



# Efficient Maximal Biclique Enumeration on GPUs

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## ABSTRACT

Maximal biclique enumeration (MBE) in bipartite graphs is an important problem in data mining with many real-world applications. All existing solutions for MBE are designed for CPUs. Parallel MBE algorithms for GPUs are needed for MBE acceleration leveraging its many computing cores. However, enumerating maximal bicliques using GPUs has three main challenges including large memory requirement, thread divergence, and load imbalance. In this paper, we propose GMBE, the first highly-efficient GPU solution for the MBE problem. To overcome the challenges, we design a node-reuse approach to reduce GPU memory usage, a pro-active pruning method using the vertex's local neighborhood size to alleviate thread divergence, and a load-aware task scheduling framework to achieve load balance among threads within GPU warps and blocks. Our experimental results show that GMBE on an NVIDIA A100 GPU can achieve 70.6× speedup over the state-of-the-art parallel MBE algorithm PARMBE on a 96-core CPU machine.

## CCS CONCEPTS

• **Mathematics of computing** → **Graph enumeration**; • **Computer systems organization** → **Multicore architectures**; • **Information systems** → **Data mining**.

## KEYWORDS

Maximal biclique enumeration, GPU

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## 1 INTRODUCTION

A *bipartite graph*  $G = (U, V, E)$  contains two disjoint vertex sets  $U$  and  $V$ , where an edge  $e \in E$  only occurs between two vertices in  $U$  and  $V$ , respectively. A *biclique* is a complete bipartite graph, i.e., there exists an edge between two vertices if and only if the two vertices are in different vertex sets. A *maximal biclique* in  $G$  is a subgraph of  $G$ , which is a biclique and can not be further enlarged to form a larger biclique. Maximal biclique enumeration (MBE) aims to find all maximal bicliques in  $G$ .

MBE is important in bipartite graph analysis, with widespread applications, such as anomaly detection in e-commerce networks [30, 35], social recommendation in social networks [29], gene expression analysis in expression datasets [34, 39], and GNN information aggregation [38]. Consider an example in an e-commerce network, where the purchasing relationships can be modeled by a bipartite graph. It is suspicious for a large group of customers to buy a set of products together because online sellers are likely to make fake purchases through illegal platforms to improve their credibility and positive ratings [10, 30, 35]. We can identify all these suspicious groups by enumerating all the maximal bicliques in the network, and then detect them.

Over the past few decades, many MBE algorithms have been proposed to speed up the enumeration of all maximal bicliques in bipartite graphs [8, 9, 14, 18, 22, 29, 32, 39]. A mainstream approach is to use the set enumeration tree [33] to recursively enumerate all candidate subgraphs, and then judge whether they are maximal bicliques. The enumeration space of this method is a powerset of  $V$  or  $U$ , so the computational overhead is very high [19], especially for large graphs. Therefore, many efforts are made to reduce the enumeration space using pruning [8, 14] and vertex ordering [14, 39]. However, they only achieve limited speedup for not exploring the parallelism of multi-core CPUs. Other works design parallelization strategies for the multi-core CPUs to speed up the enumeration process [18]. However, their performance speedup is still constrained by the limited parallelism of CPUs. For instance, the state-of-the-art parallel MBE algorithm PARMBE [18] costs over 40 minutes to finish running MBE on a medium-scale bipartite graph Github [25] on a 96-core CPU machine. In contrast, our GPU-based solution can reduce the running time to 132 seconds on an NVIDIA A100 GPU [4] because GPUs offer much higher computational throughput and parallelism than CPUs.

There are three major challenges in the design of a highly efficient MBE algorithm for GPUs. First, the existing MBE algorithms require large memory space and frequent memory allocations to store the intermediate data of each enumerated subgraph, thus they cannot efficiently run on GPUs with limited memory capacity and high dynamic memory allocation overhead [37], due to the severe shortage of memory resources.

Second, the performance of existing MBE algorithms suffers from irregular computation [13] while GPUs are suitable to perform regular computation with high parallelism. Specifically, different GPU threads in the same warp in the MBE algorithm may take different execution paths to access different vertex neighbors, causing the thread divergence problem [15]. GPUs will serialize these diverged thread operations [1], resulting in low thread utilization and poor memory access efficiency.

Lastly, existing MBE algorithms have a serious load imbalance problem on GPUs. The reason is that the maximal biclique sizes are varied significantly with real-world graphs for the power-law distribution of vertex degrees. As a result, threads assigned to each GPU core have different running times. When load imbalance happens, thousands of GPU threads have to wait for the slowest one to complete, causing degraded GPU performance.

Many existing studies have used GPUs to improve the performance of other graph enumeration problems, like maximal clique enumeration [36] and graph pattern mining [15]. The optimizations include data graph partitioning [23], two-level parallelism [17], adaptive buffering [16], hybrid order on GPU [15], etc. However, none of them is efficient for MBE using GPUs. This is because the enumerated subgraphs for MBE generally contain much more vertices than those in other graph enumeration problems. For instance, it may enumerate maximal bicliques comprising several to thousands of vertices, while the triangle counting algorithm only considers subgraphs with three vertices. The larger subgraphs generated at runtime require larger memory and computation costs and lead to more serious problems of large memory requirement, thread divergence, and load imbalance mentioned above.

To address all the challenges and achieve highly efficient maximal biclique enumeration on GPUs, we design the first GPU-based MBE algorithm (GMBE) considering the characteristics of GPU architecture and MBE computation pattern. Specifically, first, we replace the existing recursion with a stack-based iteration and reuse the memory of the root node throughout the iteration process. This approach effectively reduces memory usage because it eliminates the need to allocate additional memory space for new nodes. Second, we use the local neighborhood size of a vertex to reduce the enumeration space. The new pruning approach can reduce the number of sub-trees without visiting the nodes. Thus, it can significantly reduce thread divergence. Finally, GMBE carefully manages the size of subtrees assigned to GPU threads and schedules the tasks using two-level queues to achieve load balance within GPU warps and blocks leveraging persistent thread programming models for GPUs.

We adopt the fast CUDA primitives [28] to implement the GMBE prototype and conduct extensive experiments on real-world datasets to demonstrate the efficiency of GMBE. Our experimental results show that GMBE on an NVIDIA A100 GPU can achieve up to 70.6× speedup over the state-of-the-art parallel MBE algorithm PARMBE on a 96-core CPU machine.

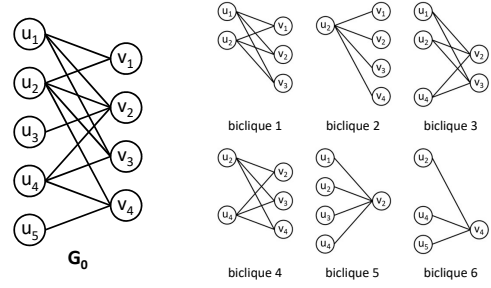


Figure 1: A bipartite graph  $G_0$  containing 6 maximal bicliques.

## 2 BACKGROUND

In this section, we briefly review the state-of-the-art MBE algorithms on CPUs and introduce the compute unified device architecture (CUDA) in modern GPUs.

### 2.1 MBE Algorithms on CPUs

**Notation definitions.** Given a bipartite graph  $G(U, V, E)$ .  $U$  and  $V$  are two disjoint vertex sets in  $G$ .  $E$  is the edge set in  $G$  and  $E \subseteq U \times V$ . For each vertex  $u$  in  $U$ ,  $N(u)$  denotes the neighbors of  $u$ , i.e.,  $N(u) = \{v \mid (u, v) \in E\}$ .  $N_2(u)$  denotes the 2-hop neighbors of  $u$ , i.e.,  $N_2(u) = \bigcup_{v \in N(u)} N(v) - \{u\}$ . For a vertex set  $X$ ,  $\Gamma(X)$  denotes the common neighbors of vertices in  $X$ , i.e.,  $\Gamma(X) = \bigcap_{u \in X} N(u)$ .  $\Delta(X)$  is the maximum degree of vertices in  $X$ , i.e.,  $\Delta(X) = \max_{u \in X} |N(u)|$ .  $\Delta_2(X)$  is the maximum 2-hop degree of vertices in  $X$ , i.e.,  $\Delta_2(X) = \max_{u \in X} |N_2(u)|$ . We have a symmetrical definition for each vertex  $v$  in  $V$ . A **biclique** is a vertex set pair  $(L, R)$  in  $G$  s.t.,  $L \subseteq U$  and  $R \subseteq V$  and  $\forall u \in L, v \in R, (u, v) \in E$ . A biclique  $(L, R)$  is a **maximal biclique** if there is no biclique  $(L', R')$  such that  $(L \cup R) \subset (L' \cup R')$ . For instance, Figure 1 shows a bipartite graph  $G_0$  and all maximal bicliques in  $G_0$ .

**Problem statement.** The MBE problem aims to enumerate all maximal bicliques in a bipartite graph.

**Baseline solution.** To efficiently solve the MBE problem, recent works [8, 14, 18, 22, 24, 29, 32, 39] **recursively** run a backtracking procedure to generate the powerset of  $V$  using a set enumeration tree [33] and then obtain all maximal bicliques correspondingly. In most works [8, 18, 29, 32], each enumeration node is represented as a 3-tuple  $(L, R, C)$ , where  $L \subseteq U$  and  $R, C \subseteq V$ .  $R$  and  $C$  are disjoint and used to generate the powerset of  $V$ .  $R$  stores the current subset of  $V$ , while  $C$  stores the candidate vertices for expanding  $R$ .  $(L, R)$  is the corresponding biclique where  $L = \Gamma(R)$ . In the following, we illustrate the basic recursive procedure for each enumeration node in existing works using Algorithm 1. Then, we provide a detailed example to demonstrate the application of Algorithm 1 to the bipartite graph  $G_0$  in Figure 1.

The procedure starts at a root node initialized as  $(U, \emptyset, V)$ . In each enumeration node, each candidate vertex  $v_c \in C$  is traversed sequentially (line #2) to generate a new biclique  $(L', R')$  (line #3). The parent node allows expanding  $R$  with untraversed candidate vertices in  $C$  (lines #5,6). The new candidate set  $C'$  contains all vertices in  $C - R'$  that connect with any vertex in  $L'$  (lines #7,8). The new biclique is maximal if and only if  $R'$  is equal to  $\Gamma(L')$  (line #9). All maximal bicliques are enumerated exactly once (line

**Algorithm 1: Recursive MBE Algorithm**


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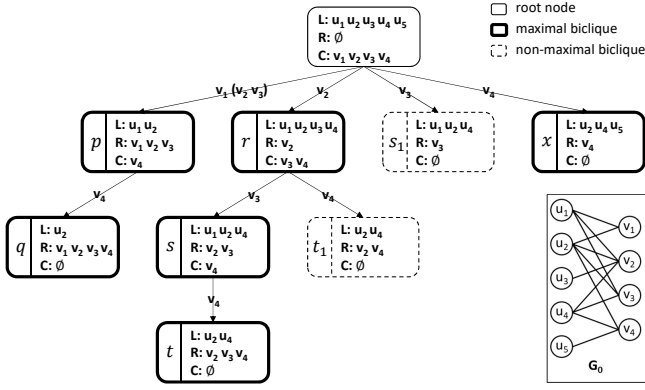
**Data:** Bipartite graph  $G(U, V, E)$   
**Input:** Set  $L \subseteq U$ , disjoint sets  $R, C \subseteq V$   
**Output:** All maximal bicliques

```

1 procedure recursively_search( $L, R, C$ ):
2   foreach  $v' \in C$  do
3      $L' \leftarrow L \cap N(v')$ ;  $R' \leftarrow R$ ;  $C' \leftarrow \emptyset$ ;
4     foreach  $v_c \in C$  do // Node generation
5       if  $L' \cap N(v_c) = L'$  then
6          $R' \leftarrow R' \cup \{v_c\}$ ;
7       else if  $L' \cap N(v_c) \neq \emptyset$  then
8          $C' \leftarrow C' \cup \{v_c\}$ ;
9     if  $R' = \Gamma(L')$  then // Maximality check
10      Output( $L', R'$ ) as a maximal biclique;
11      recursively_search( $L', R', C'$ ); // Recursion
12     $C \leftarrow C \setminus \{v'\}$ ;

```

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Figure 2: An enumeration tree for bipartite graph  $G_0$ .

#10). The procedure recursively generates new nodes in a depth-first search (DFS) manner to enumerate all maximal bicliques (line #11). After processing a new node, we remove  $v_c$  from the current candidate set  $C$  (line #12). Besides, some works [14, 39] use a set  $Q$  to keep traversed candidate vertices for accelerating the node checking in line #9. Similar to many existing works [8, 18, 29, 32], we omit the set  $Q$  to reduce the memory consumption.

**EXAMPLE 2.1.** Figure 2 depicts an enumeration tree for a bipartite graph  $G_0$  using Algorithm 1 without any optimization. The vertices on the edge between two nodes are used to expand  $R$  of the new node, including a traversed vertex  $v_c$  and the other untraversed candidate vertices in parenthesis. For presentation convenience, we always use the subscript to denote the corresponding vertex set of an enumeration node. For instance,  $L_{(p)}$  denotes  $L$  set of node  $p$ .

We start from the root node and recursively search the subspaces in a DFS manner by traversing candidate vertices in  $C$  following a pre-imposed order. By traversing  $v_1$ , we enter node  $p$ . We know  $L_{(p)} = L_{(root)} \cap N(v_1) = \{u_1, u_2, u_3, u_4, u_5\} \cap \{u_1, u_2\} = \{u_1, u_2\}$ . We then expand  $R_{(p)}$  with candidate vertices  $v_1, v_2$ , and  $v_3$  because  $\Gamma(L_{(p)}) \cap C_{(root)} = \{v_1, v_2, v_3\} \cap \{v_1, v_2, v_3, v_4\} = \{v_1, v_2, v_3\}$ . Node  $p$  generates a maximal biclique  $(L_{(p)}, R_{(p)})$  because  $R_{(p)} = \Gamma(L_{(p)})$ .  $v_4$  is in the

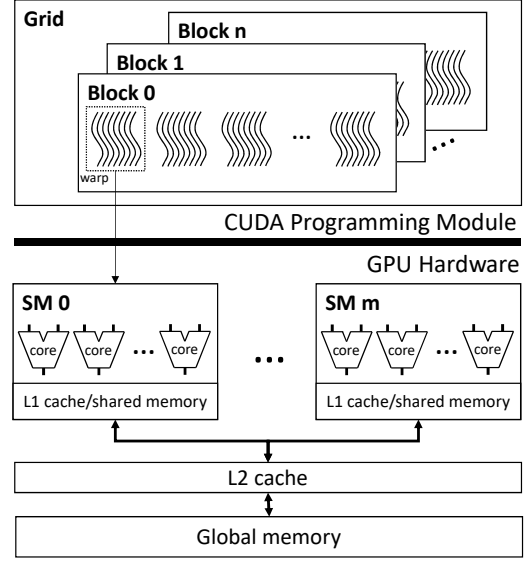


Figure 3: GPU architecture.

new candidate vertex set  $C_{(p)}$  because  $L_{(p)} \cap N(v_4) = \{u_1, u_2\} \cap \{u_2, u_4, u_5\} = \{u_2\} \neq \emptyset$ . Continuing this process, the root node can generate node  $s_1$  by traversing  $v_3$ . We then know  $L_{(s_1)} = \{u_1, u_2, u_4\}$  and  $R_{(s_1)} = (\Gamma(L_{(s_1)}) \cap C_{(root)}) \cup R_{(root)} = \{v_3\}$ . Compared to node  $s$ , node  $s_1$  generates a non-maximal biclique because the parent node of node  $s_1$  (i.e., the root node) fails to expand  $R_{(s_1)}$  with vertex  $v_2$  since  $v_2$  has been traversed by root node to generate node  $r$ . Other nodes can be generated similarly as shown in the figure.

**Recent optimizations.** To reduce the computational overhead, researchers mainly focus on reducing the enumeration space, using various vertex ordering and pruning approaches [8, 14, 39]. To achieve further speedup, other works parallelized MBE algorithms on multicore CPUs [18] or distributed architectures [32]. Specifically, existing parallel MBE algorithms distribute all vertices  $v$  in  $V$  across CPU threads, and each thread generates a subtree correspondingly using 1-hop and 2-hop neighbors of  $v$ . However, their performance speedup is constrained by the limited parallelism of CPUs. For instance, the state-of-the-art parallel MBE algorithm PARMBE [18] costs over 40 minutes to enumerate all maximal bicliques on a medium-scale bipartite graph Github (180k vertices and 440k edges) on a 96-core CPU machine, as shown in Section 6.2.

## 2.2 GPU Architecture and Programming

**GPU architecture.** Figure 3 shows the general architecture of a modern GPU. A GPU generally consists of a global memory, a shared L2 cache, and multiple streaming multiprocessors (SMs). Each SM contains an individual L1 cache, a programmable multi-bank shared memory, and multiple computing cores. A modern GPU can be equipped with thousands of lightweight cores in total, thus providing massive computing power. Unlike computing resources, memory resources on the GPU are relatively limited. For example, the recently popular NVIDIA A100 [4] can provide up to 6,912 cores, but only up to 80 GB of memory.

**CUDA programming model.** The CUDA (Compute Unified Device Architecture) programming model [1, 2] provides a parallel computing platform and a set of APIs that allows users to efficiently use GPUs for general-purpose processing. It adopts the SIMT (Single instruction, multiple threads) execution model [7] to manage numerous threads. CUDA divides GPU kernels into multiple grids, each consisting of multiple blocks. A block includes multiple threads and is assigned to an SM during execution. The SM groups 32 parallel threads into a warp and executes multiple warps concurrently. For each warp, all threads execute one common instruction at a time. In this way, thousands of GPU cores can efficiently work in parallel to achieve high performance.

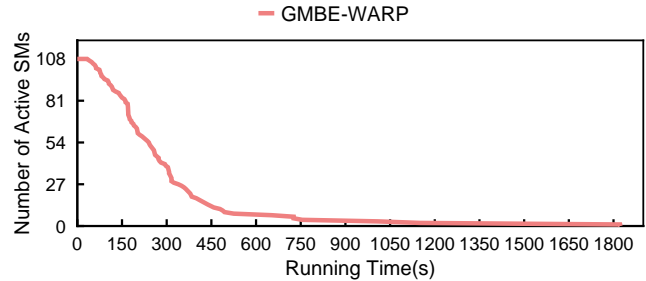
**GPU usage guidelines.** There are several notable guidelines for improving the efficiency of GPU programs. First, dynamic memory allocations on GPUs are expensive because many threads may allocate new memory at the same time, causing problems such as thread contention, synchronization overhead, and memory fragmentation [37]. Therefore, we should carefully manage the valuable GPU memory and avoid frequent memory allocations. Second, we should minimize the thread divergence, i.e., threads in a warp take different execution paths on the GPU. Because GPU has to serialize the different execution paths, thread divergence can severely degrade the execution performance [1]. Third, we should pay more attention to the load balancing among multiple cores on the GPU, because thousands of cores have to wait for the slowest thread to run on a lightweight core, which is costly.

### 3 CHALLENGES OF MBE ON GPUS

A naive approach to perform MBE on the GPU is to divide the whole enumeration tree into multiple subtrees and assign each subtree to an individual SM for execution. For example, for the enumeration tree in Figure 2, we can divide the whole tree into four subtrees, then assign SM 0 to conduct the execution for enumeration nodes  $p$  and  $q$ , and assign SM 1 to conduct the execution for enumeration nodes  $r$ ,  $s$ ,  $t$  and  $t_1$ , and so on. However, this naive approach faces the following three problems, thus making MBE on GPUs challenging.

#### 3.1 Large Memory Requirement

Existing MBE algorithms usually dynamically allocate memory for new enumeration nodes (lines #3-8 in Algorithm 1), which is very expensive on GPUs as mentioned in Section 2.2. To achieve high performance, a typical approach is to pre-allocate enough memory space on GPUs to accommodate all enumeration nodes before execution. In this case, each enumeration node requires  $O(|L|+|R|+|C|)$  memory bounded by  $O(\Delta(V) + \Delta_2(V))$ , and each subtree activates at most  $\Delta(V)$  nodes at the same time for backtracking. Therefore, the total amount of memory space that needs to be pre-allocated for each subtree traversal procedure is  $\Delta(V) \times (\Delta(V) + \Delta_2(V)) \times \text{sizeof}(\text{vertex})$ . For example, when using the NVIDIA A100 GPU with 40 GB memory to perform MBE on a real-world bipartite graph BookCrossing [25], the memory requirement for each subtree traversal procedure is  $13,601 \times (13,601 + 53,915) \times \text{sizeof}(\text{int}) \text{ B} = 3.67 \text{ GB}$ . We need more than  $108 \times 3.67 \text{ GB} = 397 \text{ GB}$  memory to fully utilize the 108 SMs, which exceeds the maximum memory space on the GPU (i.e., 40 GB), thus facing a severe memory shortage.



**Figure 4: Load imbalance for the MBE problem. 86 SMs waste over 1,458 seconds waiting for the slowest one on BookCrossing if we assign each enumeration tree to a warp based on our new algorithm GMBE as shown in Section 6.3.**

#### 3.2 Massive Thread Divergence

As mentioned in Section 2.2, thread divergence can significantly degrade thread performance on GPUs. In the case of MBE on GPUs, thread divergence becomes more prominent due to two reasons. First, during the enumeration process of Algorithm 1, threads within the same warp may traverse different vertices to generate new nodes (line #2) and use different neighbors to compute different sets of  $L$ ,  $R$ , and  $C$  (lines #3-9). This leads to each thread accessing different memory areas and executing different control flows. Second, several pruning techniques [8, 14, 39] have been proposed to reduce the enumeration space and accelerate the MBE process (refer to Section 2.1). For example, oomBEA [14] traverses the 2-hop neighbors of all candidate vertices to identify its defined batch pivots and discard enumerated branches that do not belong to those batch pivots. However, this further exacerbates thread divergence by generating more divergent threads through the traversal of 2-hop neighbors. Hence, it is challenging to mitigate thread divergence during enumeration and crucial to develop a GPU-friendly pruning approach.

#### 3.3 Load Imbalance

The parallel MBE algorithm is likely to generate severe imbalanced workloads on GPUs for two main reasons. First, the running time for processing each enumeration node in the enumeration tree varies greatly since nodes contain various numbers of candidate vertices. Second, the number of nodes in subtrees differs significantly because different maximal bicliques  $(L, R)$  contain various numbers of vertices in  $R$ . The enumeration tree grows as  $R$  increases as shown in Algorithm 1. As a result, most cores in the GPU will spend a large portion of time waiting for the processing of the largest enumeration tree, which aggravates the load imbalance. Related graph pattern mining algorithm  $G^2$ Miner [15] always assigns each enumeration tree to a warp in the GPU. However, Figure 4 shows that if we assign each enumeration tree to a warp based on our new algorithm GMBE, over 80% of SMs (i.e., 86 SMs / 108 SMs) waste 80% of running time (i.e., 1,458s / 1,822s) waiting for the slowest one on the BookCrossing dataset. It is necessary to balance workloads for MBE in a finer granularity.

## 4 DESIGN OF GMBE

We propose a new GPU-based MBE algorithm (GMBE) with three novel techniques. (1) We design a stack-based iterative MBE algorithm to replace the original recursive algorithm. The new MBE algorithm reuses the nodes of enumeration trees to reduce memory usage. (2) It uses local neighborhood sizes for pro-actively pruning for less thread divergence. And (3) it supports load-aware task scheduling to balance workloads among threads in warps and blocks. We describe them in detail in the following sections.

### 4.1 Stack-Based Iteration with Node Reuse

To reduce memory usage, we propose a stack-based iteration with node reuse. The key idea is that we can reuse a node  $x$  to represent its child nodes with additional metadata stored in the node  $x$ . By doing this, we save the memory space allocated for the child nodes of  $x$ . We can efficiently support node reuse because during enumeration the  $L \cup R \cup C$  of the child nodes of  $x$  is always a subset of the  $L \cup R \cup C$  of its parent node  $x$ . For instance, vertices  $\{u_2, v_1, v_2, v_3, v_4\}$  in child node  $q$  ( $\{u_2\}, \{v_1, v_2, v_3, v_4\}, \emptyset$ ) is the subset of vertices  $\{u_1, u_2, v_1, v_2, v_3, v_4\}$  in its parent node  $p$  ( $\{u_1, u_2\}, \{v_1, v_2, v_3\}, \{v_4\}$ ) as shown in Figure 2. We describe this stack-based MBE algorithm with node reuse in Algorithm 2.

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#### Algorithm 2: Stack-based Iterative MBE Algorithm

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**Data:** Bipartite graph  $G(U, V, E)$   
**Input:** Set  $L_r \subseteq U$ , disjoint sets  $R_r, C_r \subseteq V$   
**Output:** All maximal bicliques

```

1 procedure iteratively_search( $L_r, R_r, C_r$ ):
2    $node\_buf$ .init_and_push( $(L_r, R_r, C_r)$ );
3   while  $node\_buf$  is not empty do           // Iteration
4      $(L_p, R_p, C_p) \leftarrow node\_buf$ .pop();
5     if  $C_p$  is not empty then
6        $v' \leftarrow$  the smallest vertex in  $C_p$ ;
7        $node\_buf$ .push( $(L_p, R_p, C_p \setminus \{v'\})$ );
8        $L' \leftarrow L_p \cap N(v')$ ;  $R' \leftarrow R_p$ ;  $C' \leftarrow \emptyset$ ;
9       foreach  $v_c \in C_p$  do                 // Node generation
10        if  $L' \cap N(v_c) == L'$  then
11           $R' \leftarrow R' \cup \{v_c\}$ ;
12        else if  $L' \cap N(v_c) \neq \emptyset$  then
13           $C' \leftarrow C' \cup \{v_c\}$ ;
14        if  $R' == \Gamma(L')$  then               // Maximality check
15          Output  $(L', R')$  as a maximal biclique;
16         $node\_buf$ .push( $(L', R', C')$ );

```

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Specifically, instead of dynamically creating and freeing nodes for recursions in Algorithm 1, we **replace recursions** (line #11 in Algorithm 1) **with iterations** (lines #2-5, 7, 16 in Algorithm 2) and explicitly manage nodes with a stack-like structure  $node\_buf$ . The  $node\_buf$  structure and its update process can refer to Figure 5. A  $node\_buf$  consists of a root node  $(L_r, R_r, C_r)$ , the attribute  $depth$  of each vertex in  $L_r \cup R_r \cup C_r$ , and the *traversed vertices* from the root node to the current node. The  $depth$  of each vertex is updated according to the depth of the current node (i.e., the number of ancestor nodes of the current node). We can apply the node reuse

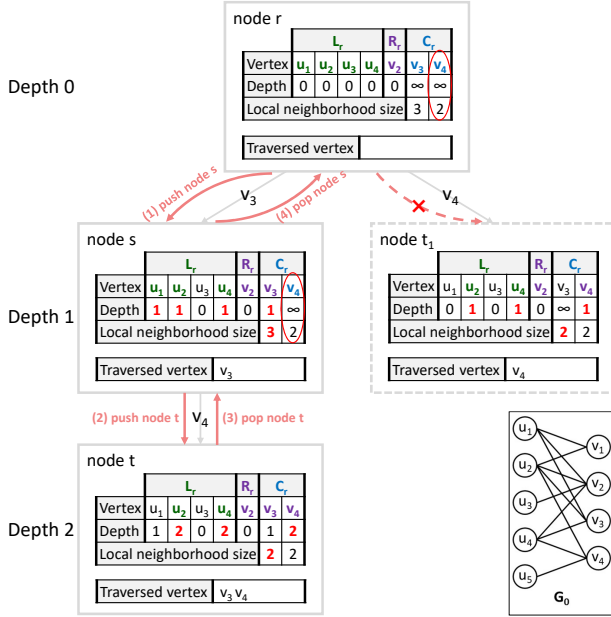
strategy on  $node\_buf$  by actively updating the  $depth$  field, and backtrack to ancestor nodes using the *traversed vertices* field. In this way, GMBE can derive all nodes in a fixed memory region during iteration, minimizing the need for dynamic allocation in GPUs. Next, we illustrate the key functions designed for node reuse.

- **init\_and\_push**( $(L_r, R_r, C_r)$ ): this function is used for creating and initializing a  $node\_buf$  using the root node  $(L_r, R_r, C_r)$  of a subtree (line #2 in Algorithm 2). The  $node\_buf$  stores all vertices in  $L_r \cup R_r \cup C_r$  and actively tracks the  $depth$  for each vertex. We initialize the  $depth$  for vertices in  $L_r \cup R_r$  to 0 and the  $depth$  for vertices in  $C_r$  to  $\infty$ .
- **push**( $(L', R', C')$ ): we use this function to reuse the  $node\_buf$  of a parent node to generate a child node (line #16 in Algorithm 2). When we push a new node  $(L', R', C')$  at depth  $D$ , we first update the  $depth$ s of vertices in  $L_r$  to  $D$  if they are also in  $L'$ , and then update the  $depth$ s of vertices in  $C_r$  to  $D$  if they are also in  $R'$  and their current  $depth$ s are  $\infty$ . Therefore, the new  $(L', R', C')$  can be found in the original  $(L_r, R_r, C_r)$  by the  $depth$  field, as  $L'$  contains all vertices in  $L_r$  whose  $depth$  is equal to  $D$ , and  $R'$  contains all vertices in  $R_r \cup C_r$  whose  $depth$  is not greater than  $D$ , and  $C'$  contains all vertices in  $C_r$  whose  $depth$  is  $\infty$ . Finally, we append the *traversed vertices* with the chosen vertex  $v'$ .
- **pop**(): we use this function to get the current node  $(L_p, R_p, C_p)$  and then backtrack to its parent node using  $node\_buf$  (line #4 in Algorithm 2). When we pop a node  $(L_p, R_p, C_p)$  at depth  $D$ , we first remove the latest traversed vertex  $v'$  in  $node\_buf$ . Then, we update the  $depth$  for vertices in  $L_p$  to  $D - 1$  and update the  $depth$  for vertices in  $C_p$  with  $depth$   $D$  to  $\infty$ .

**Discussion.** Compared to the original node structure which only tracks the  $(L, R, C)$ , our proposed  $node\_buf$  tracks more information with more memory consumption for a single node, which is bounded by  $3 \times \Delta(V) + 2 \times \Delta_2(V)$ . Whereas, a single  $node\_buf$  is adequate to be reused for all enumeration nodes for running a subtree traversal procedure shown in Algorithm 2, which significantly reduces the memory requirement and enables running thousands of MBE procedures on GPUs in parallel. For instance, an A100 GPU of 40 GB memory is adequate to run over 10k of subtree traversal procedures on the BookCrossing dataset [25] because each procedure requires only  $(3 \times 13,601 + 2 \times 53,915) \times \text{sizeof}(\text{int}) \text{ B} = 595 \text{ KB}$ . Compared to the naive implementation discussed in Section 3.1 which requires  $13,601 \times (13,601 + 53,915) \times \text{sizeof}(\text{int}) \text{ B} = 3.67 \text{ GB}$ , this node reuse approach saves  $6,178 \times$  memory space for running each MBE procedure on BookCrossing.

### 4.2 Pruning using Local Neighborhood Size

The existing pruning approaches are inefficient on GPUs because of thread divergence. To address this issue, we propose a new pruning approach to reduce enumeration space and thread divergence. Specifically, given a node  $(L, R, C)$ , we define **local neighbors** of a vertex  $v \in V$  as  $N_L(v)$ , where  $N_L(v)$  is equal to  $N(v) \cap L$ . We further define *local neighborhood size* for a vertex  $v$  as the number of local neighbors of  $v$ . We can obtain local neighborhood sizes for candidate vertices without additional overhead because they are intermediate results for computing the candidate set (line #12 in Algorithm 2).



**Figure 5: Illustration of GMBE with pruning using local neighborhood size.**

An interesting observation is that we can safely prune nodes generated by vertices whose local neighborhood size does not change after traversing any of its child nodes. We will formulate it in Theorem 4.1. Thus, we further optimize Algorithm 2 by maintaining *local neighborhood sizes* for all candidate vertices in `node_buf`. We can then prune useless candidates if their local neighborhood sizes do not change after popping a traversed child node. The new pruning approach has low thread divergence because threads in a warp always check elements in the same candidate set. We use an example to detail the stack-based iteration algorithm with pruning using local neighborhood size.

**EXAMPLE 4.1.** Figure 5 illustrates that GMBE iteratively enumerates all maximal bicliques in the subtree rooted by node  $r$  in Figure 2 with the fixed memory in `node_buf`. Specifically, GMBE initializes `node_buf` using node  $r$ , where  $L_r = L(r)$ ,  $R_r = R(r)$ , and  $C_r = C(r)$ . `node_buf` then initializes the depth for vertices in  $L_r \cup R_r$  to 0 and initializes the depth for vertices in  $C_r$  to  $\infty$ . `node_buf` initializes the local neighborhood size (i.e.,  $|N_L|$ ) for vertices in  $C_r$  according to the definition. For instance,  $|N_L(v_3)|$  at node  $r$  is  $|N(v_3) \cap L(r)| = |\{u_1, u_2, u_4\} \cap \{u_1, u_2, u_3, u_4\}| = 3$ .

Next, we generate node  $s$  by traversing  $v_3$ . We know  $L(s) = L(r) \cap N(v_3) = \{u_1, u_2, u_4\}$ . The depth of node  $s$  is 1, `node_buf` updates the depth for  $u_1, u_2, u_4$ , and  $v_3$  to 1 because they belong to  $L(s) \cup R(s)$ . The depth for other vertices (i.e.,  $u_3$  and  $v_4$ ) in  $L_r$  and  $R_r$  remains 0. The depth for  $v_4$  in  $C_s$  remains as  $\infty$ .

We then update the local neighborhood size for  $v_3$  and  $v_4$  using  $L(s)$ . We know  $|N_L(v_3)| = |N(v_3) \cap L(s)| = |\{u_1, u_2, u_4\} \cap \{u_1, u_2, u_4\}| = 3$ .  $|N_L(v_4)| = |N(v_4) \cap L(s)| = |\{u_2, u_4, u_5\} \cap \{u_1, u_2, u_4\}| = 2$ . We can generate other nodes similarly.

After processing the subtree rooted by node  $s$ , `node_buf` pops node  $s$  and recovers the parent node  $r$ . `node_buf` resets depth for  $u_1, u_2$ , and

### Algorithm 3: Naive approach for launching GPU tasks

```

foreach  $v_s \in V$  do // Initialize  $|V|$  tasks independently
1   $L_s \leftarrow N(v_s); R_s \leftarrow \{v_s\}; C_s \leftarrow \emptyset$ ; // Node generation
2  foreach  $v_c \in N_2(v_s)$  do
3    if  $L_s \cap N(v_c) == L_s$  then
4       $R_s \leftarrow R_s \cup \{v_c\}$ ;
5    else if  $v_c$  is with later order than  $v_s$  then
6       $C_s \leftarrow C_s \cup \{v_c\}$ ;
7  if  $v_s$  is the smallest vertex in  $R_s$  then // Maximality check
8    // Map each task to a warp or a block
   iteratively_search( $L_s, R_s, C_s$ );

```

$u_4$  to 0 and resets the depth for  $v_3$  to  $\infty$  because their original depth is the same as the depth of node  $s$ . GMBE proactively prunes node  $t_1$  by removing useless candidate vertex  $v_4$  at node  $r$  because the local neighborhood size (i.e., 2) for  $v_4$  does not change after popping node  $s$ .

**THEOREM 4.1.** Assume the current node is  $p$  in the enumeration tree. GMBE can safely prune its child nodes generated by choosing a candidate vertex  $v_c$  in  $C_{(p)}$  if  $|N_L(v_c)|$  for the current node  $p$  is equal to  $|N_L(v_c)|$  for any of the child nodes of  $p$ .

**PROOF.** To prove the theorem, we assume that node  $s$  has traversed  $v_m$  to generate node  $t$  and node  $s$  will traverse  $v_n$  to generate node  $r$ . We further assume that  $|N_L(v_n)|$  for node  $s$  is equal to  $|N_L(v_n)|$  for node  $t$ . Because  $|N_L(v_n)|$  for node  $s$  is equal to  $|N_L(v_n)|$  for node  $t$ ,  $|N(v_n) \cap L(s)| = |N(v_n) \cap L(t)|$ . Because  $L(t) = L(s) \cap N(v_m)$ ,  $|N(v_n) \cap L(s)| = |N(v_n) \cap L(t)| = |N(v_n) \cap L(s) \cap N(v_m)|$ . Then, we know  $N(v_n) \cap L(s) \subseteq N(v_m)$ . Because  $L(r) = N(v_n) \cap L(s)$ ,  $L(r) \subseteq N(v_m)$ . Thus, we know that  $v_m$  connects with all vertices in  $L(r)$ . Node  $r$  fails to expand  $R(r)$  with  $v_m$  because GMBE has traversed  $v_m$  generate node  $t$  and we can safely prune node  $r$ , which produces a non-maximal biclique.  $\square$

### 4.3 Load-Aware Task Scheduling

For exploring the massive parallelism of GPUs, a naive approach is assigning a task to manage each enumeration tree whose root node is  $v_s$  ( $v_s \in V$ ). The naive approach is shown in Algorithm 3. We can then map these tasks to warps [15] or blocks [11] in GPUs. We denote these schemes as the *warp-centric scheme* and the *block-centric scheme*, respectively. Our research shows that the naive approach is insufficient to balance the loads among GPU SMs for the MBE problem because the parallel GPU tasks created at line #8 in Algorithm 3 are highly unbalanced and their loads are determined by the various sizes of the enumeration trees assigned to the tasks. We observe that the slowest tasks running in a warp or a block frequently block other tasks, resulting in up to 97.8% performance degradation on the EuAll dataset. More results can be found in Figure 9 in Section 6.2.

To address this issue, we propose a **load-aware task-centric** approach for MBE using the persistent thread (PT) [21] programming model for GPUs. Specifically, we create thread groups, each of which consists of multiple warps. Each thread group is mapped to a GPU SM. We denote the number of warps on each SM as  $\text{WarpPerSM}$  and will discuss its impact on system performance in

**Algorithm 4:** Load-aware task-centric scheme in GMBE

---

```

processing_v is a global variable initialized as 0;
SM_task_queue is a global concurrent queue for load balance;
// For each warp
1 procedure warp_kernel:
2   while true do
3     if SM_task_queue is not empty then
4       (L, R, C) ← SM_task_queue.dequeue();
5     else
6       v_s = atomicInc(processing_v);
7       if v_s ∈ V then
8         L ← N(v_s); R ← {v_s}; C ← ∅;
9         foreach v_c ∈ N_2(v_s) do
10          if L ∩ N(v_c) == L then
11            R ← R ∪ {v_c};
12          else if v_c is with later order than v_s then
13            C ← C ∪ {v_c};
14        else // All tasks on GPU have been processed
15          return;
16      if R == Γ(L) then // Maximality check
17        if min{|L|, |C|} × |C| > bound_size and
18          min{|L|, |C|} > bound_height then
19          foreach v_t ∈ C do // Node generation
20            L_t ← N(v_t); R_t ← R; C_t ← ∅;
21            foreach v_c ∈ C do
22              if L_t ∩ N(v_c) == L_t then
23                R_t ← R_t ∪ {v_c};
24              else if L_t ∩ N(v_c) ≠ ∅ then
25                C_t ← C_t ∪ {v_c};
26            SM_task_queue.enqueue((L_t, R_t, C_t));
27            C ← C \ {v_t};
28        else
29          iteratively_search(L, R, C);

```

---

Section 6.4. We develop a GPU kernel, which can create load-aware tasks. Any tasks of processing larger enumeration trees are divided into smaller tasks recursively at runtime. The load-aware tasks are then added to a global structure *SM\_task\_queue* for each SM. When a task is finished, the software scheduler of PT dequeues a task from *SM\_task\_queue* and executes *iteratively\_search()* on its corresponding SM. We denote it as the *task-centric scheme*.

The key question to answer is how to detect tasks having heavier loads than others. We use the tree heights and the total number of nodes in a tree. Specifically, the height of an enumeration tree with the root node  $(L, R, C)$  is  $\min\{|L|, |C|\}$ . The number of nodes of the tree can be estimated as  $\min\{|L|, |C|\} \times |C|$ , where  $|C|$  indicates the maximum number of child nodes for each node in the enumeration tree. We empirically set two thresholds including *bound\_height* and *bound\_size*. Only when  $\min\{|L|, |C|\}$  is larger than *bound\_height* and  $\min\{|L|, |C|\} \times |C|$  is larger than *bound\_size*, we divide the task into multiple subtasks to achieve better load balance.

**Putting them together.** Algorithm 4 describes GMBE, which is a load-aware task-centric algorithm for GPUs. When *SM\_task\_queue* is not empty, it obtains a node from the queue (lines #3-4). If the size of the enumeration tree generated from the node is within bounds (line #17), GMBE directly launches a GPU task (line #28). If the size of the enumeration tree is larger than the bounds, it enqueues the root node of sub-trees to be enumerated (lines #18-26). These nodes will be processed later when they are dequeued. If *SM\_task\_queue* is empty, it will use *processing\_v* to obtain the current  $v_s$  that needs to be processed (line #6). Then, it generates a node  $(L, R, C)$  (lines #7-13) and launches its corresponding tree onto GPUs.

## 5 IMPLEMENTATION ISSUES

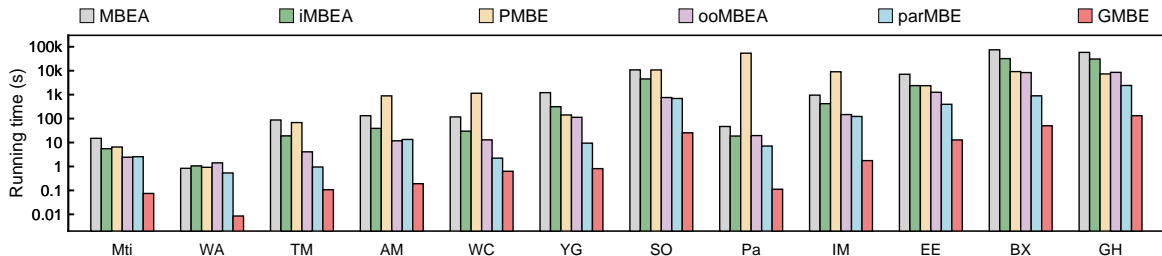
**Pre-processing.** We represent the input bipartite graph  $G$  in the compressed sparse row (CSR) format. We first load graph  $G$  into CPU memory and quickly extract important characteristics from  $G$ , such as  $|U|$ ,  $|V|$ ,  $|E|$ ,  $\Delta(V)$ , and  $\Delta_2(V)$ . Since  $U$  and  $V$  are symmetrical in the bipartite graph, we always select the vertex set with fewer vertices as  $V$  similar to [14]. We then pre-process  $G$  by sorting all vertices in  $V$  using the increasing order of their neighborhood sizes [29, 39] and sorting all neighbor lists of each vertex using increasing order of vertex IDs similar to most of the related works [11, 15]. We finally transfer the whole bipartite graph  $G$  to the global memory of the GPU and enumerate all maximal bicliques on GPUs without transferring any extra data from hosts.

**Lock-free task queue.** To reduce the synchronization overhead, we manage the task queue in a lock-free manner using the atomicCAS primitive [28] in CUDA. We implement a two-level task queuing mechanism to further improve load balance. Specifically, we implement a local task queue for each block so that all warps in the block can balance workloads by accessing the local task queue. In addition, we implement a global task queue to balance workloads between different blocks. Each block only allows one proxy warp to manage tasks between the local task queue and the global task queue. We implement the local task queues using the shared memory and implement the global task queue in the global memory because atomic operations on shared memory are faster than atomic operations on the global memory.

**MBE on multiple GPUs.** A high-performance machine may consist of multiple GPUs to accelerate application execution performance. To support this scenario, we can easily extend GMBE algorithm to multi-GPU machines. The main idea is sharing the global variable *processing\_v* in Algorithm 4 on all GPU devices and replacing the atomicInc primitive in line #6 with atomicInc\_system [1]. Consequently, the MBE problem is divided into multiple independent sub-problems, and each GPU independently processes these sub-problems. The overall running time is determined by the GPU with the longest execution time. Experimental results in Figure 13 shows that GMBE is efficient on multiple GPUs because each warp on multiple GPUs can automatically balance workloads using atomic primitives with little synchronization overhead. Theoretically, GMBE can also be extended to a distributed computing environment, where multiple machines (each with one or more GPUs) are connected by the network. Since this work focuses on the single-machine environment, we leave the exploration of GMBE on distributed multi-machine clusters as our future work.

**Table 1: Dataset statistics**

Datasets	$ U $	$ V $	$ E $	$\Delta(U)$	$\Delta_2(U)$	$\Delta(V)$	$\Delta_2(V)$	Max. bicliques
MovieLens (Mti)	16,528	7,601	71,154	640	5,817	146	3,217	140,266
Amazon (WA)	265,934	264,148	925,873	168	635	546	903	461,274
Teams (TM)	901,130	34,461	1,366,466	17	18,516	2,671	2,838	517,943
ActorMovies (AM)	383,640	127,823	1,470,404	646	3,956	294	7,798	1,075,444
Wikipedia (WC)	1,853,493	182,947	3,795,796	54	47,190	11,593	4,629	1,677,522
YouTube (YG)	94,238	30,087	293,360	1,035	37,513	7,591	7,356	1,826,587
StackOverflow (SO)	545,195	96,680	1,301,942	4,917	146,089	6,119	31,636	3,320,824
DBLP (Pa)	5,624,219	1,953,085	12,282,059	287	7,519	1,386	2,119	4,899,032
IMDB (IM)	896,302	303,617	3,782,463	1,590	15,451	1,334	15,233	5,160,061
EuAll (EE)	225,409	74,661	420,046	930	135,045	7,631	23,844	12,306,755
BookCrossing (BX)	340,523	105,278	1,149,739	2,502	151,645	13,601	53,915	54,458,953
Github (GH)	120,867	59,519	440,237	3,675	29,649	884	15,994	55,346,398

**Figure 6: Overall evaluation (log scaled).**

## 6 EVALUATION

In this section, we conduct extensive experiments to evaluate the performance of GMBE and the proposed techniques.

### 6.1 Experimental Setup

**Platform.** By default, we evaluate our GPU implementations on an NVIDIA A100 GPU [4] with 108 streaming multiprocessors (SMs) and 40 GB of global memory. For the comparison, we run the other CPU-based MBE algorithms on a Linux server with 96 Xeon(R) Gold 5318Y CPU @ 2.10GHz CPU cores. The operating system is Linux kernel-5.4.0.

**Datasets.** We use 12 real-world datasets to justify the performance of GMBE as shown in Table 1. It’s worth noting that for datasets that allow multiple edges between two vertices, such as MovieLens (Mti), we only retain one unique edge between each vertex pair for MBE analysis. The number of these unique edges is denoted by  $|E|$ . Since  $U$  and  $V$  are symmetrical in the bipartite graph, we always denote the vertex set with fewer vertices as  $V$ , i.e.,  $|U| > |V|$ . We obtain datasets Amazon and EuAll from the SNAP repository [26] and the other datasets from the KONECT repository [25]. Since the MBE time mainly depends on the number of maximal bicliques of the dataset, we sort all datasets in ascending order of their maximal biclique count. Datasets with more than two million maximal bicliques are referred to as large datasets in subsequent sections.

**Compared Algorithms.** Since there is no existing MBE algorithm working on GPUs, we compare GMBE to the CPU-oriented MBE algorithms, including the recent serial versions, i.e., MBEA [39], iMBEA [39], PMBE [8], and ooMBEA [14], and the cutting-edge parallel MBE algorithm, i.e., PARMBE [18]. For fair comparisons, we obtain well-optimized codes of all competitors from the authors and

run them on the same platform. We run PARMBE with 96 threads because our machine contains 96 CPU cores.

**Measures.** We measure the running time of each algorithm excluding the time spent reading the graph from the disk. Without specification, GMBE iteratively enumerates all maximal bicliques with node reuse, prunes useless nodes using local neighborhood sizes, and applies the load-aware task-centric scheme for load balance. By default, GMBE sets the thresholds for *bound\_height* and *bound\_size* to 20 and 1,500 respectively, sets *WarpPerSM* to 16, and sorts  $V$  in ascending order based on the vertex degree before the enumeration. We also implement other variants to evaluate the techniques proposed in this paper. We will detail those variants in the corresponding experiments.

### 6.2 Overall Evaluation

Figure 6 compares the running time of GMBE to state-of-the-art MBE algorithms on real-world datasets. The experimental results show that GMBE is 3.5×–69.8× faster than any next-best competitor on CPUs on all testing datasets because GMBE efficiently utilizes extensive computational resources on GPUs. Specifically, GMBE on a single A100 GPU outperforms the state-of-the-art parallel MBE algorithm PARMBE on a 96-core CPU machine by up to 70.6× on ActorMovies. Compared to all the existing MBE algorithms that cost over 40 minutes to enumerate all maximal bicliques on Github, GMBE needs only 132 seconds and thus is helpful for MBE on the large datasets in practice. In addition, we conduct a performance analysis of GMBE using the NVIDIA Nsight Compute software [5]. The profiling results indicate that the average warp execution efficiency is 64%, and the memory utilization is 12% across all real-world datasets. These outcomes can be attributed to the inherent irregularity present in the MBE problem [13].



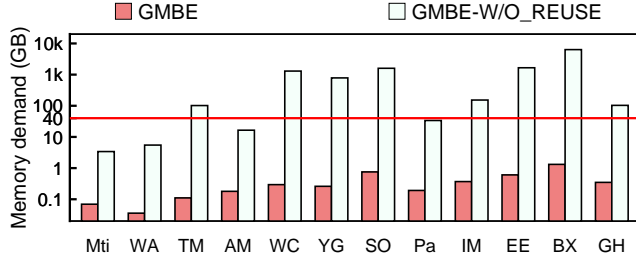


Figure 7: Effect of the node reuse approach (log scaled). The red line indicates the GPU memory capacity of NVIDIA A100.

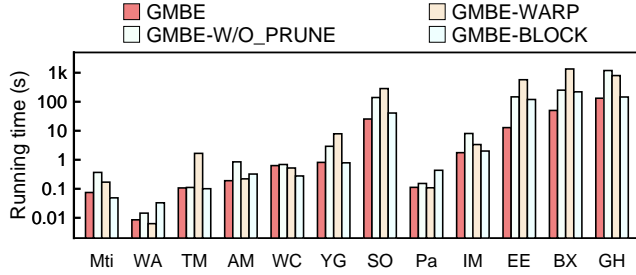


Figure 8: Effect of pruning approach and task scheduling approach (log scaled).

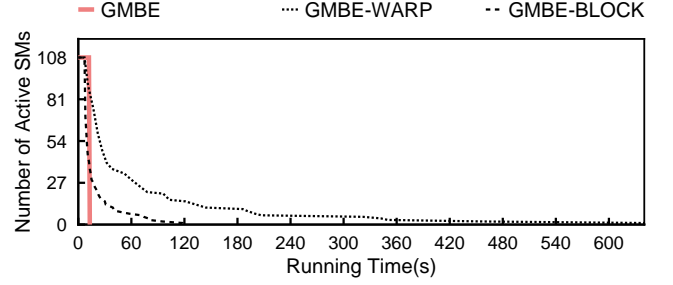
Table 2: Comparison of the ratio of generated non-maximal bicliques to maximal bicliques between GMBE and GMBE-w/o\_PRUNE.

Datasets	Mti	WA	TM	AM	WC	YG
GMBE	9.04	0.734	1.63	12.9	0.71	2.11
GMBE-w/o_PRUNE	66.0	3.68	3.88	53.0	2.89	20.1
Datasets	SO	Pa	IM	EE	BX	GH
GMBE	89.4	0.362	15.5	4.04	3.40	11.1
GMBE-w/o_PRUNE	174	1.43	74.4	56.0	27.3	51.4

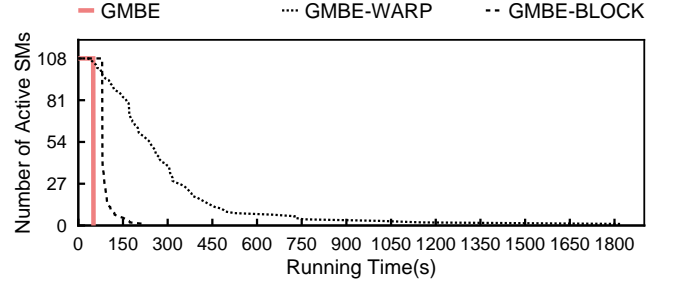
### 6.3 Effect of Optimizations

**Effect of the node reuse approach.** To study the effect of the node reuse approach in Section 3.1, we design a variant GMBE-w/o\_REUSE that pre-allocates memory on GPUs according to Section 3.1. We estimate the memory requirement allocated by the cudaMalloc primitive for GMBE with and without the node reuse approach. This memory requirement includes the pre-allocated memory for the input bipartite graph and the runtime subtrees. Figure 7 shows that the node reuse approach significantly reduces the memory requirement by  $49\times$ – $4,819\times$  on all testing datasets while GMBE-w/o\_REUSE is impractical because its memory requirements exceed the memory capacity of the A100 GPU on multiple datasets.

**Effect of the pruning approach.** To study the effect of the local-neighborhood-size-based pruning approach in Section 4.2, we design a variant GMBE-w/o\_PRUNE that only disables the pruning function of GMBE. As shown in Figure 8, GMBE constantly outperforms GMBE-w/o\_PRUNE because the pruning approach prunes enumeration space for MBE at runtime with less thread divergence.



(a) EuAll

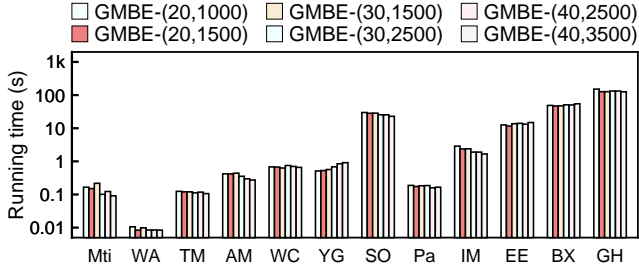


(b) BookCrossing

Figure 9: Comparison of runtime loads on SMs among GMBE, GMBE-WARP, and GMBE-BLOCK.

In addition, the pruning approach enhances memory access coalescence by comparing the sizes of local neighborhoods among vertices in the same candidate vertex set. To further explore the pruning efficiency, we use  $\alpha$  to represent the number of maximal bicliques and  $\delta$  to represent the number of pruned non-maximal bicliques generated by node checking (line #14 in Algorithm 2). Since  $\alpha$  remains constant for each dataset, we use the ratio  $\delta/\alpha$  to indicate the pruning efficiency for both GMBE and GMBE-w/o\_PRUNE, as presented in Table 2. By comparing the ratio  $\delta/\alpha$  of both GMBE and GMBE-w/o\_PRUNE, we observe that the proposed pruning approach can avoid 48.7%–92.8% non-maximal biclique checks among all testing datasets. This pruning technique plays a crucial role, particularly for larger datasets where the enumeration space grows with an increasing number of maximal bicliques. Consequently, the pruning approach significantly reduces the running time on Github from 1,191 seconds to 132 seconds.

**Effect of the task scheduling approach.** To study the effect of the load-aware task scheduling approach in Section 4.3, we design two variants GMBE-WARP and GMBE-BLOCK that apply warp-centric and block-centric schemes respectively. As shown in Figure 8, GMBE is significantly faster than GMBE-WARP and GMBE-BLOCK on large datasets, including EuALL, Github, BookCrossing, StackOverflow, and IMDB, and spends less than one second on the other datasets. GMBE is more performant on large datasets because it dynamically detects and partitions the tasks having heavy loads and manages lock-free task queues to rebalance the workloads in finer granularity. As a result, GMBE is  $44.7\times$  and  $9.3\times$  faster than GMBE-WARP and GMBE-BLOCK on EuAll, respectively.



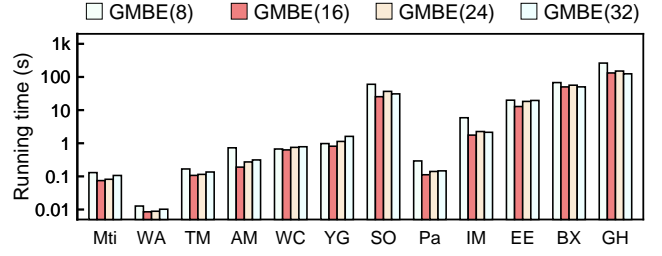
**Figure 10: Impact of thresholds *bound\_height* and *bound\_size* for load-aware task scheduling (log scaled).**

To further explore the load imbalance problem for MBE on GPUs, we record the number of active SMs while running GMBE, GMBE-WARP, and GMBE-BLOCK. Figure 9 reports the comparison of runtime loads on SMs on the BookCrossing and EuAll datasets. Because of the imbalanced workloads, SMs with light workloads may finish early and wait for the SM with the heaviest workload, which is costly. GMBE-WARP gains the worst performance because the number of active SMs decreases rapidly due to the load imbalance. GMBE-BLOCK obtains better performance than GMBE-WARP because GMBE-BLOCK could spend more resources on each workload than GMBE-WARP (i.e., a block vs. a warp) which reduces the time waiting for the SM with the heaviest workload. However, GMBE-BLOCK is insufficient because the workloads for the MBE problem could be severely imbalanced. For instance, over 80% of SMs (86 SMs / 108 SMs) waste over 80% of running time (98s / 118s) waiting for the slowest SM on EuALL. GMBE always achieves the best performance because GMBE works in the finest granularity and each SM finishes its work roughly at the same time. GMBE completes even before the number of active SMs of GMBE-BLOCK starts to decrease on BookCrossing because GMBE activates all warps in each SM while GMBE-BLOCK may use only use a small portion of warps in each SM at runtime.

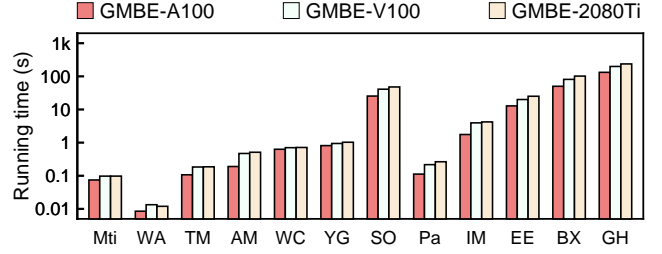
### 6.4 Sensitivity Analysis

**Impact of thresholds for load-aware task scheduling.** To explore the efficient configuration for thresholds *bound\_height* and *bound\_size* in Section 4.3, we design multiple variants GMBE-(*m*, *n*), where *m* and *n* represent *bound\_height* and *bound\_size* respectively. We always set *m* larger than *n*<sup>2</sup> because  $|L| \times |C|$  is always greater or equal to  $(\min\{|L|, |C|\})^2$ . The selection of thresholds is a trade-off between the parallel granularity and synchronization overhead. We require smaller thresholds to balance workloads in finer granularity. However, the thresholds should not be too small. Otherwise, we have to manage more tasks with the huge synchronization overhead. Figure 10 shows that the variant GMBE-(20, 1500) is empirically better than the others in most cases. Thus, GMBE applies this configuration by default.

**Impact of the number of warps in each SM.** To determine the parameter WarpPerSM in the PT model in Section 4.3, we design variants that set WarpPerSM to 8, 16, 24, and 32, respectively. The selection of WarpPerSM is a trade-off between parallelism and the resources for each warp. Intuitively, we expect WarpPerSM



**Figure 11: Impact of the parameter WarpPerSM (log scaled).**



**Figure 12: Adaptability on different GPU (log scaled).**

to be larger so that we can run more MBE tasks in parallel. However, WarpPerSM should not be too large since the computational resources (e.g., registers) in each SM are limited. A larger WarpPerSM may decrease the performance of GMBE because each warp will have fewer resources to run MBE tasks. Figure 11 shows that the variant GMBE(16) outperforms the other variants by up to 3.83× on most large datasets, such as BookCrossing, StackOverflow, IMDB, DBLP, and EuAll. In addition, GMBE(16) is 0.94× slower than GMBE(32) on Github due to its extensive enumeration space that requires more warps to enumerate maximal bicliques in parallel. Considering its efficiency in most cases, GMBE sets WarpPerSM to 16 by default.

**Adaptability on different GPUs.** To explore the adaptability of GMBE, we evaluate GMBE on an NVIDIA A100 GPU (108 SMs, 40 GB global memory), an NVIDIA V100 GPU (80 SMs, 32 GB global memory) [6], and an NVIDIA 2080Ti GPU (68 SMs, 11 GB global memory) [3], respectively. Figure 12 shows that GMBE is adaptive on all three GPUs. GMBE-A100 is slightly faster than both GMBE-V100 and GMBE-2080Ti because an A100 GPU contains more computational resources than other GPUs.

**Scalability on multi-GPU.** To explore the scalability of GMBE on multi-GPU, we conduct experiments on a machine with 8 NVIDIA V100 GPUs. To optimize GMBE for multi-GPU configurations, we divide the problem into multiple independent sub-problems, and the total execution time is determined by the longest-running sub-problem. Figure 13 shows that GMBE scales out linearly on Github and BookCrossing datasets as we increase the number of GPUs because each GPU finishes its execution almost at the same time. With the help of multiple GPUs, GMBE can enumerate over 55 million maximal bicliques on Github dataset within 31 seconds, meaning a performance speedup of 77× compared to the state-of-the-art parallel MBE algorithm PARMBE on a 96-core CPU machine (i.e., 2,411 seconds).

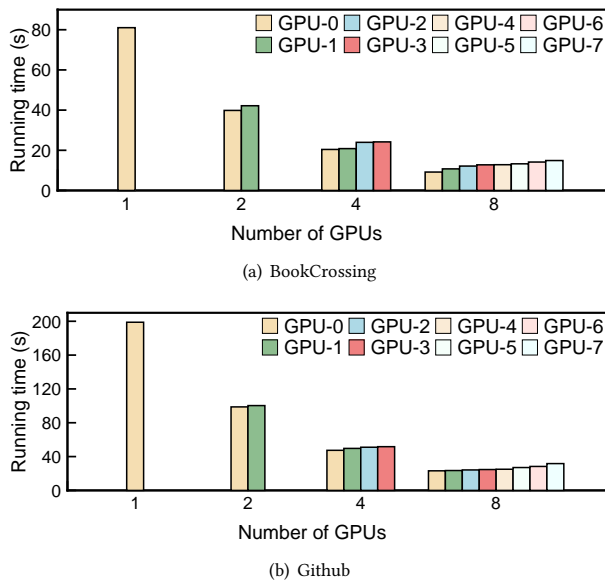


Figure 13: Scalability of GMBE on a machine with multi-GPU.

## 7 RELATED WORK

**Maximal Biclique Enumeration on CPUs.** The most efficient class of MBE algorithms [8, 14, 18, 29, 31, 32, 39] is based on backtracking with recursions on CPUs. Most of the research efforts [8, 14, 29, 39] applied various sorting and pruning techniques on a serial MBE algorithm to reduce the search space for the MBE problem. Some other works [18, 32] improved the efficiency of MBE by parallelizing MBE algorithms on multicore CPUs or distributed architectures. However, the existing MBE algorithms do not work on GPUs due to the challenges in Section 3, and thus achieve limited performance with limited computational resources on CPUs.

**Related Problems on GPUs.** Although GPUs are widely used to accelerate related graph algorithms, such as maximal clique enumeration (MCE) [12, 27, 36] and graph pattern mining (GPM) [11, 15, 20], it is still challenging for MBE on GPUs. Specifically, MCE on GPUs suffers from similar performance issues as MBE described in Section 3. As a result, the latest GPU-based MCE algorithm GBK [36] only obtained a suboptimal performance, which is comparable to a single-thread sequence algorithm on the CPU. Because the enumerated subgraphs (i.e., maximal bicliques) for MBE generally contain more vertices than those for other GPM problems, optimizations in the latest GPU-based GPM framework  $G^2$ Miner [15] cannot address the memory issue and severe load imbalance for MBE on GPUs.

## 8 CONCLUSION

In this paper, we present GMBE, the first highly-efficient GPU solution for the MBE problem. MBE on GPUs faces serious challenges, including large memory requirement, thread divergence, and severe load imbalance. To address these problems, we design a node-reuse approach to reduce GPU memory usage, a pro-active pruning method using local neighborhood size, and a load-aware

task scheduling framework to achieve load balance among threads within warps and blocks. We conduct comprehensive evaluations using 12 real-world datasets and three different GPUs. Our experimental results show that GMBE outperforms the state-of-the-art parallel MBE algorithm PARMBE by 70.6%.

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# Appendix: Artifact Description/Artifact Evaluation

## ARTIFACT DOI

<https://doi.org/10.5281/zenodo.8079259>

## ARTIFACT IDENTIFICATION

In this paper, we propose GMBE, the first highly-efficient GPU solution for the maximal biclique enumeration (MBE) problem. The computational artifact provides the source code and scripts to reproduce all experimental results. GMBE is implemented in C++ and CUDA. The artifact includes all comparison baselines (i.e., MBEA, iMBEA, PMBE, ooMBEA, and ParMBE) for justifying the performance of GMBE and all variants of GMBE for evaluating all proposed techniques. This artifact enables others to conduct their own studies on any Linux machine with NVIDIA GPUs.

## REPRODUCIBILITY OF EXPERIMENTS

### 1 ARTIFACT CHECKLIST

- **Platforms:**
  - An NVIDIA A100 GPU
  - An NVIDIA 2080Ti GPU
  - 8 NVIDIA V100 GPUs
  - A machine with 96 Xeon(R) Gold 5318Y CPU @ 2.10GHz CPU cores
- **System Details:**
  - 18.04-Ubuntu x86\_64 GNU/Linux
  - Linux kernel: 5.4.0
- **Software Dependencies:**
  - GNU Make 4.2.1
  - CMake 3.22.0
  - CUDA toolkit 11.7
  - GCC/G++ 10.3.0
  - Python 2.7.18
  - Python packages: zplot 1.41, pathlib 1.0.1
  - C++ library: libtbb-dev 2020.1-2
  - Ubuntu apt package: Ghostscript, Texlive-extra-utils
  - Nvidia driver 510.85.02
  - Docker 20.10.10

### 2 DESCRIPTION

The computational artifact provides scripts to reproduce all experimental results of GMBE, including Figures 6-13 and Table 2. Users need to download the source code and scripts from <https://github.com/fhXu00/MBE-GPU.git>. The following is the directory structure of the repository:

- **README.md:** This file contains a detailed step-by-step "Try out GMBE" guide.
- **src/:** This directory contains the core source code of GMBE implementation.
- **scripts/:** This directory has scripts to run experiments.
- **baselines/:** This directory contains the zipped source codes of the comparison baselines, i.e., MBEA, iMBEA, PMBE, ParMBE, and ooMBEA.

- **preprocess/:** This directory has scripts to preprocess the graph datasets.
- **fig/:** This directory contains the scripts to generate figures.

After downloading the source code and scripts, users need to prepare graph datasets and compile the code using our provided scripts. We provide a docker image in the repository <https://hub.docker.com/r/fhXu00/gmbe> for the convenience to deploy this project. Please refer to the corresponding *README.md* for a detailed guide.

### 3 EXPERIMENTAL WORKFLOW

#### Step 1: Prepare datasets.

We use 12 real-world datasets for our evaluation. Users can download and preprocess them with the script in the directory *preprocess/*, which will take roughly 10 minutes. After executing the script, users can find the preprocessed datasets in the new directory *datasets/*.

#### Step 2: GPU Setup.

We provide the specifications for setting up the project on A100, V100, and 2080Ti GPUs in the *CMakeList.txt* file located in the project directory. Users can deploy GMBE on other GPUs by setting the "CUDA\_NVCC\_FLAGS" and "DMAX\_SM" flags for the designated GPU, following the guidelines provided in the GPU manufacturer's documentation. The "DMAX\_SM" flag indicates the maximum number of SMs on the GPU.

#### Step 3: Execute scripts to generate results.

We provide the scripts to automatically generate the experimental results of Figures 6-13 and Table 2 in *scripts/*. It will take roughly 9 hours to generate all the results. The running progress and the expected running time of each experiment would be printed to the file *scripts/progress.txt* automatically. All the experimental results would be printed to the file *scripts/results.txt* and the results required to generate the figure would be printed to the data file in the corresponding subdirectory in *fig/*.

#### Step 4: Execute scripts to generate figures.

We provide the script to generate figures in the directory *fig/*, executing which will take roughly 30 seconds. The generated figures would be found under the directory *fig/*.

### 4 EVALUATION AND EXPECTED RESULTS

Users can reproduce all experimental results with the scripts, which should roughly match the figures from the paper. Note that, the results may be not the exact ones presented in the paper. Because in our subsequent experiments, we found that different states of the GPUs may also impact the performance.

### ARTIFACT DEPENDENCIES REQUIREMENTS

To conduct the experiments for GMBE, a Linux machine with Nvidia GPUs is required. The experiments of ParMBE require a machine with multi-core CPU and the C++ library libtbb-dev. The input datasets used can be obtained from two open-source repositories:

SNAP and KONECT. These datasets were collected from the real world and are widely used in MBE research.

## **ARTIFACT INSTALLATION DEPLOYMENT PROCESS**

The process about how to install, compile and deploy the code is detailed in the 'README.md' of our source code. It takes no more than 10 minutes to complete the process.

## **OTHER NOTES**

The project has also been deployed on ChameleonCloud, which can be found at <https://chi.uc.chameleoncloud.org/project/instances/b28ae6f2-aec0-4583-8126-789efcaefb2f/>.