

Coarse Grained Parallel Algorithms for Detecting Convex Bipartite Graphs

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Abstract. In this paper, we present parallel algorithms for the *coarse grained multicomputer* (CGM) and the *bulk synchronous parallel computer* (BSP) for solving two well known graph problems: (1) determining whether a graph G is *bipartite*, and (2) determining whether a bipartite graph G is *convex*.

Our algorithms require $O(\log p)$ and $O(\log^2 p)$ communication rounds, respectively, and linear sequential work per round on a CGM with p processors and N/p local memory per processor, $N=|G|$. The algorithms assume that $\frac{N}{p} \geq p^\epsilon$ for some fixed $\epsilon > 0$, which is true for all commercially available multiprocessors. Our results imply BSP algorithms with $O(\log p)$ and $O(\log^2 p)$ supersteps, respectively, $O(g \log(p) \frac{N}{p})$ communication time, and $O(\log(p) \frac{N}{p})$ local computation time.

Our algorithm for determining whether a bipartite graph is convex includes a novel, coarse grained parallel, version of the *PQ tree* data structure introduced by Booth and Lueker. Hence, our algorithm also solves, with the same time complexity as indicated above, the problem of testing the consecutive-ones property for $(0, 1)$ matrices as well as the chordal graph recognition problem. These, in turn, have numerous applications in graph theory, DNA sequence assembly, database theory, and other areas.

1 Introduction

In this paper, we study the problem of detecting bipartite graphs and convex bipartite graphs. That is, given an arbitrary graph G , determine whether G is a *bipartite* graph and, given a bipartite graph G , determine whether G is a *convex* bipartite graph. Bipartite and convex bipartite graphs are formally defined as follows.

Definition 1. A graph $G = (V, E)$ is a *bipartite graph* if V can be partitioned into two sets A and B such that $A \cap B = \emptyset$, $A \cup B = V$ and $E \subseteq ((A \times B) \cup (B \times A))$. A *bipartite graph* G is also denoted as $G = (A, B, E)$.

Definition 2. A bipartite graph $G = (A, B, E)$ is a convex bipartite graph if there exists an ordering $(b_1, b_2, \dots, b_{|B|})$ of B such that, for all $a \in A$ and $1 \leq i < j \leq |B|$, if $(a, b_i) \in E$ and $(a, b_j) \in E$ then $(a, b_k) \in E$ for all $i \leq k \leq j$.

These, and closely related, problems has been extensively studied for the sequential [1,15] and the shared memory (PRAM) parallel [4,5,11,12,13,14] domain. Unfortunately, theoretical results from PRAM algorithms do not necessarily match the speedups observed on *real* parallel machines. In this paper, we present parallel algorithms that are more practical in that the assumptions and cost model used reflects better the reality of commercially available multiprocessors. More precisely, we will use a version of the BSP model, referred to as the *coarse grained multicomputer* (CGM) model. In contrast to the BSP model, the CGM [6,7,8,9] allows only bulk messages in order to minimize message overhead costs. A CGM is comprised of a set of p processors P_1, \dots, P_p with $O(N/p)$ local memory per processor and an arbitrary communication network (or shared memory). All algorithms consist of alternating local computation and global communication rounds. Each communication round consists of routing a single h -relation with $h = O(N/p)$, i.e. each processor sends $O(N/p)$ data and receives $O(N/p)$ data. We require that all information sent from a given processor to another processor in one communication round is packed into one long message, thereby minimizing the message overhead. A CGM computation/communication round corresponds to a BSP superstep with communication cost $g \frac{N}{p}$ (plus the above “packing requirement”). Finding an optimal algorithm in the coarse grained multicomputer model is equivalent to minimizing the number of communication rounds as well as the total local computation time. The CGM model has the advantage of producing results which correspond much better to the actual performance of implementations on commercially available parallel machines. In addition to minimizing communication and computation volume, it also minimizes important other costs like message overheads and processor synchronization.

In this paper, we present parallel CGM algorithms for detecting bipartite graphs and convex bipartite graphs. The algorithms require $O(\log p)$ and $O(\log^2 p)$ communication rounds, respectively, and linear sequential work per round. They assume that the local memory per processor, N/p , is larger than p^ϵ for some fixed $\epsilon > 0$. This assumption is true for all commercially available multiprocessors. Our results imply BSP algorithms with $O(\log p)$ supersteps, $O(g \log(p) \frac{N}{p})$ communication time, and $O(\log(p) \frac{N}{p})$ local computation time.

The algorithm for detecting bipartite graphs is fairly simple and is essentially a combination of tools developed in [3]. The larger part of this paper deals with the problem of detecting *convex* bipartite graphs. This is clearly a much harder problem. It has been extensively studied in the literature and is closely linked to the *consecutive ones* problem for $(0,1)$ -matrices as well as chordal graph recognition [1,4,5,11,12,13,14,15].

Our algorithm for determining whether a bipartite graph is convex includes a novel, coarse grained parallel, version of the *PQ tree* data structure introduced by Booth and Lueker [1]. Hence, our algorithm also solves, with the same time complexity as indicated above, the problem of testing the consecutive-ones pro-

perty for $(0, 1)$ -matrices as well as the chordal graph recognition problem. These, in turn, have numerous applications in graph theory, DNA sequence assembly, database theory, and other areas. [1,4,5,11,12,13,14,15]

2 Detecting *Bipartite* Graphs

In this section, we present a simple CGM algorithm for detecting bipartite graphs. It is a straight-forward combination of tools developed in [3].

Algorithm 1 Detection of Bipartite Graphs

Input: A Graph $G = (V, E)$ with vertex set V and edge set E , $|G| = N$, stored on a CGM with p processors and $O(N/p)$ memory per processor; $N/p \geq p^\epsilon$ for some fixed $\epsilon > 0$. V and E are arbitrarily distributed over the memories of the CGM. **Output:** A Boolean indicating whether G is a bipartite graph and, if it is, a partition of V into two disjoint set A and B such that $E \subseteq ((A \times B) \cup (B \times A))$.

- (1) Compute a spanning forest of G [3].
- (2) For each tree in the forest, select one arbitrary node as the root. Apply the CGM Euler Tour algorithm in [3] to determine the distance between each node and the root of its tree. Classify the nodes into two groups: the nodes with an odd numbered distance to the root, and the nodes with an even numbered distance to the root.
- (3) Each processor examines the edges stored in its local memory. If any such edge has two vertices that belong to the same group, the result for that processor is "failure"; otherwise, the result is "success".
- (4) By applying CGM sort [10] to all "failure" / "success" values, it is determined whether there was any processor with a "failure" result. If there was any "failure", the graph G is *not* bipartite. Otherwise, G is a bipartite graph, and the two groups of vertices identified in Step 2 are the sets A and B .

Theorem 1. *Algorithm 1 detects whether $G = (V, E)$, $|G| = N$, is a bipartite graph and, if so, partitions E into sets A and B such that $E \subseteq ((A \times B) \cup (B \times A))$ in $O(\log p)$ communication rounds and $O(\frac{N}{p})$ local computation per round on a CGM with p processors and $O(\frac{N}{p})$ memory per processor, $\frac{N}{p} \geq p^\epsilon$ for some fixed $\epsilon > 0$.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. □

3 Detecting *Convex* Bipartite Graphs

We now turn our attention to the problem of testing whether a given bipartite graph is a convex bipartite graph. The sequential solution, presented by Booth and Lueker [1], introduced a data structure called *PQ-tree*. Our coarse grained parallel solution will include a novel coarse grained parallel version of the PQ-tree. We will first review Booth and Lueker's PQ-tree definition.

Definition 3. A tree T is a PQ-tree if every internal node of T can be classified as either a P-node or a Q-node. A P-node is an internal node that has at least 2 children, and the children can be permuted arbitrarily. A Q-node is an internal node that has at least 3 children, and the children can only be permuted in two ways: the original order or the reverse order. The leaves of the PQ-tree are elements of a universal set $S = \{a_1, \dots, a_n\}$, usually called the ground set.

The order of the ground set in the PQ-tree, from left to right, is called its *frontier*. The frontier of a PQ-tree is clearly a permutation of the ground set. Given a PQ-tree T and using only permissible permutations of its internal nodes, we can generate a number of permutations of S . We will denote with $L(T)$ the set of all these permissible permutations. A PQ-tree T' is *equivalent* to T if T' can be transformed into T using only permissible permutations of the internal nodes (if $L(T')$ and $L(T)$ have the same elements).

Given a set $A \subset S$, we say that $\lambda \in L(T)$ *satisfies* A if all elements of A appear consecutively in λ . The main operation on a PQ-tree T is called *reduce*: given a *reduction* set $\mathcal{A} = \{A_1, \dots, A_k\}$ of subsets of S and a PQ-tree T , we want obtain a PQ-tree T' , if it exists, such that each permutation in $L(T')$ satisfies every A_i , $1 \leq i \leq k$.

Let $m = \sum_{i=1}^k |A_i|$ and $N = n + m$. In order to store T and \mathcal{A} , we require a coarse grained multicomputer with p processors and N/p local memory per processor.

Two particular PQ-trees are the *universal* and the *empty* tree: the first one has only one internal node (the root of T) and that internal node is a P-node; the second one (also called a *null* PQ-tree) is used to represent an impossible reduction, that is when it is impossible to reduce a PQ-tree with respect to a given reduction set.

3.1 Multiple *Disjoint* Reduce Operations on a PQ-Tree

In this section, we will present a coarse grained parallel algorithm for the special case of performing multiple *disjoint* reductions on a PQ-tree. We will then use this solution to develop the general algorithm in the subsequent section. More precisely, given a PQ-tree T we will first study how to perform the *reduce* operation for a set $\mathcal{A} = \{A_1, \dots, A_k\}$ of subsets of the universal set S where A_1, \dots, A_k are disjoint. We shall refer to our algorithm as *Algorithm MDReduce*. For ease of discussion, each set A_i is assigned a unique color, and we color the leaves of the PQ-tree accordingly. Some of the PQ-tree definitions used are from [1,12].

We start with a pre-processing phase which extends the coloring δ of the leaves to a coloring Δ of all nodes of the PQ-tree T . For an internal node v of T , we say that a color is *complete* at v if all the leaves with that color are descendants of v . We say a color is *incomplete* at v if some, but not all, of the leaves of that color are descendants of v . We say that a color *covers* v if all the leaves below v are of that color, and that v is *uncovered* if no color covers v . Let $LCA(c)$ be the lowest common ancestor of all leaves with color c . Let

$COLORS(v)$ denote the set of colors assigned to leaves that are descendants of v . Let $INC(v)$ be the set of colors which are incomplete at v . Then $INC(v) = COLORS(v) - \{c: LCA(c) \text{ is a descendent of } v\}$.

Algorithm 2 Pre-Processing the PQ-Tree

Input: The original PQ-tree T .

Output: The original PQ-tree T in which each node is assigned a "coloring" Δ , or, if failure occurs, a null tree.

- (1) Apply the coarse grained parallel Lowest Common Ancestor (LCA) algorithm [3].
- (2) Expand T into a binary tree B .
- (3) Perform tree contraction on B ; see [3]. For each node v_b in B , let v_p be the node in T from which v_b is created. Let w_1 and w_2 be the children of v_b . The operation for the tree contraction is $INC(v_b) = INC(w_1) \cup INC(w_2) - \{c: LCA(c) \text{ is a descendent of } v_p\}$. If at any point the size of INC is more than two, stop and return a null tree.
- (4) Let c_v be a new color unique to node v . Each processor, for all its nodes, v , calculates $\Delta(v) = \langle c_1, c_2 \rangle$ as follows: If two colors are incomplete at v , then c_1 and c_2 are these colors. If only one color c is incomplete at v but c does not cover v , then $c_1 = c$ and $c_2 = c_v$. If one color c is incomplete at v and covers v , then $c_1 = c_2 = c$. If no color is incomplete at v , then $c_1 = c_2 = c_v$.

Lemma 1. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm 2 can be completed in $O(\log p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. □

A node v in a PQ-tree is *orientable* if it is a Q-node and the two colors in its $\Delta(v) = \langle c_1, c_2 \rangle$ are different, i.e. $c_1 \neq c_2$.

For a color c , define $h_v(c) = \begin{cases} c & \text{if } c \in INC(v) \\ c_v & \text{if } c \notin INC(v) \end{cases}$

For a PQ-tree T where w_1 and w_k are the leftmost and rightmost elements, respectively, of the frontier $fr_T(v)$, let $l_T = h_v(\delta(w_1))$ and $r_T = h_v(\delta(w_k))$. If $lr_T[v] = \langle l_T[v], r_T[v] \rangle$ then we use the following notation: $\langle a, b \rangle \sim \langle a', b' \rangle$ if $\{a, b\} = \{a', b'\}$.

Algorithm 3 Processing P-Nodes

Input: The PQ-Tree output from Algorithm 2.

Output: The original PQ-tree T in which all the P-nodes have been processed, or, if failure occurs, a null tree.

- (1) If the input PQ-tree T is a null tree, return T .
- (2) Each processor sets variable *FAILURE* to *FALSE*
- (3) Each processor, for each P-node v , reorder the children of v such that for each color c all children covered by c are consecutive.
- (4) Each processor, for each P-node v and each color c , if there are at least two children covered by c (and at least one child not covered by c) then insert a new P-node w_c between these c -covered children and v .

- (5) Each processor, for each P-node v , constructs an auxiliary graph G_v whose nodes are the children of v and where for each color c there is an edge between children v_i and v_j at which c is incomplete if v_i or v_j is covered by c , or there is no child covered by c . If any node has more than 2 neighbors, set *FAILURE* to *TRUE* to indicate a failure condition.
- (6) Perform a multi-broadcast of the variable *FAILURE*. If any of the broadcast values is *TRUE*, return a null tree.
- (7) Each processor uses list-ranking to identify the connected components of each G_v and verifies that each of these connected components is a simple path. If any of these components is a cycle, set *FAILURE* to *TRUE* to indicate a failure condition. We call these paths *color chains*.
- (8) Perform a multi-broadcast of the variable *FAILURE*. If any of the broadcast values is *TRUE*, return a null tree.
- (9) Each processor, for each color chain χ containing at least 2 nodes, chooses one of the 2 orientations of χ arbitrarily. Reorder the children of v so that the nodes of χ are consecutive, and insert a new Q-node between these nodes of v .
- (10) Each processor, for each P-node v , let $S = \{v_i : v_i \text{ is a child of } v, \text{ and } INC(v_i) = \emptyset\}$. If every child of v is in S , then return. Otherwise, reorder the children of v to make S consecutive, insert a new P-node v' between v and the subset S (if $|S| > 1$), and rename v to be a Q-node.

Lemma 2. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm 3 can be completed in $O(\log p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. \square

For each Q-node v , we define an orientation $\overline{LR}(v)$ which is either $\Delta(v_i)$ or $\Delta(v_i)^R$. Note that if $\Delta(v_i) = \langle c_1, c_2 \rangle$ than $\Delta(v_i)^R = \langle c_2, c_1 \rangle$. For $\langle a, b \rangle \sim \langle a', b' \rangle$ and $a \neq b$, we define $\langle a, b \rangle \text{ swap } \langle a', b' \rangle$ equals *TRUE* if $\langle a, b \rangle = \langle b', a' \rangle$, *FALSE* if $\langle a, b \rangle \neq \langle a', b' \rangle$. For a Q-node v , *flip* is defined as the operation which re-orders all its children in reverse order.

Algorithm 4 Processing Q-Nodes

Input: The PQ-tree output from Algorithm 3.

Output: The original PQ-tree T in which all the Q-nodes have been processed, or, if failure occurs, a null tree.

- (1) If the input PQ-tree T is a null tree, return T .
- (2) Each processor sets variable *FAILURE* to *FALSE*
- (3) Each processor, for each Q-node v and children be v_1, \dots, v_s , assign to each $\overline{LR}[v_i]$ either $\Delta(v_i)$ or $\Delta(v_i)^R$ such that every color in the sequence $\overline{LR}[v_1], \dots, \overline{LR}[v_s]$ occurs consecutively, and such that $h_v(\langle \overline{L}[v_1], \overline{R}[v_s] \rangle) \sim \Delta(v)$. If this is impossible, set *FAILURE* to *TRUE* to indicate a failure condition, otherwise, set $LR[v]$ to $h_v(\langle \overline{L}[v_1], \overline{R}[v_s] \rangle)$.
- (4) Perform a multi-broadcast of the variable *FAILURE*. If any of the broadcast values is *TRUE*, return a null tree.
- (5) Each processor for each node v : if v is orientable, then set $OPP[v]$ to $LR[v]$ swap $\overline{LR}[v]$, otherwise, set $OPP[v]$ to *FALSE*.

- (6) Each processor for each node v : set $REV[v]$ to $\bigoplus_{u \text{ is an ancestor of } v} OPP[u]$ (Note: \bigoplus denotes "exclusive-or").
- (7) For each orientable node v , if $REV[v]$ is $TRUE$, then flip v .

Lemma 3. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm 4 can be completed in $O(\log p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. \square

Algorithm 5 Post-Processing the PQ-Tree

Input: The PQ-tree output from Algorithm 4, with all R-nodes renamed.

Output: Result of Algorithm *MDReduce*.

- (1) If T is a null tree, return.
- (2) Each processor temporarily cuts the links of its Q-nodes to their parents.
- (3) Each processor performs pointer jumping for all its nodes that are children of R-nodes to determine their lowest Q-node ancestor.
- (4) Each processor restores the links cut in Step 2.
- (5) Each processor eliminates its R-nodes by setting the parents of their children to their lowest Q-node ancestors.

Lemma 4. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm 5 can be completed using in $O(\log p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. \square

Theorem 2. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm *MDReduce* performs a multiple disjoint reduce for a PQ-tree T in $O(\log p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

Proof. Omitted due to page restrictions. To be included in the full version of this paper. \square

3.2 Multiple *General Reduce* Operations on a PQ-Tree

Using the coarse grained parallel *MDreduce* algorithm presented in the previous section, we will now develop coarse grained parallel algorithm for the general *MReduce* operation: given a PQ-tree T over the ground set S with n elements, perform the reduce operation for an arbitrary reduction sets $\mathcal{A} = \{A_1, \dots, A_k\}$

Our CGM algorithm for the general *MReduce* operation consists of two phases. In the first phase, we execute $3 \log p$ times an algorithm which is a CGM

implementation of a PRAM algorithm proposed by Klein [12]. We call this operation $\text{MREDUCE}_1(T, \{A_1, \dots, A_k\}, 0)$. Our contribution here is the implementation of the various shared memory PRAM steps on a distributed memory CGM, which is non trivial. After this first phase, we have reduced the problem to one in which we are left with a set of smaller PQ-trees over ground sets whose size is at most n/p . Hence, each tree can be stored in the local memory of one processor. However, we can not guarantee that all the *reduction sets* of these PQ-trees do also fit in the local memory of one processor. In the second phase of our algorithm, we use a merging strategy to complete the algorithm. We will refer to this phase as the *Merging Phase*.

First Phase: For a node v of a PQ-tree, $\text{leaves}_T(v)$ denotes the set of pendant leaves of v , i.e. leaves of T having v as ancestor. Let $\text{lca}_T(A)$ denote the least common ancestor in T of the leaves belonging to A . Suppose that $v = \text{lca}_T(A)$ has children v_1, \dots, v_s in order. We say A is *contiguous* in T if either (1) v is a Q-node, and for some consecutive subsequence v_p, \dots, v_q of the children of v , $A = \bigcup_{p \leq i \leq q} \text{leaves}_T(v_i)$, or (2) v is a P-node or a leaf, and $A = \text{leaves}_T(v)$.

Suppose that E is contiguous in T . $T|E$ denotes the subtree consisting of $\text{lca}_T(E)$ and those children of $\text{lca}_T(E)$ whose descendants are in E (it is still a PQ-tree whose ground set is E). For a set A , define

$$A_i|E = \begin{cases} A_i \cap E & \text{if } A_i \cap E \neq E \\ \emptyset & \text{if } A_i \cap E = E \end{cases}$$

Let \star_E denote $\text{lca}_T(E)$. T/E denotes the subtree of T obtained by omitting all the proper descendants of $\text{lca}_T(E)$ that are ancestors of elements of E (it is still a PQ-tree whose ground set is $S - E \cup \{\star_E\}$). For a set A , define

$$A_i/E = \begin{cases} A_i - E \cup \{\star_E\} & \text{if } A_i \supseteq E \\ A_i - E & \text{otherwise} \end{cases}$$

Algorithm 6 $\text{MREDUCE}_1(T, \{A_1, \dots, A_k\}, i)$:

- (1) If $i = 3 \log p$, return.
- (2) Purge the collection of input sets A_i of empty sets. If no sets remain, return.
- (3) Let n be the size of the ground set of T . If $n \leq 4$, carry out the reduction one by one. If the size of the input is smaller than the size of the local memory of the processors, than solve the problem sequentially using the Booth and Lueker's algorithm.
- (4) Otherwise, let \mathcal{A} be the family of (nonempty) sets A_i . Let \mathcal{S} consist of the sets A_i such that $|A_i| \leq n/2$. We call such sets "small". Let \mathcal{L} be the remaining, "large", sets in \mathcal{A} . Find the connected components of the intersection graph of \mathcal{A} , find a spanning forest of the intersection graph of \mathcal{S} , and find the intersection $\cap \mathcal{L}$ of the large sets.
- (5) Proceed according to one of the following cases:
 - (a) The intersection graph of \mathcal{A} is disconnected. In this case, let $\mathcal{C}_1, \dots, \mathcal{C}_r$ be the connected components of \mathcal{A} . For $i = 1, \dots, r$, let E_i be the union of sets in the connected component \mathcal{C}_i . Call MDREDUCE to reduce T with respect to the disjoint sets E_1, \dots, E_r . Next, for each $i = 1, \dots, r$ in parallel, recursively call $\text{MREDUCE}_1(T|E_i, \mathcal{C}_i, i + 1)$.

- (b) The union of sets in some connected component of \mathcal{S} has cardinality at least $n/4$. In this case, from the small sets making up this large connected component, select a subset whose union has cardinality between $n/4$ and $3n/4$. Let E be this union, and call $\text{SUBREDUCE}(T, E, \{A_1, \dots, A_k\}, i)$.
- (c) The cardinality of the intersection of the large sets is at most $3n/4$. In this case, from the large sets choose a subset whose intersection has cardinality between $n/4$ and $3n/4$. Let E be this intersection, and call $\text{SUBREDUCE}(T, E, \{A_1, \dots, A_k\}, i)$.
- (d) The other case do not hold. In this case, let E be the intersection of the large sets, and call $\text{SUBREDUCE}(T, E, \{A_1, \dots, A_k\}, i)$.

In the full version of this paper, we show how to implement the above on a coarse grained multicomputer with p processors and $O(\frac{n}{p})$ storage per processor in $O(\log p)$ communication rounds. The non trivial parts are Step 4, Step 5b, the computation of E , T/E , and $T|E$, as well as the SUBREDUCE operation. The latter involves another operation called GLUE . Due to page restrictions, we can not present this part of our result in the extended abstract. Instead, we give one example which shows the coarse grained parallel computation of the set E in Step 5(b) of Algorithm 6.

Algorithm 7 Computation of E .

Input: The set \mathcal{S} and the spanning forest of its intersection graph.

- (1) In order to find a connected component \mathcal{C} in the spanning forest of \mathcal{S} , such that the union of its sets has cardinality at least $n/4$, order all the components according to the labeling given by the coarse grained parallel *spanning forest* algorithm [3].
- (2) Sort each component with respect to the values of its elements and mark as "valid" only one element per distinct value.
- (3) Sort again with respect to the components' labels. Compute the cardinality of the union of the elements of each component (that is the *size* of each component), with a prefix-sum computation, counting only the "valid" elements. (Hence, we do not count twice the elements with same values and compute correctly the cardinality of the union.)
- (4) If a processor finds a component whose size is $\geq n/4$, then it broadcasts the label of this component. Otherwise it broadcast a "not-found" message.
- (5) If everybody sent "not-found", go to step 4(c) of MREDUCE algorithm. Otherwise, among all the labels received in the previous step, choose as \mathcal{C} the component with the smallest label.
- (6) For each of the sets comprising \mathcal{C} , compute the distance in the spanning tree (from the root) using the coarse grained parallel Euler-tour technique [3].
- (7) Sort the sets according to distance, and let B_1, \dots, B_s be the sorted sequence. Sort each sets with respect to the values of its elements and mark as "valid" only one element per distinct value. Sort the sets again, according to distance, and let \hat{i} be the minimum i such that $|\bigcup_{j=1}^i B_j| \geq n/4$. (\hat{i} can be found with a prefix-sum computation on the "valid" elements.) Broadcast \hat{i} .
- (8) Mark all "valid" elements in $B_1, \dots, B_{\hat{i}}$ as elements of E .

Second Phase: Consider the tree R of recursive calls in MREDUCE_1 . We observe that, after $l = 3 \log_{4/3} p$ levels of R (when the first part of our algorithm

stops), the sizes of the ground sets associated with the nodes in R at level l are at most n/p . This is due to the fact that the descendants of a node u in R that are 3 levels below u are *smaller* than u by approximately a factor $3/4$. More precisely, if $n(u)$ denotes the size of the ground set of $T(u)$ (the subtree rooted at u) then, for every node w three levels below u , $n(u) \leq 3n(w)/4 + 1$. Hence, each PQ-tree obtained at the end of the first phase fits completely into the local memory of one processor.

Unfortunately, the same argument does not hold for the reduction sets. Recall that $m = \sum_{i=1}^k |A_i|$. Let u be an internal node of R , A_{u_1}, \dots, A_{u_j} its reduction sets, and $m_u = \sum_{i=1}^j |A_{u_i}|$. Since the sizes of the reduction sets of the children of u depend strictly on the A_{u_i} and on how they intersect with the set E computed for u , it is possible that the A_{u_i} are split in an unbalanced way. That is, we can have $\sum_{i=1}^j |A_{u_i} \cap E| = O(m_u)$ and $\sum_{i=1}^j |A_{u_i} \setminus E| = O(1)$ (or vice versa). If this continues up to level $3 \log p$ of R , it is possible that for a recursive call associated with a node v at level l , $\sum_{i=1}^f |A_v i| > m/p$.

Therefore, while the ground set of $T(v)$, and hence $T(v)$, can fit in one processor, the reduction sets could possibly not. Thus, at this point of the computation, we can not simply use the sequential algorithm of Booth and Lueker [1] for completing the reduction.

Our idea for solving this problem is the following. Let us consider a node v at level l in R that has $m_u > m/p$. Since, at any level of recursion, the sum of the sizes of all reduction sets is at most $2m$, we can create α_v copies of $T(v)$, with $\alpha_v = \lfloor \frac{m_v}{m/p} \rfloor$. We observe that

$$\sum_{v \in l} \alpha_v = \sum_{v \in l} \lfloor \frac{m_v}{m/p} \rfloor \leq \sum_{v \in l} \frac{m_v}{m/p} \leq \frac{p}{m} \sum_{v \in l} m_v \leq \frac{p}{m} \cdot 2m = 2p.$$

Hence, we require at most two copies per processor. The reduction problem of each node v at level l of R will be solved by the α_v processors that have copies of $T(v)$. The next step is the *distribution* of the reduction sets associated to v among these α_v processors. Each of these α_v processors can solve locally the problem of reducing $T(v)$ with respect to the reduction sets that it has stored, using Booth and Lueker's algorithm [1]. For each processor, let $T'(v)$ refer to this reduced tree. Now, we need to merge these α_v trees, $T'(v)$. More precisely, we need to compute a PQ-tree $\widehat{T}(v)$ such that $L(\widehat{T}(v)) = L(\overline{T}(v))$, where $\overline{T}(v)$ is the PQ-tree that we would have obtained by reducing $T(v)$ directly with respect its reduction sets. For the construction of $\widehat{T}(v)$, we merge the $T'(v)$ trees in a binary tree fashion.

Algorithm 8 Merging Phase

Input: h PQ-trees $T(i)$, with $|T(i)| \leq n/p$ and $\sum_i |T(i)| \leq n$, and their reduction sets.

Output: The $T(i)$ reduced with respect their reduction sets.

- (1) Let m_i be the sum of the sizes of the reduction sets of T_i . Make $\alpha_i = \lfloor \frac{m_i}{m/p} \rfloor$ copies of each $T(i)$. Distribute the reduction sets of each T_i between the processors that have the copies of $T(i)$.
- (2) Each processor executes the sequential algorithm [1] for its PQ-trees with the reduction sets that it has stored. Let $T'(i)$ refer to the trees obtained.

- (3) The α_i processors associated with each $T(i)$ merge the $T'(v)$ trees in a binary tree fashion. More details are outlined below.

The following Theorem 3 shows that the merge operation in Step 3 of Algorithm 8 reduces to a tree intersection operation. We have designed a CGM algorithm for tree intersection which implements Step 3 of Algorithm 8. Due to page restrictions, we can not include a description of our tree intersection algorithm in this extended abstract. It will be included in the full version of this paper.

Theorem 3. *Let T be a PQ-tree over the ground set S and let T' be a copy of T . Let T^* and T'^* be the result of the reduction of T with respect to $\{A_1, \dots, A_r\}$ and of T' with respect to $\{B_1, \dots, B_t\}$, respectively. Let \bar{T} be the PQ-tree obtained by reducing T with respect to $\{A_1, \dots, A_r, B_1, \dots, B_t\}$. Then,*

$$\lambda \in L(\bar{T}) \Leftrightarrow \lambda \in L(T^*) \cap L(T'^*).$$

Proof. $L(\bar{T})$ is the intersection of the sets of all orderings that satisfy $A_1, \dots, A_r, B_1, \dots, B_t$, and $L(T)$. $L(\bar{T})$ is always the same, independently of the order in which we reduce T . If $\lambda \in L(\bar{T})$, then λ must belong to the intersection between the set of all orderings that satisfy A_1, \dots, A_r and $L(T)$ and it must also belong to the intersection between the set of all orderings that satisfy B_1, \dots, B_t and $L(T)$, that is $\lambda \in L(T^*)$, $\lambda \in L(T'^*)$ and $\lambda \in L(T)$. Hence λ belongs to $L(T^*) \cap L(T'^*)$. The reverse can be shown analogously. \square

In summary, we obtain

Theorem 4. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, Algorithm MReduce performs a reduce operation for a PQ-tree T in $O(\log^2 p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

3.3 Convex Bipartite Graphs

Recall the definition of convex bipartite graphs (Definition 2). Given a bipartite graph $G = (A, B, E)$ with $A = \{a_1, a_2, \dots, a_k\}$ and $B = \{b_1, b_2, \dots, b_n\}$. Let $\mathcal{A} = \{A_1, \dots, A_k\}$ where $A_i = \{b \in B : (a_i, b) \in E\}$, and let T be a PQ-tree over the ground set B consisting of a root with children b_1, b_2, \dots, b_n . The problem of determining whether G is convex and, if this is the case, computing the correct ordering of the elements in B is equivalent to the *MReduce* operation on T with respect to \mathcal{A} .

Theorem 5. *On a coarse grained multicomputer with p processors and $O(\frac{N}{p})$ storage per processor, the problem of determining whether G is convex (and computing the correct ordering of the elements in B) can be solved in $O(\log^2 p)$ communication rounds with $O(\frac{N}{p})$ local computation per round.*

References

1. K.S. Booth and G.S. Lueker, "Testing for the Consecutive Ones Property, Interval Graphs, and Graph Planarity Using PQ-Tree Algorithms," in *Journal of Computer and System Sciences*, vol. 13, pages 335-379, 1976.
2. P. Bose, A. Chan, F. Dehne, and M. Latzel, "Coarse grained parallel maximum matching in convex bipartite graphs," in Proc. *13th International Parallel Processing Symposium (IPPS'99)*, 1999.
3. E. Cáceres, F. Dehne, A. Ferreira, P. Flocchini, I. Rieping, A. Roncato, N. Santoro, and S.W. Song, "Efficient parallel graph algorithms for coarse grained multicomputers and BSP," in Proc. *24th International Colloquium on Automata, Languages and Programming (ICALP'97)*, Bologna, Italy, 1997, Springer Verlag Lecture Notes in Computer Science, vol. 1256, pages 390-400.
4. L. Chen and Y. Yesha, "Parallel recognition of the consecutive ones property with applications," *J. Algorithms*, vol. 12, no. 3, pages 375-392, 1991.
5. L. Chen, "Graph isomorphism and identification matrices: parallel algorithms," *IEEE Trans. on Parallel and Distr. Systems*, vol. 7, no. 3, March 1996, pages 308 ff.
6. F. Dehne (Ed.), "Coarse grained parallel algorithms," Special Issue of *Algorithmica*, vol. 24, no. 3/4, 1999, pages 173-426.
7. F. Dehne, A. Fabri, and A. Rau-Chaplin, "Scalable Parallel Geometric Algorithms for Coarse Grained Multicomputers," in Proc. *ACM 9th Annual Computational Geometry*, pages 298-307, 1993.
8. F. Dehne, A. Fabri, and C. Kenyon, "Scalable and Architecture Independent Parallel Geometric Algorithms with High Probability Optimal Time," in Proc. *6th IEEE Symposium on Parallel and Distributed Processing*, pages 586-593, 1994.
9. F. Dehne, X. Deng, P. Dymond, A. Fabri, and A.A. Kokhar, "A randomized parallel 3D convex hull algorithm for coarse grained multicomputers," in Proc. *ACM Symposium on Parallel Algorithms and Architectures (SPAA'95)*, pages 27-33, 1995.
10. M.T. Goodrich, "Communication efficient parallel sorting," *ACM Symposium on Theory of Computing (STOC)*, 1996.
11. X. He and Y. Yeshua, "Parallel recognition and decomposition of two terminal series parallel graphs," *Information and Computation*, vol. 75, pages 15-38, 1987.
12. P. Klein, *Efficient Parallel Algorithms for Planar, Chordal, and Interval Graphs* PhD. Thesis, MIT, 1988.
13. P. Klein. "Efficient Parallel Algorithms for Chordal Graphs". *Proc. 29th Symp. Found. of Comp. Sci., FOCS* 1989, pages 150-161.
14. P. Klein. "Parallel Algorithms for Chordal Graphs". In *Synthesis of parallel algorithms*, J.H. Reif (editor). Morgan Kaufmann Publishers, 1993, pages 341-407.
15. A.C. Tucker, "Matrix characterization of circular-arc graphs," *Pacific J. Mathematics*, vol. 39, no. 2, pages 535-545, 1971.
16. L. Valiant, "A bridging model for parallel computation," *Communications of the ACM*, vol. 33, no. 8, August 1990.