution

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**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All BCPC in  $G$

```

1 Let  $T = (N, \vec{E})$  be the MBE tree of  $G$ ,  $\mathcal{B}$  be the set of maximal bicliques in  $G$ ;
2  $Nei(\mathcal{B}) \leftarrow$  set of maximal bicliques that share vertices with  $\mathcal{B}$ ;
3 Let  $UF$  be an initial union-find set;
4 Let  $\mathcal{P} \leftarrow \emptyset$ ;
/* compute partial-BCPCs */
5 for  $(\alpha, \beta)$ -node  $nd$  in  $T$  do
6   ProcessNode( $nd$ );
7   for  $B \in \mathcal{P}$  do  $UF.union(B, \mathcal{P}[0])$ ;
8    $\mathcal{P} \leftarrow \emptyset$ ;
/* traverse MBAG with partial-BCPCs */
9 for  $B \in \mathcal{B}$  do
10  for  $B' \in Nei(B)$  and  $B, B'$  are in different partial-BCPCs do
11    if  $|B.X \cap B'.X| \geq \alpha, |B.Y \cap B'.Y| \geq \beta$  then  $UF.union(B, B')$ ;
12 return  $UF$ ;
/* recursively traverse the MBE tree to gather all maximal bicliques under node */
13 Procedure ProcessNode( $node$ )
14 for  $(node, node', u) \in \vec{E}$  do
15   if  $node'.C_U \cup node'.C_V = \emptyset$  then
16     if  $node'.X_U \cup node'.X_V = \emptyset$  then
17        $\mathcal{P} \leftarrow \mathcal{P} \cup \{(node'.R_U, node'.R_V)\}$ ; /*  $(node'.R_U, node'.R_V)$  is a maximal biclique */
18   else ProcessNode( $node'$ );

```

---

same partial-BCPC (Line 7). In Lines 9-11, the algorithm traverses a smaller MBAG based on the partial-BCPCs. Specifically, for each maximal biclique  $B$ , the algorithm only traverses maximal bicliques  $B'$  that reside in different partial-BCPCs from  $B$  (Line 10). The algorithm finally returns  $UF$ , which represents all detected BCPCs.

**The storage of the MBE tree.** In practice, we do not store all branches of the MBE tree. We first introduce the two types of nodes in the MBE tree.

**DEFINITION 8 (REAL AND VIRTUAL NODE).** Given a graph  $G$  and its MBE tree  $T = (N, \vec{E})$ , for any node  $nd \in N$ , if  $nd$  is a leaf node and satisfies  $nd.X_U = nd.X_V = \emptyset$ , then  $nd$  is a real node—in this case,  $(nd.R_U, nd.R_V)$  is a maximal biclique (see Definition 6). Otherwise, if  $nd$  has at least one descendant that is a real node, it is also a real node. If neither of these conditions is satisfied, then  $nd$  is a virtual node.

The ProcessNode procedure in Algorithm 4 only needs to identify the branches that contain maximal bicliques (Lines 15-17), i.e., branches composed of real nodes. Thus, in practice, we only store real nodes in the MBE tree.

**Analysis.** We first analyze the correctness of Algorithm 4, then examine the relationship between the number of partial-BCPCs obtained by Algorithm 4 and the numbers of maximal bicliques as well as BCPCs, and finally analyze the complexity of Algorithm 4.

**THEOREM 2.** Algorithm 4 correctly computes all BCPCs.

**PROOF.** We first prove that the partial-BCPCs are all correct. The procedure ProcessNode collects all maximal bicliques within the subtree rooted at the given  $node$  into  $\mathcal{P}$ . Since the parameter is always an  $(\alpha, \beta)$ -node, the maximal bicliques collected in each execution of the procedure must share an  $(\alpha, \beta)$ -biclique. Therefore, these bicliques correctly form a partial-BCPC. Then we prove that the final BCPCs are correct. In Lines 9-11, any two maximal bicliques that belong to different partial-BCPCs but satisfy the adjacency condition are connected via the union-find set  $UF$ . As a

result, each disjoint set maintained by  $UF$  corresponds to a valid BCPC.  $\square$

**THEOREM 3.** *Given a bipartite graph  $G$ , two integer  $\alpha, \beta$  and MBE tree  $T = (N, \vec{E})$ , let  $n_{biclique}$  be the number of maximal bicliques,  $ND \leftarrow \{nd | nd \in N, nd \text{ is an } (\alpha, \beta)\text{-node}, nd \text{ is a real node}\}$ ,  $pbcp$  be the number of partial-BCPCs computed by Algorithm 4,  $bcpc$  be the number of BCPCs, the following inequality holds:  $n_{biclique} \geq |ND| = pbcp \geq bcpc$ .*

**PROOF.** Based on Theorem 1, any two  $(\alpha, \beta)$ -nodes in  $ND$  can not be in the same branch, and nodes in  $ND$  are all real nodes, thus, each node in  $ND$  has at least one maximal biclique in its subtree. As a result,  $n_{biclique} \geq |ND|$ . Since Algorithm 4 connects all maximal bicliques in the subtree of each  $(\alpha, \beta)$ -node in  $ND$  into a partial-BCPC, and the maximal bicliques from different subtrees of  $(\alpha, \beta)$ -nodes are distinct, thus, each  $(\alpha, \beta)$ -node in  $ND$  corresponds to one partial-BCPC ( $|ND| = pbcp$ ). Partial-BCPCs are incomplete BCPCs according to Definition 5, thus,  $pbcp \geq bcpc$ .  $\square$

**THEOREM 4.** *The time and space complexity of Algorithm 4 are  $O((1 - \frac{1}{pbcp})n_{biclique}^2 n)$  and  $O(n_{biclique}n)$  respectively, where  $n = |U| + |V|$ ,  $pbcp$  is the number of partial-BCPCs,  $n_{biclique}$  is the number of maximal bicliques.*

**PROOF.** Let  $m = |E|$ , the time complexity of maximal biclique enumeration is  $O(2^{n/2}m)$  [10]. Compared to maximal biclique enumeration, Algorithm 4 introduces additional operations: traversing  $(\alpha, \beta)$ -nodes and collecting maximal biclique branches from the subtrees rooted at these nodes. Since  $(\alpha, \beta)$ -nodes do not reside in the same branch (Theorem 1), the time complexity of these additional operations does not exceed  $O(2^{n/2}m)$ . In Lines 9–11, in the worst case, Algorithm 4 only skips the inner mutual visits of maximal bicliques in each initial partial-BCPC. Let  $x = n_{biclique}$ ,  $y = pbcp$  and  $p_1, p_2, \dots, p_y$  be the number of maximal bicliques in each partial-BCPC, the time complexity of Lines 9–11 is  $O(x^2n - (p_1^2 + p_2^2 + \dots + p_y^2)n) = O(x^2n - (\frac{x}{y})^2yn)$  (based on Cauchy-Schwarz inequality), which can be written as  $O((1 - \frac{1}{pbcp})n_{biclique}^2 n)$ . As a result, the time complexity of Algorithm 4 is  $O(2^{n/2}m + (1 - \frac{1}{pbcp})n_{biclique}^2 n)$ . Since  $O(n_{biclique}) = O(2^{n/2})$  [10], the time complexity can be rewritten as  $O((1 - \frac{1}{pbcp})n_{biclique}^2 n)$ .

As for the space complexity, the main space consumption in Algorithm 4 comes from the MBE tree and maximal bicliques. Since only the real nodes in the MBE tree are stored, the number of branches in the MBE tree is equal to the number of maximal bicliques, thus, the space complexity of Algorithm 4 is  $O(n_{biclique}n)$ .  $\square$

**Discussion.** The inequality  $n_{biclique} \geq pbcp$  in Theorem 3 forms the basis for reducing the MBAG using partial-BCPC. This is because maximal bicliques within the same partial-BCPC are treated as a single vertex in the MBAG, which leads to an MBAG with formally fewer vertices. This can also be seen from the time complexity in Theorem 4: if  $pbcp$  is replaced with  $n_{biclique}$ , the result is exactly the complexity of the existing method in Section 3 for traversing the complete MBAG ( $O(n_{biclique}^2 n)$ ). Experiments (Figure 5 (c)) show that  $pbcp$  can be an order of magnitude smaller than  $n_{biclique}$ .

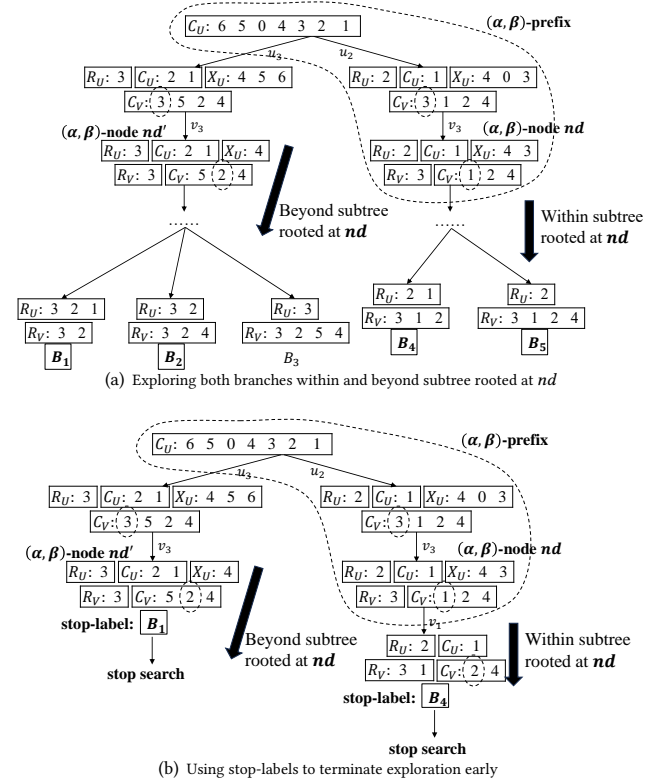


Figure 3: Examples of optimization

## 4.2 Optimizations

**Beyond subtree rooted at  $(\alpha, \beta)$ -node.** The basic framework in the previous section directly merges the maximal bicliques in the subtree rooted at an  $(\alpha, \beta)$ -node into a partial-BCPC. However, this approach is insufficient because, for an  $(\alpha, \beta)$ -node  $nd$ , not all maximal bicliques sharing  $nd.R_U$  and  $nd.R_V$  are located within the subtree rooted at  $nd$ . For example, in Figure 2, if  $\alpha = \beta = 1$ ,  $nd(B_4, h_2)$  is an  $(\alpha, \beta)$ -node, but  $B_1$  and  $B_2$  are not in the subtree rooted at  $nd(B_4, h_2)$ , even though both  $B_1$  and  $B_2$  share  $nd(B_4, h_2).R_U$  and  $nd(B_4, h_2).R_V$ . This phenomenon occurs because some vertices in  $B_1$  and  $B_2$  (e.g.,  $u_3$ ) appear early in the candidate set  $C_U$  of the root node, causing  $B_1$  and  $B_2$  to be enumerated much earlier.

To merge all relevant maximal bicliques within and beyond the subtree of an  $(\alpha, \beta)$ -node, it is essential to fully utilize all the information maintained in the  $(\alpha, \beta)$ -node, i.e.,  $R_U, R_V, C_U, C_V, X_U$ , and  $X_V$ , to systematically search on the MBE tree. Specifically, for each  $(\alpha, \beta)$ -node  $nd$ , we use  $nd.RC_X_U$  and  $nd.RC_X_V$  to search, within the MBE tree, all branches of maximal bicliques that contain  $nd.R_U$  and  $nd.R_V$ , and connect them into the same partial-BCPC.

**EXAMPLE 4.** Let  $\alpha = 1, \beta = 1$ , Figure 3 (a) illustrates that when processing  $(\alpha, \beta)$ -node  $nd$ , both the branches within and beyond subtree rooted at  $nd$  are taken into account. It can be observed that the maximal biclique branches sharing  $(nd.R_U, nd.R_V)$  include not only  $B_4$  and  $B_5$  under  $nd$  but also the maximal biclique branches  $B_1$  and  $B_2$  beyond the subtree rooted at  $nd$ .

**Stop-Label.** The above method may result in fully traversing all the relevant maximal biclique branches when processing each  $(\alpha, \beta)$ -node. To reduce the overhead of traversing all maximal biclique branches that share  $nd.R_U$  and  $nd.R_V$  ( $nd$  is an  $(\alpha, \beta)$ -node) without affecting correctness, we assign a stop-label to certain real nodes in the MBE tree. When such a real node is visited, the stop-label allows immediate retrieval of all maximal biclique information within its subtree, thus eliminating the need to further traverse its descendants. Given  $\alpha, \beta$ , the following presents the details of the stop-label:

- (1) Stop-labels are assigned only to real nodes in the MBE tree.
- (2) For a descendant node  $nd$  of an  $(\alpha, \beta)$ -node, the prerequisite for setting stop-label to  $nd$  is to connect all maximal bicliques within the subtree rooted at  $nd$  into the same partial-BCPC. The stop-label of  $nd$  is set to one of the maximal bicliques in that partial-BCPC.
- (3) For an  $(\alpha, \beta)$ -node  $nd$ , the prerequisite for setting stop-label to  $nd$  is to connect all maximal bicliques sharing  $nd.R_U$  and  $nd.R_V$  into the same partial-BCPC. The stop-label of  $nd$  is set to one of the maximal bicliques in that partial-BCPC.

We now describe how stop-labels contribute to pruning in identifying branches.

- (1) For a descendant node  $nd$  of an  $(\alpha, \beta)$ -node, before setting a stop-label to  $nd$ , it is sufficient to merge the stop-labels of all its child nodes (also real nodes) into the same partial-BCPC using union-find set. This eliminates the need to traverse all maximal biclique branches within the subtree rooted at  $nd$ .
- (2) For  $(\alpha, \beta)$ -node  $nd$ , in identifying all maximal biclique branches that share  $nd.R_U$  and  $nd.R_V$ , the search terminates as soon as it encounters a node with a stop-label and directly adds the stop-label to the set  $\mathcal{P}$ .

**EXAMPLE 5.** Let  $\alpha = 1, \beta = 1$ , when processing  $nd$  in Figure 3 (b): During the search of subtree rooted at  $nd$ , the process terminates upon encountering the stop-label of child node of  $nd$ . This stop-label is  $B_4$ , with  $B_5$  belonging to the same partial-BCPC as  $B_4$ .

During the search beyond the subtree rooted at  $nd$ , the search process terminates upon detecting the stop-label of  $nd$ .

Finally, the two stop-labels ( $B_1$  and  $B_4$ ) are connected; concurrently, the partial-BCPCs they belong to are merged into a larger partial-BCPC.

**Implementation.** Based on the above strategies, we revise the original framework and present Algorithm 5. Algorithm 5 performs an initialization step in Line 1 that is similar to that of Algorithm 4. In Line 3,  $ST(nd)$  denotes the stop-label of  $nd$ . The core component of Algorithm 5 is the Postorder function (Lines 4 and 7). This function performs a post-order traversal of the MBE tree, a process that can be executed during the MBE progress. The Postorder function is responsible for handling both the  $(\alpha, \beta)$ -node itself—by invoking  $\text{ProcessNode+}(node, UV, RUV)$  in Line 10—and the real nodes among its descendant nodes—by invoking  $\text{ProcessNode+}(node)$  in Line 12. Both processing routines may add certain maximal bicliques to  $\mathcal{P}$ . Finally, in Lines 13–15, Algorithm 5 connects all the maximal bicliques in  $\mathcal{P}$  into the same partial-BCPC, and it set the first biclique in  $\mathcal{P}$  as the stop-label of  $node$  (Line 15). Similar to Algorithm 4, Algorithm 5 traverses the MBAG based on partial-BCPCs to obtain the final BCPC (Line 5).

When processing an  $(\alpha, \beta)$ -node (Line 10), the  $\text{ProcessNode+}$  function starts traversing the MBE tree from its root node. It

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#### Algorithm 5: Partial-BCPC based solution with optimizations

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Input: Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$ 
Output: All BCPC in  $G$ 
1 Lines 1-3 of Algorithm 4;
2 Let  $\mathcal{P} \leftarrow \emptyset$ ,  $root \leftarrow$  root node of  $T$ ;
3 Let  $ST(nd)$ ,  $nd \in N$  be the stop-label of  $nd$ ;
  /* compute partial-BCPCs */
4 Postorder( $root$ );
  /* traverse MBAG with partial-BCPCs */
5 Lines 9-11 of Algorithm 4;
6 return  $UF$ ;

7 Procedure Postorder( $node$ )
8 foreach child of  $node$ ,  $node'$ , in creation order of  $node'$  do Postorder( $node'$ );
9 if  $node$  is an  $(\alpha, \beta)$ -node then
10   ProcessNode+( $root$ ,  $node.RC_{X_U} \cup node.RC_{X_V}$ ,  $node.R_U \cup node.R_V$ );
11 else if  $node$  is a real node and a descendant of an  $(\alpha, \beta)$ -node then
12   ProcessNode+( $node$ );
13 if  $\mathcal{P}$  is not empty then
14   for  $B \in \mathcal{P}$  do  $UF.union(B, \mathcal{P}[0])$ ;
15    $ST(node) \leftarrow \mathcal{P}[0]$ ,  $\mathcal{P} \leftarrow \emptyset$ ;

16 Procedure ProcessNode+( $node$ ,  $UV$ ,  $RUV$ )
17 foreach child of  $node$ ,  $node'$ , in creation order of  $node'$  do
18   Let  $(node', u) \in E$ ;
19   if  $node'$  is a real node then
20     if  $u \in UV$  then
21       if  $ST(node') = \text{None}$  then ProcessNode+( $node'$ ,  $UV$ ,  $RUV$ );
22       else  $\mathcal{P} \leftarrow \mathcal{P} \cup \{ST(node')\}$ ;
23   if  $u \in RUV$  then break;

24 Procedure ProcessNode+( $node$ )
25 if  $node$  is a leaf node then  $\mathcal{P} \leftarrow \mathcal{P} \cup \{(node.R_U, node.R_V)\}$ ;
26 else
27   foreach  $node'$ ,  $node'$  is a child of  $node$  and a real node do
28      $\mathcal{P} \leftarrow \mathcal{P} \cup \{ST(node')\}$ 

```

---

mainly explores real nodes (Line 19), since any branch containing a maximal biclique must be composed of real nodes. Moreover, any vertex  $u$  along the traversal path must belong to the set  $UV$  (Line 20). The recursive calls end immediately upon encountering a node that has a stop-label assigned (Lines 22). After processing a child node  $node'$ , if the vertex  $u$  along the traversal path belongs to the set  $RUV$ , there is no need to explore the remaining child nodes and their corresponding branches (Line 23). This is because, according to the enumeration principle of the MBE algorithm, any maximal biclique that includes  $RUV$  can not be found in those subsequent branches. Since the goal of this function is to identify maximal bicliques that contain  $RUV$ , further traversal becomes unnecessary in this case.

When processing the descendant nodes of an  $(\alpha, \beta)$ -node (Line 12), if the parameter  $node$  is already a leaf node, the maximal biclique it represents can be directly added to  $\mathcal{P}$ . Otherwise, the stop-labels of all its child nodes are added to  $\mathcal{P}$ .

**Analysis.** We now proceed to analyze the correctness of Algorithm 5. The key lies in demonstrating that the partial-BCPCs computed in Postorder (Line 3) are correct.

**THEOREM 5.** For each  $(\alpha, \beta)$ -node processed in Algorithm 5 (Line 9),  $\text{ProcessNode+}$  connects all maximal bicliques in the bipartite graph that share  $(node.R_U, node.R_V)$  into the same partial-BCPC.

**PROOF.** Consider a sequence of  $(\alpha, \beta)$ -nodes  $(nd_1, nd_2, \dots, nd_n)$ , which follow the processing order in Postorder (i.e., post-order traversal). We first prove that during the processing of  $nd_1$ ,  $\text{ProcessNode+}$  connects all maximal bicliques that share  $(nd_1.R_U, nd_1.R_V)$  into the same partial-BCPC. Since  $nd_1$  is the first  $(\alpha, \beta)$ -node, it must also be in the first enumerated maximal



biclique branch (if  $nd_1$  appears in an earlier branch, then that earlier or an even earlier branch is likely the first one containing a maximal biclique; if  $nd_1$  appears in a later branch, then it can not be the first  $(\alpha, \beta)$ -node). Therefore, all maximal biclique branches that contain  $(nd_1.R_U, nd_1.R_V)$  must lie in the subtree rooted at  $nd_1$ , and can not appear in any subtree processed before  $nd_1$ .

Using mathematical induction, assume that ProcessNode+ connects all maximal bicliques that share  $(nd_i.R_U, nd_i.R_V)$  when processing  $nd_1, nd_2, \dots, nd_i$ . Next, we prove the case for  $i + 1$ . For  $nd_{i+1}$ , let  $R_U = nd_{i+1}.R_U$  and  $R_V = nd_{i+1}.R_V$ . According to the MBE process [10], the branches containing maximal bicliques with  $R_U$  and  $R_V$  are either generated before  $nd_{i+1}$ , or located in the subtree rooted at  $nd_{i+1}$ . For the maximal biclique branches generated before  $nd_{i+1}$ , their corresponding  $(\alpha, \beta)$ -node  $nd_j$  must satisfy  $j < i + 1$ , so we only need to collect  $ST(nd_j)$ , as it represents the partial-BCPCs of those maximal bicliques. For the maximal bicliques in the subtree of  $nd_{i+1}$ , we only need to collect the stop-labels of the child nodes of  $nd_{i+1}$ , because the maximal bicliques under each child node have already been connected into partial-BCPCs (since the traversal is post-order). Therefore, the case for  $nd_{i+1}$  is proved.  $\square$

**THEOREM 6.** *Algorithm 5 correctly computes all BCPCs.*

**PROOF.** The proof of this theorem is similar to that of Theorem 2. Since Theorem 5 proves that the partial-BCPCs in Algorithm 5 are obtained by connecting maximal bicliques that share the  $R_U$  and  $R_V$  of  $(\alpha, \beta)$ -nodes, these partial-BCPCs are correct. Moreover, the process of computing the final BCPCs in Algorithm 5 (Line 5) is the same as in Algorithm 4, thus, Algorithm 5 can correctly compute the BCPCs.  $\square$

It should be noted that the partial-BCPCs computed by Algorithm 5 are not the same as those computed by Algorithm 4. We illustrate this with the following theorem.

**THEOREM 7.** *Given a bipartite graph  $G$ , two integer  $\alpha, \beta$ , let  $pbcp$  be the number of partial-BCPCs computed by Algorithm 4,  $pbcp+$  be the number of partial-BCPCs computed by Algorithm 5,  $bcpc$  be the number of BCPCs, the following inequality holds:  $pbcp \geq pbcp+ \geq bcpc$ .*

**PROOF.** We only need to prove  $pbcp \geq pbcp+$ . When processing an  $(\alpha, \beta)$ -node, Algorithm 5 not only connects maximal bicliques in the subtree rooted at that node, but also further connects maximal bicliques that were enumerated in earlier branches and share  $R_U, R_V$  of the  $(\alpha, \beta)$ -node. Moreover, the ProcessNode+ procedure in Algorithm 5 also processes  $(\alpha, \beta)$ -nodes that belong to virtual nodes. Therefore,  $pbcp \geq pbcp+$ .  $\square$

**THEOREM 8.** *The time and space complexity of Algorithm 5 are  $O((1 - \frac{1}{pbcp+})n_{biclique}^2n)$  and  $O(n_{biclique}n)$  respectively, where  $n = |U| + |V|$ ,  $pbcp+$  is the number of partial-BCPCs,  $n_{biclique}$  is the number of maximal bicliques.*

**PROOF.** Let  $m = |E|$ , the time complexity of maximal biclique enumeration is  $O(2^{n/2}m)$  [10]. Unlike Algorithm 4, Algorithm 5 handles not only  $(\alpha, \beta)$ -nodes (Lines 9-10) but also their child nodes (Lines 11-12). For processing the child nodes of  $(\alpha, \beta)$ -nodes, Algorithm 5 only needs to traverse one layer of their child nodes, resulting in an overall time complexity not exceeding  $O(2^{n/2}m)$ . Regarding the processing of  $(\alpha, \beta)$ -nodes, let  $x$  be the total number of such nodes (including both real nodes and virtual nodes),  $d$  be

#### Algorithm 6: $(\alpha, \beta)$ -Biclique based solution

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**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All BCPC in  $G$

- 1 Let  $UF$  be an initial union-find set;
- 2  $\mathcal{B} \leftarrow$  set of all maximal bicliques in  $G$ ;
- 3  $\mathcal{B} \leftarrow \{B \in \mathcal{B}, |B.X| \geq \alpha, |B.Y| \geq \beta\}$ ;
- 4 Construct 2-hop graph  $\tilde{H} = (U, \tilde{E}_H)$  on vertex set  $U$  [30];  $\tilde{H} = (U, \tilde{E}_H)$ : a directed graph,  $(u, v) \in \tilde{E}_H$  indicates that  $u < v$ ,  $u, v$  have common neighbors in  $V$  \*/
- 5 BicliqueListing( $\tilde{H}, \emptyset, V$ );
- 6 return  $UF$ ;

---

7 **Procedure** BicliqueListing( $\tilde{H}, R_U, C_V$ )

- 8 if  $|R_U| = \alpha$  then
- 9   foreach  $R_V \subseteq C_V, |R_V| = \beta$  do Connect maximal bicliques sharing  $(R_U, R_V)$  in  $\mathcal{B}$  with  $UF.union(*)$ ;
- 10   return;
- 11 foreach  $u \in \tilde{H}$  do
- 12    $C'_V \leftarrow C_V \cap Nei_G(u)$ ;
- 13   if  $|C'_V| < \beta$  or  $Nei_{\tilde{H}}(u) < \alpha - |R_U| - 1$  then continue;
- 14   Let  $\tilde{H}'$  be induced subgraph of  $\tilde{H}$  on  $Nei_{\tilde{H}}(u)$ ;
- 15   BicliqueListing( $\tilde{H}', R_U \cup \{u\}, C'_V$ );

---

the maximum depth of real  $(\alpha, \beta)$ -nodes. When processing each of these nodes (Line 10), the procedure ProcessNode+ will access at most all real  $(\alpha, \beta)$ -nodes. Since the number of child nodes of each node in the MBE tree does not exceed  $n$ , the total time complexity of the above process is  $O(xn^d) = O(2^{n/2}mn^d)$ . The time complexity analysis for the traversal of MBAG in Algorithm 5 is similar to that of Algorithm 4. As a result, the time complexity of Algorithm 5 is  $O(2^{n/2}mn^d + (1 - \frac{1}{pbcp+})n_{biclique}^2n)$ . Since  $O(n_{biclique}) = O(2^{n/2})$  [10?], the time complexity can be rewritten as  $O((1 - \frac{1}{pbcp+})n_{biclique}^2n)$ .

Similar to Algorithm 4, the space complexity of Algorithm 5 is also  $n_{biclique}n$ .  $\square$

**Discussion.** The inequality  $pbcp \geq pbcp+$  in Theorem 7 forms the basis for Algorithm 5 outperforming Algorithm 4, since it implies that Algorithm 5 can traverse a smaller MBAG. Experiments (Figure 5) show that  $pbcp+$  is significantly smaller than  $pbcp$ —for example, smaller by an order of magnitude, and Algorithm 5 can be an order of magnitude faster than Algorithm 4 (Table 2).

## 5 Novel $(\alpha, \beta)$ -Biclique based Solution

Existing BCPC detection methods are all based on the MBAG approach. In the previous section, we reduce the MBAG using partial-BCPCs, but it is still essentially based on MBAG and remains inefficient for certain datasets where MBAG is extremely large. This section introduces a novel BCPC detection framework, which can also leverage the partial-BCPCs proposed in Section 4 to accelerate the detection process. We first introduce this framework.

### 5.1 The Basic Framework

**Key idea.** Note that in the previous section, each edge in the MBAG essentially represents an  $(a, b)$ -biclique, where  $a \geq \alpha$  and  $b \geq \beta$ . A straightforward idea is to enumerate all  $(\alpha, \beta)$ -bicliques in the bipartite graph, and for each enumerated  $(\alpha, \beta)$ -biclique  $B$ , connect all maximal bicliques containing  $B$  using union-find set.

**Implementation.** Algorithm 6 presents the basic framework of the  $(\alpha, \beta)$ -biclique based solution. It is built upon the  $(\alpha, \beta)$ -biclique enumeration framework proposed in [30], where the core idea is to



enumerate combinations of vertices in the  $U$  set while maintaining their common neighbors in the  $V$  set. This framework constructs an ordinary directed graph, called the *2-hop graph*, on the vertices of the  $U$  set to accelerate the enumeration process.

In Lines 2–3, Algorithm 6 prepares the maximal bicliques that need to be connected. In Line 4, it constructs the 2-hop graph according to [30]. Then, in Line 5, it executes the  $(\alpha, \beta)$ -biclique enumeration framework. Within BicliqueListing, for each enumerated  $(\alpha, \beta)$ -biclique (Line 9), all maximal bicliques that share this biclique are connected together.

**Analysis.** Now we proceed to analyze the correctness and complexity of Algorithm 6.

**THEOREM 9.** *Algorithm 6 correctly computes all BCPCs.*

**PROOF.** This theorem can be easily proven as follows: Algorithm 6 is capable of enumerating all  $(\alpha, \beta)$ -bicliques and connecting all maximal bicliques that share these bicliques. Consequently, the final BCPCs are correct.  $\square$

**THEOREM 10.** *The time and space complexity of Algorithm 6 are  $O(a(\vec{H})^{\alpha-2}|\vec{E}_h||V| + \Delta n_{\text{biclique}})$  and  $O(n_{\text{biclique}}n)$  respectively, where  $n = |U| + |V|$ ,  $a(\vec{H})$  is the arboricity of  $\vec{H}$  [30],  $\Delta$  is the number of  $(\alpha, \beta)$ -bicliques,  $n_{\text{biclique}}$  is the number of maximal bicliques.*

**PROOF.** The time complexity of maximal biclique enumeration is  $T_{\text{biclique}} = 2^{n/2}|E|$  [10]. The time complexity of enumerating  $(\alpha, \beta)$ -biclique is  $O(a(\vec{H})^{\alpha-2}|\vec{E}_h||V| + \Delta)$  [30]. Algorithm 6 connects all maximal bicliques sharing an  $(\alpha, \beta)$ -biclique (Line 9), thus, the time complexity of Algorithm 6 is  $O(T_{\text{biclique}} + a(\vec{H})^{\alpha-2}|\vec{E}_h||V| + \Delta n_{\text{biclique}})$ . Since  $O(n_{\text{biclique}}) = O(2^{n/2})$  [10?],  $O(\Delta) = O(n^{\alpha+\beta})$  [30], the time complexity can be rewritten as  $O(a(\vec{H})^{\alpha-2}|\vec{E}_h||V| + \Delta n_{\text{biclique}})$ .

Algorithm 6 does not store all  $(\alpha, \beta)$ -bicliques, thus, the space complexity of Algorithm 6 is  $O(n_{\text{biclique}}n)$ .  $\square$

## 5.2 Pruning with Maximal Bicliques and Partial-BCPCs

**Challenges in the basic framework.** It is evident that the basic framework in Algorithm 6 is highly inefficient, as the number of  $(\alpha, \beta)$ -bicliques grows exponentially with respect to  $\alpha$  and  $\beta$  [30]. Therefore, enumerating all  $(\alpha, \beta)$ -bicliques and connecting the corresponding maximal bicliques is impractical.

**Pruning with Maximal Bicliques.** The key to addressing the above challenge is to reduce the search space in the enumeration process of  $(\alpha, \beta)$ -bicliques — that is, to prune in advance those branches that do not contribute to connecting multiple maximal bicliques.

**THEOREM 11.** *Given a bipartite graph  $G$ , for a biclique  $B$ , if the number of maximal bicliques sharing it is 0, or all those maximal bicliques are in the same BCPC, then for any other biclique  $B'$  such that  $B \subseteq B'$ , the number of maximal bicliques sharing  $B'$  is also 0, or they are in the same BCPC.*

**PROOF.** It is easy to see that any maximal biclique containing  $B'$  must also contain  $B$ , so the set of maximal bicliques containing  $B'$  is a subset of those containing  $B$ . Therefore, Theorem 11 can be easily proved.  $\square$

### Algorithm 7: $(\alpha, \beta)$ -Biclique based solution with maximal biclique

---

**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All BCPC in  $G$

---

```

1 Let  $UF$  be an initial union-find set;
2  $\mathcal{B} \leftarrow$  set of all maximal bicliques in  $G$ ;
3  $\mathcal{B} \leftarrow \{B | B \in \mathcal{B}, |B.X| \geq \alpha, |B.Y| \geq \beta\}$ ;
4 Construct 2-hop graph  $\vec{H} = (U, \vec{E}_h)$  on vertex set  $U$  [30]; /*  $\vec{H} = (U, \vec{E}_h)$ : a directed graph,  $(u, v) \in \vec{E}_h$  indicates that  $u < v$ ,  $u, v$  have common neighbors in  $V$  */
5 BicliqueListing+ ( $\vec{H}, \emptyset, V$ );
6 return  $UF$ ;

7 Procedure BicliqueListing+ ( $\vec{H}, R_U, C_V$ )
8  $RB \leftarrow \{B | R_U \subseteq B.X, B \in \mathcal{B}\}$ ; /* maximal bicliques sharing  $R_U$  */
9 if  $|\{UF.find(B) | B \in RB\}| \leq 1$  then return; /* there is only 1 or 0  $UF$  disjoint set in  $RB$  */
10 if  $|R_U| = \alpha$  then
11   foreach  $R_V \subseteq C_V, |R_V| = \beta$  do
12      $RB \leftarrow \{B | R_V \subseteq B.Y, B \in \mathcal{B}\}$ ;
13     if  $|\{UF.find(B) | B \in RB\}| \leq 1$  then continue; /* there is only 1 or 0  $UF$  disjoint set in  $RB$  */
14     Connect maximal bicliques in  $RB$  with  $UF.union(*)$ ;
15   return;
16 foreach  $u \in \vec{H}$  do
17    $C_V \leftarrow C_V \cap Nei_G(u)$ ;
18   if  $|C_V| < \beta$  or  $Nei_{\vec{H}}(u) < \alpha - |R_U| - 1$  then continue;
19   Let  $\vec{H}'$  be induced subgraph of  $\vec{H}$  on  $Nei_{\vec{H}}(u)$ ;
20   BicliqueListing+ ( $\vec{H}', R_U \cup \{u\}, C_V$ );

```

---

Theorem 11 reveals the core idea of pruning the  $(\alpha, \beta)$ -biclique enumeration space: during the enumeration process, for each intermediate small biclique generated, we can maintain the set of maximal bicliques containing it. If this set is empty or all the maximal bicliques in the set already belong to the same BCPC, then the current enumeration branch can be safely pruned.

**Implementation.** Algorithm 7 presents the implementation of pruning the  $(\alpha, \beta)$ -biclique enumeration process using maximal bicliques. The difference between Algorithm 7 and Algorithm 6 lies in the BicliqueListing+ function. Specifically, in Line 8, BicliqueListing+ collects maximal bicliques that share  $(R_U, \emptyset)$ . In Line 9, if the maximal bicliques in  $RB$  belong to only one or zero union-find sets, then the entire enumeration branch can be pruned. In Lines 12–13, the algorithm further checks whether the maximal bicliques in  $RB$  that share  $R_V$  belong to more than one union-find set. If so, it connects them using  $UF.union(*)$ .

**Analysis.** Now we proceed to analyze the correctness and complexity of Algorithm 7.

**THEOREM 12.** *Algorithm 7 correctly computes all BCPCs.*

**PROOF.** Compared with Algorithm 6, Algorithm 7 prunes the branches that can not connect multiple  $UF$  sets based on Theorem 11. Therefore, Algorithm 7 maintains the same correctness as Algorithm 6.  $\square$

**THEOREM 13.** *The time and space complexity of Algorithm 7 are  $O((a(\vec{H}_m)^{\alpha-2}|\vec{E}_{hm}||V_m| + \Delta_m)n_{\text{biclique}})$  and  $O(n_{\text{biclique}}n)$  respectively.  $n = |U| + |V|$ . Let  $U_m$  and  $V_m$  be the vertex sets shared by any two maximal bicliques,  $G_m$  be the induced subgraph on  $U_m$  and  $V_m$ , and  $\vec{H}_m = (U_m, \vec{E}_{hm})$  be the 2-hop graph based on  $G_m$ .  $a(\vec{H}_m)$  is the arboricity of  $\vec{H}_m$  [30],  $\Delta_m$  is the number of  $(\alpha, \beta)$ -bicliques in  $G_m$ ,  $n_{\text{biclique}}$  is the number of maximal bicliques.*

**Algorithm 8:**  $(\alpha, \beta)$ -Biclique based solution with partial-BCPC

---

**Input:** Bipartite graph  $G = (U, V, E)$ ; Two integer  $\alpha, \beta$   
**Output:** All BCPC in  $G$

---

```

1 Lines 1-3 of Algorithm 5; /* partial-BCPC computation */
2 Lines 2-5 of Algorithm 7; /* connect maximal bicliques by  $(\alpha, \beta)$ -biclique listing */
3 return  $UF$ ;
```

---

**PROOF.** The time complexity of maximal biclique enumeration is  $T_{biclique} = 2^{n/2}|E|$  [10]. Compared with Algorithm 6, Algorithm 7 prunes the branches that can not connect multiple  $UF$  sets based on Theorem 11. This implies that vertices initially contained in only one maximal biclique need not be considered during the  $(\alpha, \beta)$ -biclique enumeration process. Therefore, Algorithm 7 can be regarded as performing  $(\alpha, \beta)$ -biclique enumeration on  $G_m$ . Considering Lines 8-9 and Lines 12-13 of Algorithm 7, its time complexity is  $O(T_{biclique} + (a(\vec{H}_m)^{\alpha-2}|\vec{E}_{hm}||V_m| + \Delta_m)n_{biclique})$ . Since  $O(n_{biclique}) = O(2^{n/2})$  [10?],  $O(\Delta_m) = O((U_m + V_m)^{\alpha+\beta})$  [30], the time complexity can be rewritten as  $O((a(\vec{H}_m)^{\alpha-2}|\vec{E}_{hm}||V_m| + \Delta_m)n_{biclique})$ .

Similarly, the space complexity of Algorithm 7 is still  $O(n_{biclique}n)$ .  $\square$

**Pruning with Partial-BCPC.** As shown in Section 4, a partial-BCPC is an incomplete BCPC. Based on the partial-BCPCs, we can determine in advance that certain maximal bicliques belong to the same BCPC. Therefore, they enable the  $(\alpha, \beta)$ -biclique enumeration process in Algorithm 7 to reach pruning conditions earlier.

**Implementation.** Algorithm 8 is the algorithm that leverages partial-BCPCs, and it consists of two main parts. First, it computes the partial-BCPC (Line 1), which are recorded in the union-find structure ( $UF$ ). Then, in Line 2, these partial-BCPCs are used to further prune the  $(\alpha, \beta)$ -biclique listing process.

**Analysis.** Now we proceed to analyze the correctness and complexity of Algorithm 8.

**THEOREM 14.** *Algorithm 8 correctly computes all BCPCs.*

**THEOREM 15.** *The time and space complexity of Algorithm 8 are  $O((a(\vec{H}_p)^{\alpha-2}|\vec{E}_{hp}||V_p| + \Delta_p)n_{biclique})$  and  $O(n_{biclique}n)$  respectively.  $n = |U| + |V|$ . Let  $U_p$  and  $V_p$  be the vertex sets shared by any two partial-BCPCs,  $G_p$  be the induced subgraph on  $U_p$  and  $V_p$ , and  $\vec{H}_p = (U_p, \vec{E}_{hp})$  be the 2-hop graph based on  $G_p$ .  $a(\vec{H}_p)$  is the arboricity of  $\vec{H}_p$  [30],  $\Delta_p$  is the number of  $(\alpha, \beta)$ -bicliques in  $G_p$ ,  $n_{biclique}$  is the number of maximal bicliques.*

The proves of Theorem 14 and Theorem 15 are similar to those of Theorem 12 and Theorem 13.

## 6 Experiments

### 6.1 Experimental Setup

**Datasets.** We use 10 real-world bipartite graph datasets, whose detailed information is presented in Table 1. All the datasets are downloaded from <http://konect.cc/>.

**Algorithms.** We first implement the state-of-the-art algorithm in [7, 15] as MBAG.

**Table 1: Datasets,  $\bar{d}_U$  is the average degree of vertices in  $U$ ,  $\bar{d}_V$  is the average degree of vertices in  $V$ ,  $N_m$  is the number of maximal bicliques,  $N_o$  is the number of edges in maximal biclique adjacent graph,  $K=10^3$ ,  $M=10^6$ ,  $B=10^9$ , ‘-’ means timeout**

Datasets	$ U $	$ V $	$ E $	$\bar{d}_U$	$\bar{d}_V$	$N_m$	$N_o$
Youtube	94,238	30,087	293,360	3.11	9.75	1,769,331	2B
Bookcrossing	77,802	185,955	433,652	5.57	2.33	155,391	1B
Github	56,519	120,867	440,237	7.79	3.64	55,260,550	-
Citeseer	105,353	181,395	512,267	4.86	2.82	54,083	962K
Stackoverflow	545,195	96,678	1,301,942	2.39	13.47	2,922,148	26B
Twitter	175,214	530,418	1,890,661	10.79	3.56	5,102,542	13B
Imdb	685,568	186,414	2,715,604	3.96	14.57	1,809,175	1B
Actor2	303,617	896,302	3,782,463	12.46	4.22	3,761,666	2B
Amazon	2,146,057	1,230,915	5,743,258	2.68	4.67	5,702,211	108B
DBLP	1,953,085	5,624,219	12,282,059	6.29	2.18	1,281,831	85M

- MBAG (Algorithm 3) is directly based on the maximal biclique adjacency graph (also abbreviated as MBAG). It first enumerates all maximal bicliques, then constructs the maximal biclique adjacency graph using these bicliques, and finally traverses this adjacency graph to compute the BCPC.

We then implement the following partial-BCPC based solution and  $(\alpha, \beta)$ -biclique based solution.

- PBCPC (Algorithm 4) is the basic partial-BCPC based solution. It first enumerates maximal bicliques, and builds the MBE tree. For each  $(\alpha, \beta)$ -node in the MBE tree, it identifies all maximal bicliques associated with that node and connects them into a single partial-BCPC. It then traverses the MBAG reduced by the partial-BCPCs to obtain the final BCPC.
- PBCPC+ (Algorithm 5) is an improved version of PBCPC. The key difference lies in its use of stop-labels to reduce the search overhead when identifying all maximal bicliques associated with an  $(\alpha, \beta)$ -node.
- Biclique (Algorithm 6) is the basic solution based on  $(\alpha, \beta)$ -biclique enumeration. Its core idea is to enumerate all  $(\alpha, \beta)$ -bicliques, and for each such biclique, connect all maximal bicliques that contain it in order to construct the BCPC.
- BicliqueM (Algorithm 7) is an improved version of Biclique. It prunes  $(\alpha, \beta)$ -biclique enumeration branches by checking whether the maximal bicliques associated with the currently enumerated biclique already belong to the same BCPC.
- BicliqueP (Algorithm 8) is also an improved version of Biclique. Unlike BicliqueM, it leverages the method from PBCPC+ to compute partial-BCPCs in advance, thereby identifying some maximal bicliques that already belong to the same BCPC. This enables a stronger pruning effect.

**Parameters.** There are two parameters used in the experiments:  $\alpha$  and  $\beta$ . Their default values are both set to 4, and the possible values they can take are 2, 4, 6, and 8.

All the algorithms are implemented in C++. The experiments are conducted on a system running Ubuntu 20.04.4 LTS with an AMD Ryzen 3900X 2.2GHz CPU and 256GB of memory.

### 6.2 Experimental Results

**Exp-1: Efficiency of various algorithms.** In this experiment, we evaluated the performance of all algorithms across all datasets, and the results are presented in Table 2.

Overall, MBAG and PBCPC algorithms perform the worst in terms of runtime across most datasets and  $\alpha, \beta$ . This is mainly because the MBAG algorithm traverses the original maximal

**Table 2: Performance (s) of all algorithms on all datasets. The best and sub-best results are bold and underlined respectively. “-” denotes out of time ( $\geq 7$  days)**

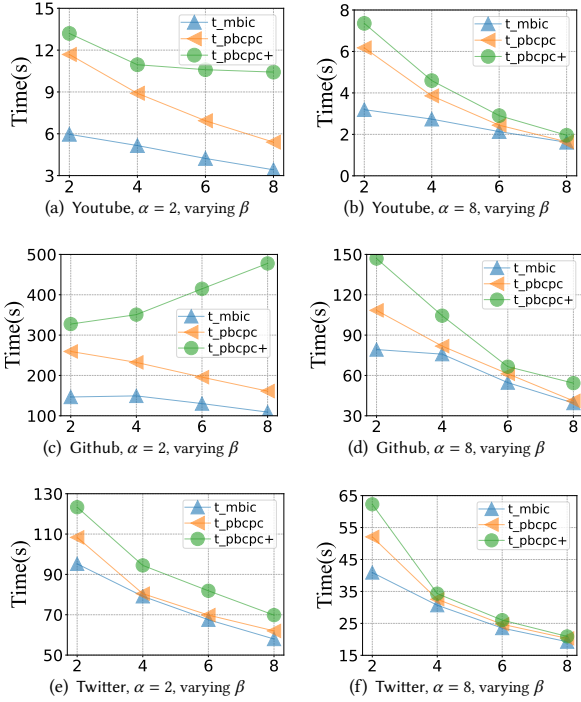
Datasets		Youtube				Bookcrossing				Github				Citeseer				Stackoverflow			
$\alpha$	$\beta$	2	4	6	8	2	4	6	8	2	4	6	8	2	4	6	8	2	4	6	8
2	Algo																				
	MBAG	13,376	11,492	6,072	1,902	35	2	0.9	0.7	-	-	-	-	1	0.8	0.7	0.6	32,598	17,967	4,287	844
	PBCPC	4,651	7,283	6,942	3,207	42	3	1	1	-	-	-	-	1	0.8	0.7	0.6	30,051	25,220	8,811	2,177
	PBCPC+	197	1,914	2,787	1,484	20	3	1	0.8	44,276	-	-	-	1	0.9	0.8	0.6	2,564	7,144	3,467	987
	Biclique	135	358	2,995	-	3	26	93,773	-	-	-	-	-	1	1	64	6,999	2,038	3,118	-	-
4	BicliqueM	59	70	43	22	2	1	0.8	0.6	24,581	-	-	-	1	0.8	0.7	0.6	486	208	119	80
	BicliqueP	34	43	31	20	3	2	1	1	7,718	34,478	33,948	14,272	1	0.9	0.8	0.6	339	159	121	103
	MBAG	13,113	11,358	5,941	1,793	3	0.7	0.5	0.4	-	-	-	-	0.1	0.08	0.07	0.06	27,582	14,496	3,308	544
	PBCPC	7,748	9,568	7,802	3,197	4	0.9	0.6	0.5	-	-	-	-	0.1	0.09	0.08	0.06	37,632	25,336	7,496	1,476
	PBCPC+	1,450	2,964	3,404	1,618	4	0.8	0.6	0.5	-	-	-	-	0.1	0.08	0.07	0.06	13,222	9,955	3,857	864
6	Biclique	3,754	5,445	5,522	2,877	2	1	0.8	0.6	-	-	-	-	0.1	0.1	0.1	0.3	1,210	5,446	748	11,595
	BicliqueM	54	235	140	50	1	0.8	0.6	0.5	-	-	-	412,886	0.1	0.09	0.07	0.06	306	539	199	56
	BicliqueP	28	66	49	24	1	0.9	0.6	0.5	11,483	145,121	424,317	157,465	0.1	0.09	0.08	0.07	190	159	76	34
	MBAG	10,936	9,409	4,635	1,312	1	0.6	0.5	0.4	-	-	-	-	0.03	0.03	0.03	0.03	9,142	4,150	561	45
	PBCPC	7,736	8,370	5,855	2,080	1	0.7	0.5	0.4	-	-	-	-	0.05	0.04	0.04	0.04	12,482	6,893	1,134	89
8	PBCPC+	2,500	3,622	3,133	1,121	1	0.7	0.5	0.4	-	-	-	-	0.04	0.03	0.04	0.03	5,758	3,833	775	76
	Biclique	-	3,289	34,084	3,352	36	0.9	0.5	0.4	-	-	-	-	0.07	0.04	0.04	0.04	-	1,764	2,724	26
	BicliqueM	44	165	178	35	1	0.6	0.5	0.4	13,182	444,343	-	55,394	0.04	0.04	0.04	0.04	90	131	40	12
	BicliqueP	24	57	56	16	1	0.7	0.5	0.4	9,450	141,003	158,184	20,785	0.05	0.04	0.04	0.04	75	72	22	10
	MBAG	7,236	6,102	2,832	744	1	0.5	0.4	0.3	-	-	-	-	0.03	0.03	0.03	0.03	1,377	398	21	7
10	PBCPC	5,585	5,707	3,596	1,241	1	0.6	0.4	0.3	-	-	-	-	0.03	0.03	0.03	0.03	1,721	592	32	6
	PBCPC+	2,327	2,878	1,991	713	1	0.6	0.4	0.3	-	-	-	-	0.03	0.03	0.03	0.03	1,110	447	29	7
	Biclique	-	109,345	4,728	8,555	2,315	6	0.4	0.3	-	-	401,158	23,651	0.09	0.04	0.04	0.04	-	77	20	14
	BicliqueM	33	117	101	38	1	0.6	0.4	0.3	5,219	146,064	103,754	2,324	0.03	0.03	0.03	0.03	30	22	9	7
	BicliqueP	19	45	33	13	1	0.6	0.4	0.3	4,066	61,855	36,416	1,252	0.04	0.04	0.04	0.03	30	16	9	6
Datasets		Twitter				Imdb				Actor2				Amazon				DBLP			
$\alpha$	$\beta$	2	4	6	8	2	4	6	8	2	4	6	8	2	4	6	8	2	4	6	8
2	Algo																				
	MBAG	11,652	5,290	1,817	657	859	706	622	409	7,303	6,650	5,217	3,502	16,282	3,974	580	137	18	12	10	9
	PBCPC	47,374	24,237	6,055	2,335	323	188	293	418	1,820	2,285	2,377	2,041	20,703	6,687	1,297	342	28	19	15	13
	PBCPC+	10,580	11,843	3,162	1,337	102	58	90	160	266	524	772	780	2,413	3,067	706	228	23	15	12	10
	Biclique	685	5,339	4,584	-	55	147	58,284	-	190	875	39,764	-	964	727	-	-	20	176	296,803	-
4	BicliqueM	510	777	208	108	35	26	23	21	88	56	48	47	618	157	79	55	17	13	10	9
	BicliqueP	518	402	152	94	42	33	28	25	79	51	44	39	590	176	108	82	25	19	16	14
	MBAG	4,339	2,737	1,236	457	762	687	614	403	7,107	6,586	5,178	3,468	7,856	1,720	227	57	7	5	4	3
	PBCPC	11,436	6,764	3,176	1,482	408	355	446	514	1,622	2,251	2,462	2,125	9,234	2,496	427	100	8	5	4	3
	PBCPC+	6,282	4,237	2,004	1,001	99	74	121	185	201	516	794	840	4,219	1,508	304	78	8	5	4	4
6	Biclique	-	12,028	20,544	242,804	92	1,805	1,335	25,431	626	4,996	29,168	526,141	534	2,687	474	9,676	12	7	29	1,834
	BicliqueM	134	129	94	87	22	21	26	21	57	131	51	44	303	161	61	39	8	5	4	3
	BicliqueP	134	100	69	64	23	18	16	15	47	48	34	31	280	111	61	42	9	5	4	3
	MBAG	2,184	1,596	891	298	623	600	536	340	6,859	6,376	5,085	3,343	1,576	349	63	32	3	2	2	2
	PBCPC	3,124	2,404	1,677	690	557	551	605	568	1,988	2,585	2,583	2,270	1,669	499	97	44	3	2	2	2
8	PBCPC+	2,666	1,469	1,073	506	169	164	204	237	326	641	933	955	1,035	331	71	36	3	3	2	2
	Biclique	-	-	134,078	27,608	1,748	835	11,021	10,968	194,453	22,912	49,668	137,995	33,212	1,595	525	806	10,632	6	17	30
	BicliqueM	90	81	68	52	17	17	22	23	51	121	62	52	171	80	44	32	3	2	2	2
	BicliqueP	95	58	51	41	19	14	14	13	40	43	38	32	213	79	47	33	3	2	2	2
	MBAG	1,482	1,166	651	210	400	388	340	187	6,381	5,838	4,529	2,958	445	104	41	28	1	1	1	1
10	PBCPC	1,479	1,297	994	369	440	437	450	331	2,631	2,922	2,867	2,398	539	143	54	34	1	1	1	2
	PBCPC+	880	729	669	271	187	186	209	193	625	894	1,093	1,023	338	101	45	31	2	1	1	1
	Biclique	-	-	-	190,333	355,545	2,375	1,985	11,003	-	-	-	-	-	29,402	3,606	355	-	25	61	93
	BicliqueM	74	65	53	36	15	13	16	16	46	100	151	66	143	59	38	28	1	1	1	1
	BicliqueP	79	45	36	28	16	12	12	11	35	38	46	36	188	64	40	29	2	1	1	1

biclique adjacency graph (MBAG), which is often very large (see Table 1). Although the PBCPC algorithm attempts to reduce the size of the MBAG by computing partial-BCPCs, these partial-BCPCs are not sufficiently comprehensive, leading to limited pruning effectiveness on the MBAG (we will explore the effectiveness of these partial-BCPCs in Exp-3).

PBCPC+ performs slightly better than MBAG and PBCPC, especially when  $\alpha$  and  $\beta$  are small. For instance, on dataset Youtube, when  $\alpha = \beta = 2$ , PBCPC+ outperforms MBAG and PBCPC by more than one order of magnitude. This is because when  $\alpha$  and  $\beta$  are small, there is a larger number of maximal bicliques that are larger than  $(\alpha, \beta)$ -bicliques, which increases the scale of maximal biclique adjacency graph. In such cases, the effect of PBCPC+ using partial-BCPC to reduce the size of the adjacency graph is more prominently demonstrated.

Biclique achieves relatively good performance only when both  $\alpha$  and  $\beta$  are small—for example, on datasets Youtube, Bookcrossing, and DBLP with  $\alpha = \beta = 2$ . When  $\alpha$  or  $\beta$  increases, the performance of Biclique drops sharply. This is because the number of  $(\alpha, \beta)$ -bicliques grows exponentially as  $\alpha$  and  $\beta$  increase.

BicliqueM and BicliqueP are the two optimal algorithms in almost all cases, and they can outperform MBAG by nearly three orders of magnitude. For example, on dataset Youtube, when  $\alpha = \beta = 2$ , BicliqueP only takes 34 seconds, while MBAG requires 13,376 seconds. Furthermore, BicliqueP performs better than BicliqueM, and this advantage is more pronounced on dataset Github (under many parameters, only BicliqueP can complete within 7 days, i.e., 604,800 seconds). This is because BicliqueP uses partial-BCPC for pruning, which has stronger pruning capability compared to maximal biclique (see Exp-4).

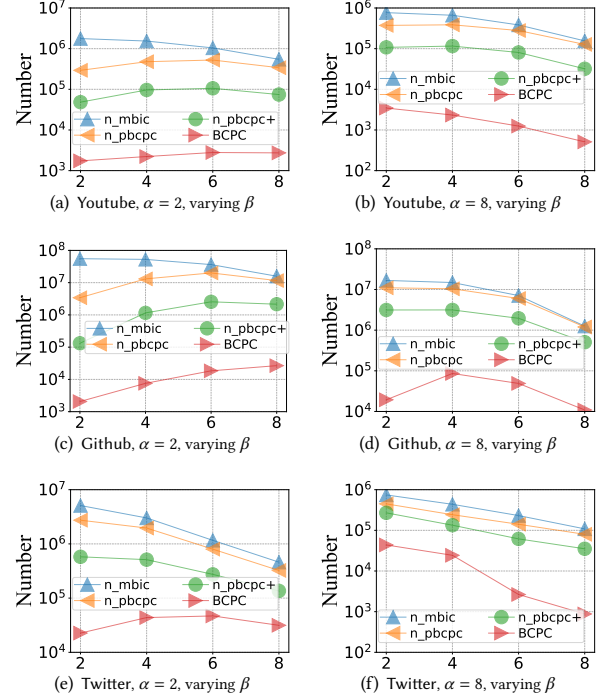


**Figure 4: Time spent on maximal biclique enumeration ( $t\_mbic$ ), partial-BCPC computation (including maximal biclique enumeration as a necessary step) in PBCPC ( $t\_pbcpc$ ) and PBCPC+ ( $t\_pbcpc+$ )**

**Exp-2: Overhead of computing partial-BCPC.** This experiment evaluates the overhead of computing partial-BCPCs (with maximal biclique enumeration included, which is a necessary step). Specifically, we compare the time spent on computing partial-BCPCs in the algorithms PBCPC and PBCPC+, and also compare it against the time spent on maximal biclique enumeration. The experimental results are shown in Figure 4.

As shown in Figure 4, although computing partial-BCPCs introduces additional overhead on top of maximal biclique enumeration, the added cost is manageable. For example, on datasets Youtube and Twitter,  $t\_pbcpc$  and  $t\_pbcpc+$  are only slightly higher than the time spent on maximal biclique enumeration alone ( $t\_mbic$ ). For the Github dataset, due to the larger number of maximal bicliques (see Table 1) and larger scale of the MBE tree,  $t\_pbcpc+$  is no longer close to  $t\_mbic$  and  $t\_pbcpc$ . However, this overhead is meaningful—our best-performing algorithm, BicliqueP, is built upon the partial-BCPC computed in PBCPC+ (the time spent is also  $t\_pbcpc+$ ). Notably, BicliqueP is the only algorithm that successfully passed all tests on the Github dataset.

**Exp-3: Effectiveness of partial-BCPC in partial-BCPC based solution.** This experiment investigates the effectiveness of partial-BCPC in PBCPC and PBCPC+. Specifically, we compare the number of partial-BCPCs obtained by the two algorithms with the number of maximal bicliques and the number of BCPCs. The results are shown in Figure 5. We can see that in all cases, the inequality in Theorem 3 and 7 holds:  $n\_mbic \geq n\_pbcpc \geq n\_pbcpc+ \geq BCPC$ . We can also observe that compared to  $n\_mbic$  and  $n\_pbcpc$ ,  $n\_pbcpc+$  is much closer to BCPC, and in some cases, it is more

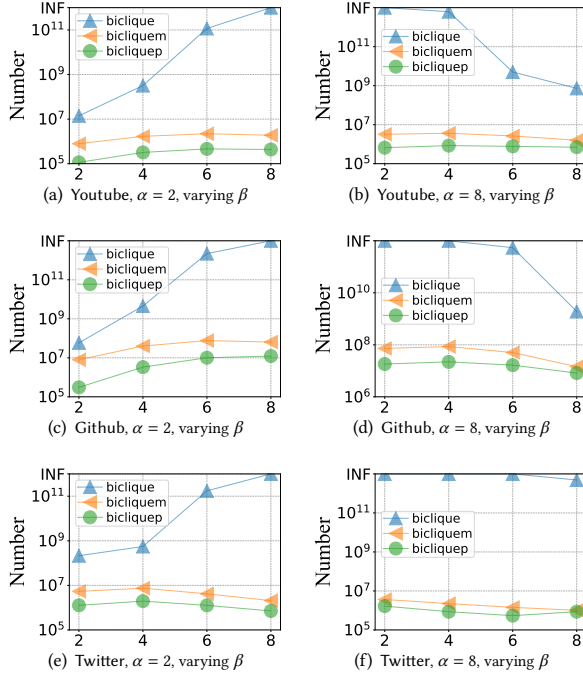


**Figure 5: Number of maximal bicliques ( $n\_mbic$ , where the size of  $X$  set and  $Y$  set is at least  $\alpha$  and  $\beta$  respectively), partial-BCPCs obtained by PBCPC ( $n\_pbcpc$ ) and PBCPC+ ( $n\_pbcpc+$ ), and number of BCPCs**

than an order of magnitude smaller than  $n\_mbic$  (see Figure 5 (a), (c)). In both the PBCPC and PBCPC+ algorithms, vertices in the MBAG are regrouped according to the partial-BCPCs, meaning that the number of vertices in the reduced MBAG equals the number of partial-BCPCs. As a result, PBCPC+ outperforms PBCPC on nearly all datasets.

**Exp-4: Effectiveness of partial-BCPC in  $(\alpha, \beta)$ -biclique based solution.** This experiment investigates the pruning effectiveness of maximal biclique and partial-BCPC in the  $(\alpha, \beta)$ -biclique enumeration process. Figure 6 shows the number of enumerated nodes in the  $(\alpha, \beta)$ -biclique enumeration trees of three algorithms: Biclique, BicliqueM, and BicliqueP. It can be observed that in Biclique (without any pruning), the number of nodes in the enumeration tree is more than four orders of magnitude higher than that in the other two algorithms, which sufficiently demonstrates the pruning capabilities of maximal biclique and partial-BCPC in  $(\alpha, \beta)$ -biclique enumeration process. Among BicliqueM and BicliqueP, the partial-BCPC in BicliqueP exhibits the strongest pruning capability, as the number of enumerated nodes in BicliqueP is another order of magnitude smaller than that in BicliqueM (e.g., when  $\beta = 2$  in Figure 6 (c)). This explains why BicliqueP achieves the best performance in most cases in Table 2, particularly with the dataset Github.

**Exp-5: Scalability.** This experiment explores the scalability of our best algorithm, BicliqueP. Table 3 presents the performance of BicliqueP on Github and four other large datasets. Specifically, we sampled 20%, 40%, 60%, and 80% of the edges from each dataset respectively. It can be observed that as the dataset size increases, the



**Figure 6: Number of enumerated nodes in the  $(\alpha, \beta)$ -biclique enumeration trees of three algorithms: Biclique (biclique), BicliqueM (bicliquem), and BicliqueP (bicliquep)**

**Table 3: Scalability test of PBCPC+ (s)**

	Github	Imdb	Actor2	Amazon	DBLP
20%	5,779.7	3.4	1.2	8.3	0.8
40%	72,202.2	7.8	5.7	54.5	1.4
60%	119,608.0	13.5	14.7	85.2	2.8
80%	142,445.8	17.5	25.8	107.3	4.5
100%	145,121.7	18.2	49.0	111.2	5.8

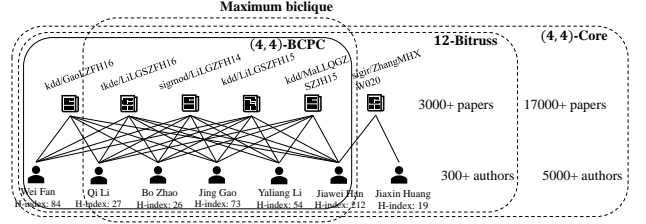
**Table 4: Memory usage (MB) of various algorithms**

	MBAG	PBCPC	PBCPC+	Biclique	BicliqueM	BicliqueP
Github	10,051	26,965	26,418	9,904	<b>9,904</b>	22,158
Imdb	<b>233</b>	364	361	260	260	343
Actor2	<b>409</b>	725	720	449	449	639
Amazon	<b>711</b>	1,228	1,207	797	772	1,130
DBLP	935	957	970	1,200	<b>935</b>	940

growth of running time of BicliqueP is manageable, demonstrating that BicliqueP has excellent scalability.

**Exp-6: Memory.** Table 4 presents the memory usage of all algorithms on the largest four datasets and dataset Github, which contains the maximum number of maximal bicliques. It can be observed that across all datasets, the three algorithms—MBAG, Biclique, and BicliqueM—consistently consume the least memory. This is because these algorithms do not require storing the MBE tree. However, although our best algorithm, BicliqueP, also stores the MBE tree, its memory consumption remains manageable.

**Exp-7: Case study.** We conduct a case study on the DBLP dataset. This dataset includes papers from major journals (TODS, TOIS, TKDE, VLDBJ) and conferences (SIGMOD, SIGKDD, ICDE, SIGIR, VLDB) in the database field, where vertices represent papers and authors respectively, and an edge  $(u, v)$  indicates that  $v$  is one of



**Figure 7: Case study on DBLP**

the authors of  $u$ . We compare  $(\alpha, \beta)$ -BCPC with maximum biclique,  $k$ -bitruss [28], and  $(\alpha, \beta)$ -core [19].

Given a bipartite graph  $G = (U, V, E)$ ,  $(\alpha, \beta)$ -core of  $G$  is a maximal subgraph of  $G$ ,  $G_1 = (U_1, V_1, E_1)$ , that  $\forall u \in U_1, |Nei_{G'}(u)| \geq \alpha, \forall v \in V_1, |Nei_{G'}(v)| \geq \beta$ .  $k$ -bitruss is a maximal subgraph of  $G$ ,  $G_2 = (U_2, V_2, E_2)$ , that each edge in  $E_2$  is contained in at least  $k$  butterflies (butterfly is a subgraph with four vertices:  $u, w \in U_2, v, x \in V_2, (u, v) \in E_2, (u, x) \in E_2, (w, v) \in E_2, (w, x) \in E_2$ ).

In this case study,  $\alpha = 4, \beta = 4, k = 12$ . By Definition 4, the smallest maximal bicliques in  $(4, 4)$ -BCPC are either  $(5, 4)$ -bicliques or  $(4, 5)$ -bicliques, and the edges within these maximal bicliques exist in at least  $3 \times 4 = 12$  butterflies, thus, we set  $k = 12$  for  $k$ -bitruss.

Figure 7 presents the results containing Jiawei Han across all models. Among them, the result from maximum biclique is the smallest. Compared with  $(4, 4)$ -BCPC, it lacks one author, Wei Fan, who is one of Jiawei Han's collaborators on Google Scholar. Although 12-bitruss and  $(4, 4)$ -core include the results of  $(4, 4)$ -BCPC, they also contain a large number of other vertices, making the usability of these two models far inferior to that of  $(4, 4)$ -BCPC.

## 7 Related Work

The research related to BCPC is the most relevant to our work. Initially, [15] proposed the BCPC model and introduced a method to detect BCPCs by traversing the maximal biclique adjacency graph (MBAG). Subsequently, [27] developed an online web toolkit, implementing the approach presented in [15]. In the context of social network analysis, [12] applied BCPC to identify clusters of users and entities. Regarding Wikipedia networks, [14] leveraged BCPC to explore the relationships between editors and articles, revealing that topics inherently aggregate editors. [16] applied BCPC to enterprise networks to analyze interaction patterns between hosts and users, thereby supporting the detection of abnormal user behaviors. Meanwhile, [13, 33] utilized BCPC to study the relationships between students and online learning resources, delving into the temporal evolution patterns between students and courses. Additionally, [7] investigated the problem of personalized BCPC search based on the BCPC model.

Our work is also related to maximal biclique enumeration and  $(p, q)$ -biclique enumeration. Existing methods for maximal biclique enumeration, a foundational task supporting BCPC model implementation, are primarily rooted in the idea of enumerating the powerset of vertex set [32]. Building on this framework, [1] introduced the pivot pruning strategy, which effectively reduces redundant search space and improves the efficiency. To further optimize performance, [6] proposed a method that optimizes the vertex enumeration order, enhancing computational speed. Additionally, [10] developed a unified enumeration framework suitable for hereditary cohesive subgraphs. This framework achieves state-of-the-art performance in maximal biclique enumeration

problem, providing a more efficient and generalizable solution for related tasks. Key works on  $(p, q)$ -biclique enumeration include [24, 30]. [24] uses  $(p, q)$ -biclique enumeration only as a sub-process for broader graph analysis, while [30] focuses specifically on optimizing this enumeration task. Additionally, studies like [31] address the exact and approximate counting of  $(p, q)$ -biclques.

There are also several cohesive subgraph models relevant to our work, such as  $(\alpha, \beta)$ -core and  $k$ -bitruss. For  $(\alpha, \beta)$ -core, key works include [18–20], which focus on efficient computation and distributed implementation of this structure. In terms of  $k$ -bitruss, studies such as [17, 28, 34] contribute to optimizing its decomposition and analysis processes. For maximum biclique, [22, 29] propose efficient methods for its identification. Beyond specific subgraph structures, modularity-based modeling approaches (e.g., [2, 4]) and label propagation methods (e.g., [21]) are also studied.

## 8 Conclusion

In this work, we tackle the problem of biclique percolation community (BCPC) detection in bipartite network. Confronting the high time complexity of existing BCPC detection methods, which stems from the potentially massive maximal biclique adjacency graph (MBAG), we introduce two innovative solutions. First, the partial-BCPC based solution reduces redundant computations by grouping maximal bicliques into incomplete BCPCs, lessening the burden of explicitly representing the entire MBAG. Second, our novel method based on  $(\alpha, \beta)$ -biclque enumeration detects BCPCs by enumerating all  $(\alpha, \beta)$ -biclques and connecting maximal bicliques that share such bicliques. Leveraging partial-BCPC, this approach significantly prunes the  $(\alpha, \beta)$ -biclque enumeration space. Experimental results demonstrate that our methods achieve remarkable performance, outperforming existing approaches by nearly three orders of magnitude. These advancements not only enhance the efficiency of BCPC detection but also pave the way for more scalable analysis of bipartite networks in diverse domains, including social networks and recommendation systems.

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