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Scalable Community Detection using Distributed Louvain Algorithm

A Thesis

Submitted to the the Graduate Faculty of the  
University of New Orleans  
in partial fulfillment of the  
requirements for the degree of

Master of Science  
in  
Computer Science  
HPC and Big Data Analytics

by

Naw Safrin Sattar

B.S. Bangladesh University of Engineering and Technology, 2016

May, 2019

My thesis work is wholeheartedly dedicated to my parents who tried throughout their life to provide me with the best education possible.

To my beloved husband without whose encouragement and support it was not possible to aspire for higher studies at abroad. I am thankful to my family for keeping faith in me and providing me with the opportunity.

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# Abstract

Community detection (or clustering) in large-scale graph is an important problem in graph mining. Communities reveal interesting characteristics of a network. Louvain is an efficient sequential algorithm but fails to scale emerging large-scale data. Developing distributed-memory parallel algorithms is challenging because of inter-process communication and load-balancing issues. In this work, we design a shared memory-based algorithm using OpenMP, which shows a 4-fold speedup but is limited to available physical cores. Our second algorithm is an MPI-based parallel algorithm that scales to a moderate number of processors. We also implement a hybrid algorithm combining both. Finally, we incorporate dynamic load-balancing in our final algorithm DPLAL (Distributed Parallel Louvain Algorithm with Load-balancing). DPLAL overcomes the performance bottleneck of the previous algorithms, shows around 12-fold speedup scaling to a larger number of processors. Overall, we present the challenges, our solutions, and the empirical performance of our algorithms for several large real-world networks.

**Keywords:** Community Detection; Louvain Method; Parallel Algorithms; MPI; OpenMP; Load-balancing; Graph Mining

# Chapter 1

## Introduction

Parallel computing plays a crucial role in processing large-scale graph data [1, 2, 3, 4]. The problem of community detection in graph data arises in many scientific domains [5], e.g., sociology, biology, online media, and transportation. Due to the advancement of data and computing technologies, graph data is growing at an enormous rate. For example, the number of links in social networks [6, 7] is growing every millisecond. Processing such graph big data requires the development of parallel algorithms [2, 8, 3, 9, 4]. Existing parallel algorithms are developed for both shared memory and distributed memory based systems. Each method has its own merits and demerits. Shared memory based systems are usually limited by the moderate number of available cores [10]. Conventional multi-core processors can exploit the advantages of shared-memory based parallel programming. The increase in physical cores is restricted by the scalability of chip sizes. Shared global address space size is also limited because of memory constraint. On the other hand, a large number of processing nodes can be used in distributed-memory systems. Although distributed memory based parallelism has the freedom of communicating among processing nodes through passing messages, an efficient communication scheme is required to overcome communication overhead. We present a compar-

ative analysis of our shared and distributed memory based parallel Louvain algorithms, their merits and demerits. We have disclosed the problems arisen in communication among processes for distributed memory based parallelism [11]. We also develop a hybrid parallel Louvain algorithm using the advantage of both shared and distributed memory based approaches. The hybrid algorithm gives us the scope to balance between both shared and distributed memory settings depending on available resources. Load balancing is crucial in parallel computing. A straight-forward distribution with an equal number of vertices per processor might not scale well [3]. We also find that load imbalance also contribute to a higher communication overhead for distributed memory algorithms [8]. A dynamic load balancing [9, 12] approach can reduce the idle times of processors leading to increased speedup. Finding a suitable load balancing technique is a challenge in itself as it largely depends on the internal properties of a network and the applications [13]. We present DPLAL, an efficient algorithm [14] for distributed memory setting based on a parallel load balancing scheme and graph partitioning.

# Chapter 2

## Related Works

There exists a rich literature of community detection algorithms [15, 16, 1, 17, 18, 19, 20, 21]. Louvain method [15] is found to be one of the most efficient sequential algorithms [18, 21]. In recent years, several works have been done for paralleling Louvain algorithm and a majority of those are shared memory based implementations. These implementations demonstrate only a moderate scalability.

A Template has been proposed to parallelize the Louvain Method for Modularity Maximization with a shared-memory parallel algorithm [16] using OpenMP. Maximum modularity has been found by parallel reduction. They have combined communities to supervertices using parallel mergesort. They run their experimental setup on two sets of LFR benchmarks of 8,000 and 10,000 vertices which is a very small number compared to dealing with large networks. Another shared-memory implementation is done using a hierarchical clustering method with adaptive parallel thread assignment [22]. They have showed the granularity of threads could be obtained adaptively at run-time based on the information to get the maximal acceleration in the parallelization of Louvain algorithm. They have computed the gained modularity of adding a neighbor node to the community by assigning some threads in parallel. Dynamic thread assignment of OpenMP has been disabled to

let the algorithm adaptively choose the number of threads. For upto 32 cores, the speedup is not significant compared to the previous implementations PLM [17] [23] and CADs [24].

One Distributed-Memory implementation has been done in [1]. Graph partitioning has been done using PMETIS . They have only parallelized the first level for speedup. Each MPI process locally ignores cross partition edges, that might be an issue with accuracy. They have used three different vertex ordering strategies but none of those are effective enough considering performance. Although they get a speedup in their approach but it has been flattened for most of the graphs after scaling up to 16 - 32 processors.

One of the fastest shared memory implementations is Grappolo software package [25, 26], which is able to process a network with 65.6M vertices using 20 compute cores. One of the MPI based parallel implementations [1] of Louvain method reported scaling for only 16 processors. Later, in [27] the authors could run large graphs with 1,000 processing cores for their MPI implementation but did not provide a comprehensive speedup results. Their MPI+OpenMP implementation demonstrated about 7-fold speedup on 4,000 processors. But the paper uses a higher threshold in lower levels in Louvain method to terminate the level earlier and thus minimized the time contributing to their higher speedup. The work also lacks on the emphasis on graph partitioning and balancing load among the processors. This is a clear contrast with our work where we focused on load balancing issue among others. Our work achieves comparable (or better in many cases) speedups using a significantly fewer number of processors than the work in [27].

# Chapter 3

## Background

Notations, Definitions, and computational model used are described in this section.

### 3.1 Notation

The network is denoted by  $G(V, E)$ , where  $V$  and  $E$  are the sets of vertices and edges, respectively. Vertices are labeled as  $V_0, V_1, \dots, V_{n-1}$ . We use the words node and vertex interchangeably as well as links and edges.  $P$  is the number of processors used in the computation, denoted by  $P_0, P_1, \dots, P_{N-1}$  where  $0, 1, 2, \dots, N-1$  refers to the rank of a processor. Terms frequently used throughout the paper, are enlisted in Table 3.1.

Table 3.1: Terminologies used in the work

| Symbol    | Meaning  |
|-----------|--|
| $G(V, E)$ | Graph network with $V$ = set of vertices and $E$ = set of edges      |
| in [c]    | sum of the weights of the links inside community, c                  |
| tot [c]   | sum of the weights of the links incident to vertices in community, c |
| n2c [i]   | community of vertex, i   |
| d(i,c)    | number of links from vertex, i to community, c                       |
| N         | total number of processors (World size)                              |
| $n =  V $ | total number of vertices (Network size)                              |

## 3.2 Louvain Algorithm for Community Detection

Louvain is a simple, heuristic method to extract the community structure of large networks based on modularity optimization [15]. It outperforms all other known community detection method in terms of computation time. Modularity,  $Q$  is calculated using Equation 3.1, where  $-1 < Q < 1$ . The meanings are described in Table 3.2. The algorithm is divided in two phases those are iteratively repeated .

$$Q = \frac{1}{2m} \sum_{ij} \left[ A_{ij} - \frac{k_i k_j}{2m} \right] \delta(c_i c_j) \quad (3.1)$$

Table 3.2: Symbols used for calculating Modularity in Equation 3.1 and Equation 3.2

| Symbol                        | Meaning  |
|-------------------------------|--|
| $Q$                           | Modularity   |
| $A$                           | Usual adjacency matrix   |
| $A_{ij}$                      | Link weight between nodes $i$ and $j$  |
| $m$                           | Total link weight in the network   |
| $k_i$                         | Sum of the link weights attached to node $i$   |
| $\frac{k_i}{2m}$              | Average fraction of weight that would be assigned to node $j$ , if node $i$ assigned its link weight randomly to other nodes in proportion to their own link weights |
| $A_{ij} - \frac{k_i k_j}{2m}$ | How strongly nodes $i$ and $j$ are connected in the real network, compared to how strongly connected we would expect them to be in a random network                  |
| $c_i$                         | Community to which node $i$ is assigned  |
| $\delta(c_i, c_j)$            | Kronecker delta. Value is 1 when nodes $i$ and $j$ are assigned to the same community. Otherwise, the value is 0   |
| $\Delta Q$                    | Gain in Modularity   |
| $\sum_{in}$                   | Sum of the weights of the links inside community $C$   |
| $\sum_{tot}$                  | Sum of the weights of the links incident to nodes in community $C$   |
| $k_{i,in}$                    | sum of the weights of the links from node $i$ to nodes in $C$  |

### 3.2.1 Modularity Optimization Phase

For each node  $i$  the neighbours  $j$  of  $i$  are considered and the gain in modularity,  $\Delta Q$  that would take place by removing  $i$  from its community and by placing it in the community of  $j$  is evaluated.  $\Delta Q$  obtained by moving an isolated node  $i$  into a community  $C$  is computed using Equation 3.2.

$$\Delta Q = \left[ \frac{\sum_{in} + k_{i,in}}{2m} - \left( \frac{\sum_{tot} + k_i}{2m} \right)^2 \right] - \left[ \frac{\sum_{in}}{2m} - \left( \frac{\sum_{tot}}{2m} \right)^2 - \left( \frac{k_i}{2m} \right)^2 \right] \quad (3.2)$$

The symbolic meanings are given in Table 3.2. When  $i$  is removed from its community  $C$  a similar equation is used. This continues repeatedly until no further improvement is achieved. This phase comes to an end when a local maxima of the modularity is attained.

### 3.2.2 Community Aggregation Phase

In this phase, a new network is formed whose nodes are the communities found during the first phase. The weights of the links between two new nodes are given by the sum of the weight of the links between nodes in the corresponding two communities. Links between nodes of the same community lead to self-loops for this community in the new network.

After completion of second phase, the first phase of the algorithm is reapplied to the resulting weighted network and iteration continues as long as positive gain in modularity is achieved.

### 3.3 Computational Model

At first, we develop share-memory based parallel algorithm using Open Multi-Processing (OpenMP) to eliminate the limitations of [16] with a different approach. Shared-memory based algorithms have some own limitations due to limited number of physical processing cores hindering high speedup. Therefore, we develop parallel algorithm for message passing interface (MPI) based distributed-memory parallel systems, where each processor has its own local memory. The processors do not have any shared memory, one processor cannot directly access the local memory of another processor, and the processors communicate via exchanging messages using MPI.

# Chapter 4

## Methodology

### 4.1 Shared Memory Parallel Louvain Algorithm

In shared memory based algorithms, there is a shared address space and multiple threads share this common address space. This shared address space can be used efficiently using lock and other synchronization techniques. The main hindrance behind the shared memory based systems is the limited number of processing cores. We parallelize the Louvain algorithm by distributing the computational task among multiple threads using Open Multi-Processing (OpenMP) framework. We parallelize the Louvain algorithm computational task-wise in a straight-forward approach. Whenever there is a need to iterate over the full network or even the neighbors of a node, considering the large network size, the work is done by multiple threads to minimize the workload and do the computation faster.

## 4.2 Distributed Memory Parallel Louvain Algorithm

Our aim is to compute the communities of the full network in a distributed manner. To serve the purpose, we distribute the full network among the processors in a balanced way, so that each processor can do its computation in a reasonable time and no one should wait for another. It is necessary because after each level of computation, they have to communicate with the root processor to generate the modularity of the full network. After each level of iteration, the network size decreases gradually and when the network size is considerably small, the final level is computed by a single processor which acts similar to the Louvain Sequential Algorithm.

### 4.2.1 Graph Partitioning

Let  $N$  be the number of processors used in the computation. The network is partitioned into  $N$  partitions, and each processor,  $P_i$  is assigned one such partition  $G_i(V_i, E_i)$ . Processor  $P_i$  performs computation on its partition  $G_i$ . The network data is given as input in a single disk file. While partitioning  $G(V, E)$ , if the vertex-id does not start with 0, then the vertices are renumbered from 0 to  $n_i - 1$ . The network is partitioned depending on the number of vertices of the network such that each processor gets equal number of vertices.  $N$  can vary depending on the system configuration and available resources.  $n$  is also of varied range. So, we cannot divide the vertices equally among processors when  $n \bmod N \neq 0$ . We distribute the remaining nodes starting with processor  $P_0$  and continue up to processor  $P_k$  ( $0 \leq k < N$ ), as long as the remainder lasts. Each processor has its own part of the network necessary to compute the communities in the partial

network. It is a very naive partitioning technique.

## 4.2.2 Community Detection

We have parallelized the sequential algorithm in such a way that each processor can compute its partial network's community with minimized communication among the processors. The following information are needed for each processor to complete its part of computation.

- Degree of each vertex within the partition
- Neighbors of each vertex
- Weight associated with each neighbor

In first phase, each processor scans through all neighboring vertices and identifies those in different processors. It then gathers the mentioned information by message passing among those processors. Then each processor locally computes the modularity of the partial network and does community detection. After computation, it sends information of each vertex's community to the processor acting as the root. The Root processor needs the value of *in*, *tot* arrays and total weight of the network to compute modularity of the full network. This full process is iterated several times as long as there is increase in modularity.

The output is stored as (node,community) for each level of iteration. It is also stored in an adjacency matrix format that is used to visualize the graph using Python's **networkx** library.

Algorithm 1 represents the pseudo-code of our approach. All the steps from Section 4.2.2.3 to 4.2.2.11 are done for each level of computation and repeated as long as we get increase in modularity value.

---

**Algorithm 1:** Our Parallel Louvain using MPI

---

**Data:** Input Graph  $G(V,E)$   
**Result:** (Vertex, Community) Pair

```
1 while increase in modularity do
2    $G(V, E)$  is divided into  $p$  processes;
3   Each  $graph\_i.bin$  contains  $\lceil \frac{n}{p} \rceil$  vertices and corresponding edges in
   adjacency list format;
4   for Each processor  $P_i$  (executing in parallel) do
5     Initialize_Graph();
6     Exchange_Starting_Node();
7     Gather_Neighbour_Info();
8     Compute_Community();
9     Exchange_Updated_Community();
10    Resolve_Community_Duality();
11    Exchange_Duality_Resolved_Community();
12    Find_Unique_Communities();
13    Compute_Modularity();
14    Generate_NextLevel_Graph();
15    if number_of_communities  $< i$  then
16      |  $i \leftarrow \frac{number\_of\_communities}{2}$ ;
17    end
18  end
19 end
```

---

#### 4.2.2.1 Graph Initialization

In this step, each processor  $P_i$  reads the input graph,  $graph\_name\_i.bin$ . If the node-id does not start with 0, then the nodes are renumbered from 0 to  $n_i - 1$ . Afterwards a quality object,  $q$  is created from graph,  $g$  which initializes the arrays  $tot$ ,  $in$  and  $n2c$ . Line 5, in Algorithm 1 does this.

#### 4.2.2.2 Exchange of Starting Node

In this step, each processor  $P_i$  sends own starting node-id to other processors,  $P_j$  in the communicating world.

$$P_i \xrightarrow{msg} P_j$$

where

$$i, j \in world, j = 0, 1, \dots, N - 1 \quad \wedge \quad i \neq j, \quad N = world\_size$$

$$msg \ni starting\_node_{P_i}$$

The number of Send and Receive operations are equal to  $N - 1$ . This step is mainly required because at later phases, we use this information to find out which node belongs to which processor at other steps of our calculation. If number of processors is always a factor of network size, we could have skipped this step by numerically figuring out the starting node for each processor. This is a very rare scenario and our world size can vary depending on the system configuration and available resources. Again, network size is also of varied range. So, we cannot divide the nodes equally among processors when

$$n \bmod N! = 0, \quad n = network\_size$$

So, we distribute the remaining nodes starting with processor  $P_0$  and continue up to processor  $P_k$ , as long as the remainder lasts. Here,  $0 \leq k < N$

#### 4.2.2.3 Collection of Neighbour Information

In this step. We scan through the neighbor list of all vertices and find out the neighbors those do not belong to current processor using the information of Section 4.2.2.2. Now, we need the following information of each vertex in the neighbor list:

- degree,

- weight,
- neighbor list with weights

These are necessary for the calculation in section 4.2.2.4. To gather this information, we need to know beforehand, how many send and receive operations are required to communicate with other processors. To calculate the number of sending and receiving processors, we iterate over full neighbor list of all vertices. Here, we find out the vertices that don't belong to current processor and identify the unique processors, we need to communicate. The number of unique processors is the size of our send-list. The number of unique vertices found from neighbor list of all the vertices, is the size of our receive-list. After receiving the desired information, we update the  $n2c$ ,  $in$  and  $tot$  array entries for the aforementioned vertices and store the neighbor list with weight information for further calculation. Below pseudo-code in Algorithm 2 represents this step. Table 4.1 shows an example simulation for this step. Figure 4.1 visualizes a network of 16 nodes.

#### 4.2.2.4 Community Computation

In this step, we do the actual computation to determine community of each vertex. Each processor,  $P_i$  completes this step individually and locally updates the community of the vertices belonging to it. This step does not require any communication among processors. A random vertex,  $V$  is chosen from the list of vertices. Then, the set of neighboring communities of that vertex is computed with the information from Section 4.2.2.3. Here, the number of links from that vertex,  $V$  to all its neighboring community is computed and stored. Next, the vertex is removed from its current community,  $C_{old}$ . Now, the modularity gain is calculated for all its neighboring communities. If the gain is maximum for community,  $C$ , then vertex,  $V$ 's community is updated to  $C$ . Otherwise,  $V$  gets back to its

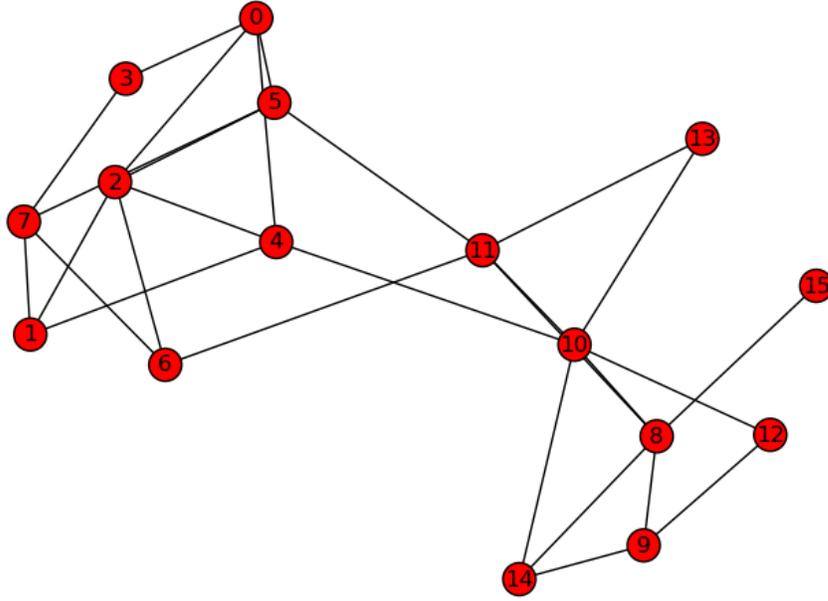


Figure 4.1: Example Graph

previous community,  $C_{old}$ . This, remove () and insert () operations update the in and tot arrays implicitly. As, each processor is locally updating the community, we need to keep trace if community C belongs to processor  $P_i$  or not. If C belongs to processor,  $P_j$ , we simply store  $V_{comm} = C$  and  $V_{link} = d_{V,C}$  where  $d_{V,C}$  = number of links from vertex V to Community C. These two values  $V_{comm}$  and  $V_{link}$  are stored separately in two arrays rem\_comm and rem\_dvc that we used later, described in section 4.2.2.6. This continues until all vertices from the list is covered. The pseudo-code is given in Algorithm 3

Table 4.1: Send and Receive Count

| Processor | Node-Id | Neighbor-List   | Send-List  | Receive-List          |
|-----------|---------|-----------------|------------|-----------------------|
| P0        | 0       | 2,3,4,5         | 1 (P1)     | 4 (4,5,7,6)           |
|           | 1       | 2,4,7           | 1 (P1)     |                       |
|           | 2       | 0,1,4,5,6       | 1 (P1)     |                       |
|           | 3       | 0,7             | 1 (P1)     |                       |
| P1        | 4       | 0,1,2,10        | 2 (P0, P2) | 6(0,1,2,10,11,3)      |
|           | 5       | 0,2,7,11        | 2 (P0, P2) |                       |
|           | 6       | 2,7,11          | 2 (P0, P2) |                       |
|           | 7       | 1,3,5,6         | 1 (P0)     |                       |
| P2        | 8       | 9,10,11,14,15   | 1 (P3)     | 7(14,15,12,4,13, 5,6) |
|           | 9       | 8,12,14         | 1 (P3)     |                       |
|           | 10      | 4,8,11,12,13,14 | 2 (P1, P3) |                       |
|           | 11      | 5,6,8,10,13     | 2 (P1, P3) |                       |
| P3        | 12      | 9,10            | 1 (P2)     | 4(9,10,11, 8)         |
|           | 13      | 10,11           | 1 (P2)     |                       |
|           | 14      | 8,9,10          | 1 (P2)     |                       |
|           | 15      | 8               | 1 (P2)     |                       |
|           |         |                 | 21         | 21                    |

#### 4.2.2.5 Update of Processor List for Communication

This is a preliminary step required for Section 4.2.2.6. After updating the community of the vertices locally, we need to circulate the update globally among all processors so that each community keeps aware of the vertices in its region. Again, we need to figure out the number of sending and receiving processors before starting the communication. It is done in this step. We iterate over all the elements of `rem_comm` array and find out the processor to which that element belonged to. For each processor, we summed the total number of vertices those required update. Again, we also send to all processors of the world a message, `msg` containing either 0 or 1 and receiving back from all. Now, the processors those have at least one vertex that required update, are inserted into our list of sending processors and the value `msg = 1`. Otherwise, `msg = 0`. So, the receive list size is computed by summing the received value of `msg`.

---

**Algorithm 2:** Gather\_Neighbour\_Info()

---

```
1 Initialize empty array receive_list;
2 for each vertex v in g do
3   Initialize empty array send_list;
4   for each neighbor of v do
5      $V_n \leftarrow$  neighboring node of v;
6     if  $V_n$  not in current processor then
7        $P_j \leftarrow$  processor having  $V_n$  ;
8       if receive_list.contains()  $\neq V_n$  then
9         receive_list.push( $V_n$ );
10      end
11      if send_list.contains()  $\neq P_j$  then
12        send_list.push( $P_j$ );
13      end
14    end
15  end
16  for times=send_list.size() do
17     $P_j \leftarrow$  send_list.pop();
18    MPI_Isend(deg(v), weight(v),  $P_j$ );
19    MP_Isend (v, n2c[v], neighbors of v (node-id, weight),  $P_j$ );
20  end
21 end
22 for times= receive_list.size() do
23   MPI_Recv(deg(v'), weight(v'), MPI_ANY_SOURCE);
24    $P_j \leftarrow$  status.MPI_ANY_SOURCE ;
25   MPI_Recv (v', n2c[v'], neighbors of v' (node-id, weight),  $P_j$ );
26   Update n2c[v'], in[v'] and tot[v'] locally;
27   Neighbor[v']. push (neighbors of v' (node-id, weight));
28 end
```

---

$$P_i \xrightarrow{\text{msg}} P_j$$

where,

$$i, j \in \text{world}, j = 0, 1, \dots, N-1 \quad \wedge \quad i \neq j,$$

$$N = \text{world\_size}$$

---

**Algorithm 3:** Compute\_Community()

---

```

1 while vertices.size() != 0 do
2   Pick random vertex, V from g ;
3   for all neighboring vertices of V do
4      $V_n \leftarrow$  neighboring vertex of V ;
5      $d_{V,V_n} \leftarrow$  number of links from vertex, V to community,  $V_n$ ;
6   end
7   /* remove node from its current community */
8   remove (V, n2c[V],  $d_{V,V_n}$ );
9   for all neighboring vertices of V do
10     $V_n \leftarrow$  neighboring vertex of V ;
11    compute_gain() ;
12    if positive gain then
13       $n2c[v] \leftarrow n2c[V_n]$  ;
14      if  $n2c[V_n]$  not in current processor then
15         $P_j \leftarrow$  processor having  $n2c[V_n]$ ;
16         $rem\_comm[V_n] \leftarrow n2c[V_n]$ ;
17         $rem\_dvc[V_n] \leftarrow d_{V,V_n}$ ;
18      end
19      /* insert locally */
20      Insert (V,  $n2c[V_n]$ ,  $d_{V,V_n}$ );
21    end
22  end

```

---

$$msg = \begin{cases} 1, & \text{verticesrequiringupdate} \geq 1 \\ 0, & \text{verticesrequiringupdate} = 0 \end{cases}$$

Algorithm 4 represents the pseudo-code.

#### 4.2.2.6 Exchange of Updated Community

In this step, the update of community is done globally among all processors. Now, Processor,  $P_i$  sends the sending processor,  $P_j$  the following data until the send-list becomes empty.

- Vertex,  $v$  belonging to Processor  $P_i$ , whose community is in Processor  $P_j$

---

**Algorithm 4:** Update\_Send\_Receive\_List()

---

```
1 for times = world_size do
2   if  $P_i \neq rank$  then
3     if  $P_i \hat{=} \checkmark$ s vertices requiring update > 0 then
4       MPI_Send (1,  $P_i$ );
5       send_list.push( $P_i$ );
6     else
7       MPI_Send (0,  $P_i$ );
8     end
9     MPI_Recv (j,  $P_i$ );
10    rcv_list_size+ = j;
11  end
12 end
```

---

- Vertex,  $v$ 's community,  $C$  belonging to Processor  $P_j$
- Number of links from vertex,  $v$  to community  $C$
- Number of self-loops of vertex,  $v$
- Weighted degree of vertex,  $v$

$P_i \xrightarrow{msg} P_j$  where,  $j \in send\_list$

$msg \ni Vertex, Vertex_{community}, Vertex_{link},$

$Vertex_{self-loop}, Vertex_{weighted-degree}$

$Vertex \in P_i, Vertex_{community} \in P_j$

Upon receiving the data, the vertex is inserted to its community and it updates in and tot arrays internally. Pseudo-code for this step is given in Algorithm 5.

---

**Algorithm 5:** Exchange\_Updated\_Community()

---

```
1 for times = send_list.size() do
2   | MPI_Send (#nodes, Pj);
3   | MPI_Send (v, n2c[Vn], dnodecomm, vself-loop, vweighted-degree, Pj);
4 end
5 for times = rcv_list_size do
6   | MPI_Recv (# nodes, MPI_ANY_SOURCE);
7   | Pi status.MPI_ANY_SOURCE;
8   | MPI_IRecv (x, y, dnodecommxy, xself-loop, xweighted-degree, Pi);
9   | /* insert x in community y global update, only update in[],
      |    tot[] of y */
10  | Insert (x, y, dnodecommxy, xself-loop, xweighted-degree);
11 end
```

---

#### 4.2.2.7 Resolving Community Duality

After the previous step, there remains an inconsistency to calculate total number of communities in the full network. The same network shown in Figure 4.1 is used to demonstrate the problem.

Table 4.2: Community Duality

| Processor | Vertex | Community | Problem  | Solution   |
|-----------|--------|-----------|--|--|
| P0        | 1      | 4         | Vertex, 1<br>switched its<br>community to<br>vertex, 4 | Vertex 1<br>retained<br>community 4                        |
| P1        | 4      | 1         | Vertex, 4<br>switched its<br>community to<br>vertex, 1 | Vertex 4's<br>community<br>switched<br>back to<br>vertex 4 |

Basically, Vertex 1 and Vertex 4 belong to the same community. It can be either 1 or 4. The community number does not have any effect on the result. But in current scenario, same community will be counted twice. So, we need to eliminate this problem. To solve this, we kept the communities with higher number

and changed the lower numbers to higher ones. So now, in Table 4.2, vertex 4's community is changed again to community 4 and vertex 1 retains its community 4. As, vertex 4 belongs to Processor 1. Processor 0 needs to communicate with it to circulate the update. In this step, we update the vertices' communities those belong to current processor and track the communities those belonged to other processors and communication is required. So, for other processors those contain communities of the vertices, we summed the total number where current community id is less than vertex id and stored it in an array, `count_community` and it will be used in Step-8. We stored the vertices whose community will be updated after the communication. We also enlisted the unique community ids whose updated community we will receive after the communication is done. Both of these are required for the step described in Section 4.2.2.9.

#### **4.2.2.8 Update of Processor List for Communication in Resolving Community Duality**

This is a preliminary step required for Section 4.2.2.9. It is similar as described in Section 4.2.2.5. Here, we iterate over the array, `count_community` and stored the indices,  $i$  in `send_list` if `count_community[i] > 0`.  $i$  represents the processor number for sending. Update of receive list is also analogous to the calculation as described in Section 4.2.2.5 for finding out the size of `receive_list`.

#### **4.2.2.9 Exchange of Duality Resolved Community**

In this step, processor,  $P_i$  send to all processors,  $P_j$  from its `send-list` the community-id to get the updated community of that id.  $P_j$  after receiving the message, sends back to  $P_i$  the corresponding community of received id. After receiving info from  $P_j$ ,  $P_i$  updates the community of the vertices those needed up-

date, using the values from Section 4.2.2.7. In this communication step, subsequent send-receive is done.

$$P_i \xrightarrow{msg1} P_j \quad \text{where, } j \in send\_list$$

$$msg1 \ni Community; Community \in P_j$$

$$P_j \xrightarrow{msg2} P_i$$

$$msg2 \ni Community, n2c[community]; Community \in P_j$$

A brief pseudo-code to represent Section 4.2.2.7, 4.2.2.9 is given in Algorithm 6.

---

**Algorithm 6:** Resolve\_Community\_Duality() and Exchange\_Duality\_Resolved\_Community()

---

```

1 for each vertex, v do
2   if n2c[v] < v then
3     if n2c[v] not in Pi then
4       Pj ← processor having n2c[v];
5       MPI_Send(n2c[v], Pj);
6       MPI_Recv (n2c[v], n2c[n2c[v]], Pj);
7       n2c[v] ← n2c[n2c[v]];
8     else
9       comm ← n2c[v];
10      n2c[v] ← n2c[comm];
11    end
12  end
13 end

```

---

#### 4.2.2.10 Finding Unique Communities and Computation of Modularity

In this step, each processor, P<sub>i</sub> , where 0 ≤ i < N finds out all the unique communities by iterating over all of its vertices belonging to it. Pseudo-code is presented in Algorithm 7.

---

**Algorithm 7:** Find\_Unique\_Communities()

---

```
1 Initialize empty array unique_community;
2 for each vertex, v do
3   comm ← n2c[v];
4   if unique_community.contains()! = comm then
5     unique_community.push(comm);
6   end
7 end
```

---

To calculate total unique communities in the world, each Processor,  $P_k$ , where  $(1 \leq k < N)$  sends its unique\_community list to root processor,  $P_0$ .  $P_0$  then merges all the unique communities received from each processor,  $P_k$  and eliminates duplicate ones.

Along with the unique\_community list,  $P_k$  also sends to  $P_0$  values  $in1$  and  $tot1$  calculating the values from  $in$  and  $tot$  arrays. These two values are required for calculation of modularity of the full network. Algorithm 8 represents the pseudo-code for this step.

---

**Algorithm 8:** Compute\_ Modularity()

---

```
1 if rank! = 0 then
2   i ← rank;
3   m ← total_weight;
4   in1 ←  $\frac{(\sum in[node])}{m}$ ;
5   tot1 ←  $(\frac{\sum tot[node]}{m})^2$ ;
6   MPI_Send(community array, in1i, tot1i, 0);
7 else
8   MPI_Recv (community array, in1i, tot1i, MPI_ANY_SOURCE);
9   Modularity ←  $\sum in1_1 - \sum tot1_i$ ;
10 end
```

---

#### 4.2.2.11 Generating Next Level Graph

This step is performed by only the root processor,  $P_0$ .  $P_0$  renumbers the communities from 0 to  $Z - 1$  to formulate the new input graph to be used for next level.

$Z =$  Total number of unique communities after merging and eliminating duplicate

So,  $Z$  is the number of vertices for input graph of next level. The connectivity of edges and corresponding weights are calculated from available data and the new graph is generated.

All the steps from Section 4.2.2.1 to 4.2.2.11 are done for each level of computation and repeated as long as we get increase in modularity value.

### 4.3 Hybrid Parallel Louvain Algorithm

We use both MPI and OpenMP together to implement the Hybrid Parallel Louvain Algorithm. The hybrid version gives us the flexibility to balance between both shared and distributed memory system. We can tune between shared and distributed memory depending on available resources. In the multi-threading environment, a single thread works for communication among processors and other threads do the computation.

### 4.4 Distributed Parallel Louvain Algorithm with Load-balancing

To implement DPLAL, we use the similar approach as described in Section 4.2. In the first phase, we have used well-known graph-partitioner METIS [28] to par-

---

**Algorithm 9:** DPLAL-Distributed Parallel Louvain Algorithm using Load-balancing

---

**Data:** Input Graph  $G(V,E)$  [Edge List Format]  
**Result:** (Vertex, Community) Pair

```

1 while increase in modularity do
2    $G(V, E)$  is partitioned using METIS into  $p'$  partitions;
3    $G'(V,E)$  pre-processed according to METIS-Output;
4    $G''(V,E)$  converted to Adjacency List format to be given to each
   processor ;
5   for Each processor  $P_i$  (executing in parallel) do
6     Gather_Neighbour_Info();
7     Compute_Community();
8     Exchange_Updated_Community();
9     Resolve_Community_Duality();
10    Exchange_Duality_Resolved_Community();
11    Find_Unique_Communities();
12    Compute_Modularity();
13    Generate_NextLevel_Graph();
14    if number_of_communities <  $i$  then
15      |  $i \leftarrow \frac{\text{number\_of\_communities}}{2}$ ;
16    end
17  end
18 end

```

---

tition our input graph to distribute among the processors. Depending on METIS output, we adjust the number of processors because METIS does not always create same number of partitions as provided in input. We use both edge-cut and communication volume minimization approaches. An empirical comparison of these approaches is described later in Section 6. After partitioning, we distribute the input graph among the processors. For second phase, we follow the same flow as described in the Algorithm 1. But we have to recompute each function that has been calculated from input graph. Runtime analysis for each of these functions being used in MPI communication has been demonstrated in Section 6. Our incorporation of graph partitioning scheme helps minimize the communication overhead of MPI to a great extent and we get an optimized performance from DPLAL.

# Chapter 5

## Experimental Setup

We describe our experimental setup and datasets below. We use large-scale compute cluster for working on large real-world graph datasets.

### 5.1 Execution Environment

We use Louisiana Optical Network Infrastructure (LONI) QB2 [29] compute cluster to perform all the experiments. QB2 is a 1.5 Petaflop peak performance cluster containing 504 compute nodes with over 10,000 Intel Xeon processing cores of 2.8 GHz. We use at most 50 computing nodes with 1000 processors for our experiments.

### 5.2 Description of Datasets

We have used real-world networks from SNAP [30] depicted in Table 5.1. We have performed our experimentation on different types of network including social networks, internet, peer-to-peer networks, road networks, network with ground truth communities, and Wikipedia networks. All these networks show different

structural and organizational properties. This gives us an opportunity to assess the performance of our algorithms for worst case inputs as well. The size of graphs used in our experiments ranges from several hundred thousands to millions of edges.

Table 5.1: Datasets used in our experimental evaluation.

| <b>Network</b>   | <b>Vertices</b> | <b>Edges</b> | <b>Description</b>   |
|------------------|-----------------|--------------|--|
| email-Eu-core    | 1,005           | 25,571       | Email network from a large European research institution   |
| ego-Facebook     | 4,039           | 88,234       | Social circles ('friends lists') from Facebook   |
| wiki-Vote        | 7,115           | 103,689      | Wikipedia who-votes-on-whom network  |
| p2p-Gnutella08   | 6,301           | 20,777       | A sequence of snapshots of the Gnutella peer-to-peer file sharing network for different dates of August 2002 |
| p2p-Gnutella09   | 8,114           | 26,013       |  |
| p2p-Gnutella04   | 10,876          | 39,994       |  |
| p2p-Gnutella25   | 22,687          | 54,705       |  |
| p2p-Gnutella30   | 36,682          | 88,328       |  |
| p2p-Gnutella31   | 62,586          | 147,892      |  |
| soc-Slashdot0922 | 82,168          | 948,464      | Slashdot social network from February 2009   |
| com-DBLP         | 317,080         | 1,049,866    | DBLP collaboration (co-authorship) network   |
| roadNet-PA       | 1,088,092       | 1,541,898    | Pennsylvania road network  |

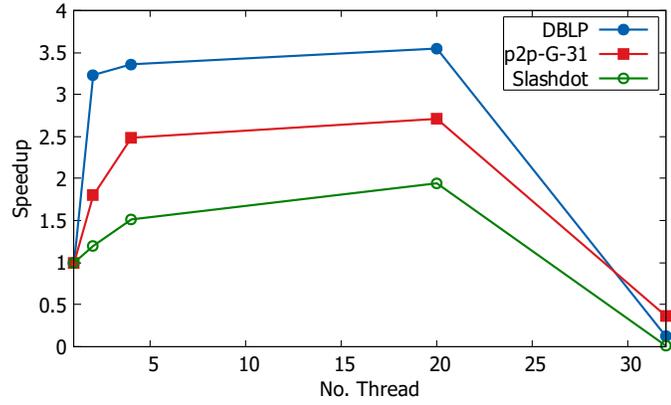
# Chapter 6

## Results

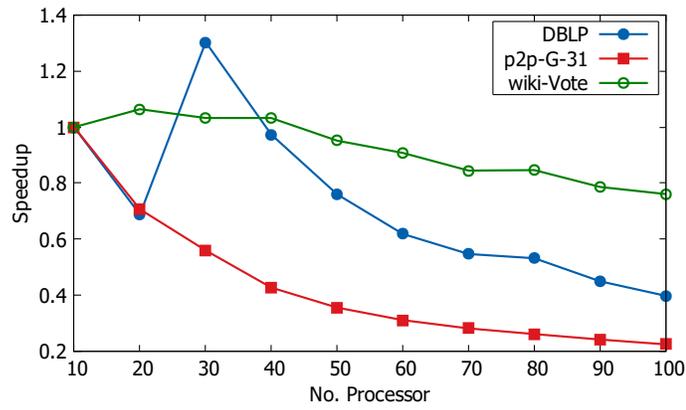
We present the scalability and runtime analysis of our algorithms below. We discuss the trade-offs and challenges alongside.

### **6.1 Speedup factors of shared and distributed memory algorithms**

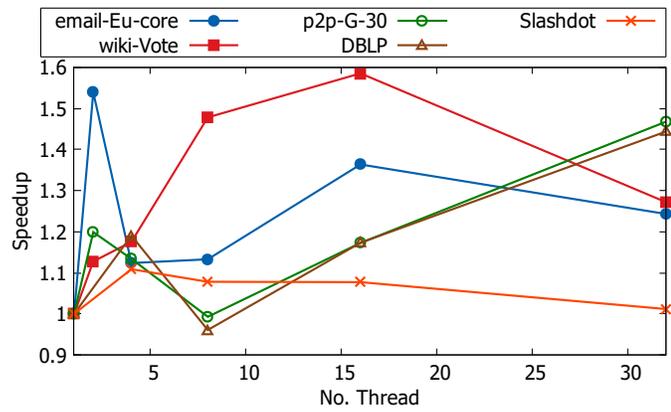
We design both shared and distributed memory based algorithms for Louvain methods. The speedup results are shown in Fig. 6.1a and fig. 6.1b. Our shared memory and distributed memory based algorithms achieve speedups of around 4 and 1.5, respectively. The number of physical processing core available to our system is 20. Our shared memory algorithm scales well to this many cores. However, due to the unavailability of large shared memory system, we also design distributed memory algorithm. Further, shared memory algorithms show a limited scalability to large networks as discussed in [16]. Our distributed memory algorithm demonstrates only a minimal speedup for 30 processors . The inter-processor communication severely affects the speedup of this algorithm. We strive to overcome such com-



(a) Shared Memory Algorithm



(b) Distributed Memory Algorithm



(c) Hybrid Algorithm

Figure 6.1: Speedup factors of our parallel Louvain algorithms for different types of networks. Our hybrid algorithm strikes a balance between shared and distributed memory based algorithms.

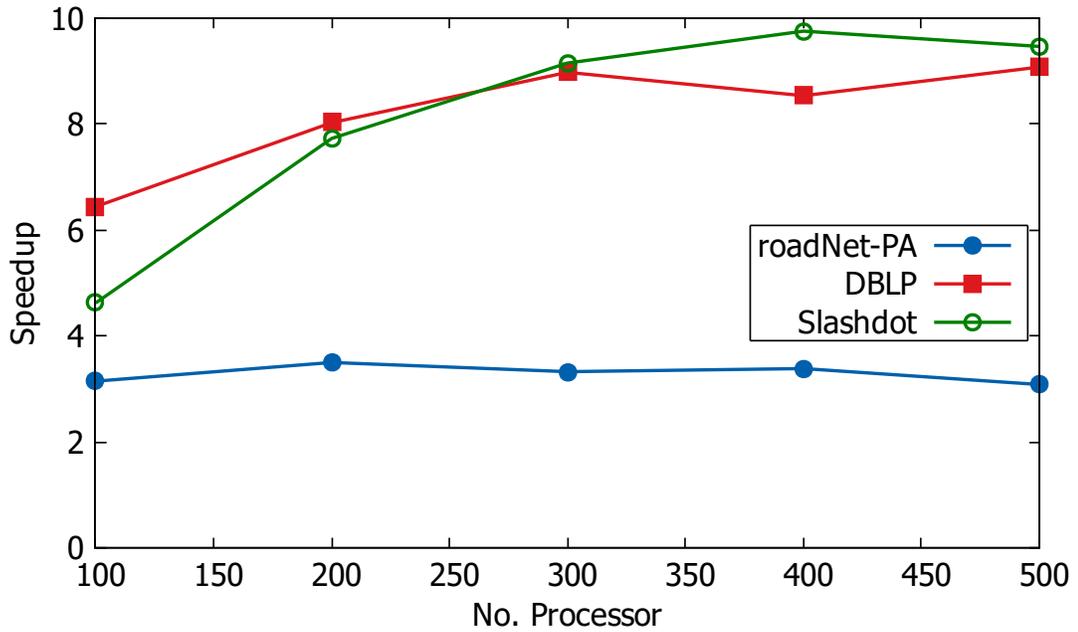
munication bottleneck by designing hybrid algorithm.

## **6.2 Speedup factors of our hybrid parallel algorithm**

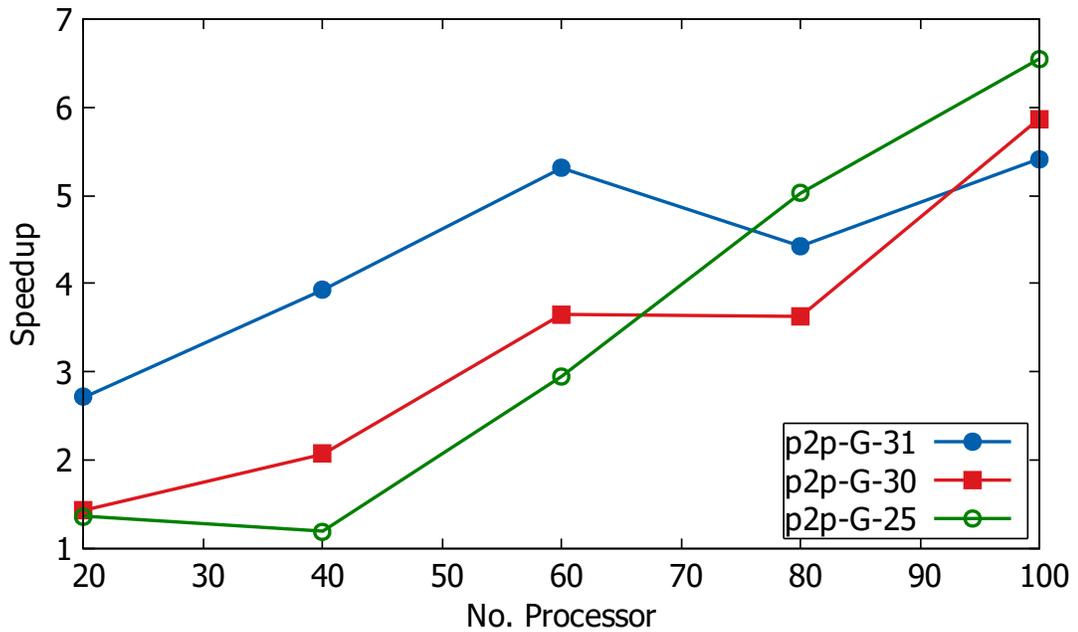
Our hybrid algorithm tends to find a balance between the above two approaches, shared and distributed memory. As shown in Fig. 6.1c, we get a speedup of around 2 for the hybrid implementation of Louvain algorithm. The speedup is similar to the MPI implementation. It is evident that in multi-threading environment runtime will decrease as workload is distributed among the threads. But we observe that in some cases, both single and multiple threads take similar time. Even sometimes multiple threads take more time than a single thread. It indicates that hybrid implementation also suffers from the communication overhead problem alike MPI. Communication overhead of distributed memory setting limits the performance of hybrid algorithm as well.

## **6.3 Speedup factors of our improved parallel algorithm DPLAL**

Our final parallel implementation of Louvain algorithm is DPLAL. This algorithm achieves a speedup factor up-to 12. We reduce the communication overhead in message passing setting to a great extent by introducing a load balancing scheme during graph partitioning. The improved speedup for DPLAL is presented in Figure 6.2. For larger networks, our algorithm scales to a larger number of processors. We are able to use around a thousand processors. For smaller networks, the algorithm scales to a couple of hundred processors. It is understandable that for



(a) Speedup results for large graphs



(b) Speedup results for relatively small graphs

Figure 6.2: Speedup factors of DPLAL algorithm for different types of networks. Larger networks scale to a larger number of processors.

smaller networks, the communication overhead gradually offsets the advantage obtained from parallel computation. However, since we want to use a larger number of processors to work on larger networks, our algorithm in fact has this desirable property. Overall, DPLAL algorithm scales well with the increase in the number of processors and to large networks.

## 6.4 Runtime analysis: a breakdown of execution times

We present a breakdown of executions times. Fig. 6.3 shows the runtime analysis for our largest network RoadNet-PA. We observe that communication time for *gathering neighbor information* and *exchanging duality resolved community* decreases with increasing number of processors. Communication time for both *exchanging updated community* and *gathering updated community* increases up-to a certain number of processors and after decreasing, the time becomes almost constant. Among all these communications, time to gather communities at the root processor takes maximum time and contribute to the high runtime.

## 6.5 Network size versus execution time.

For many large networks that we experimented on (including the ones in Fig. 6.2a), we find that those can scale to up to  $\approx 800$  processors. We call this number as the *optimum number of processors* for those networks. This optimum number depends on network size. As our focus is on larger networks, to find out the relationship between runtime and network size, we keep the number of processor 800 fixed and run an experiment. As shown in Fig. 6.4, the communication

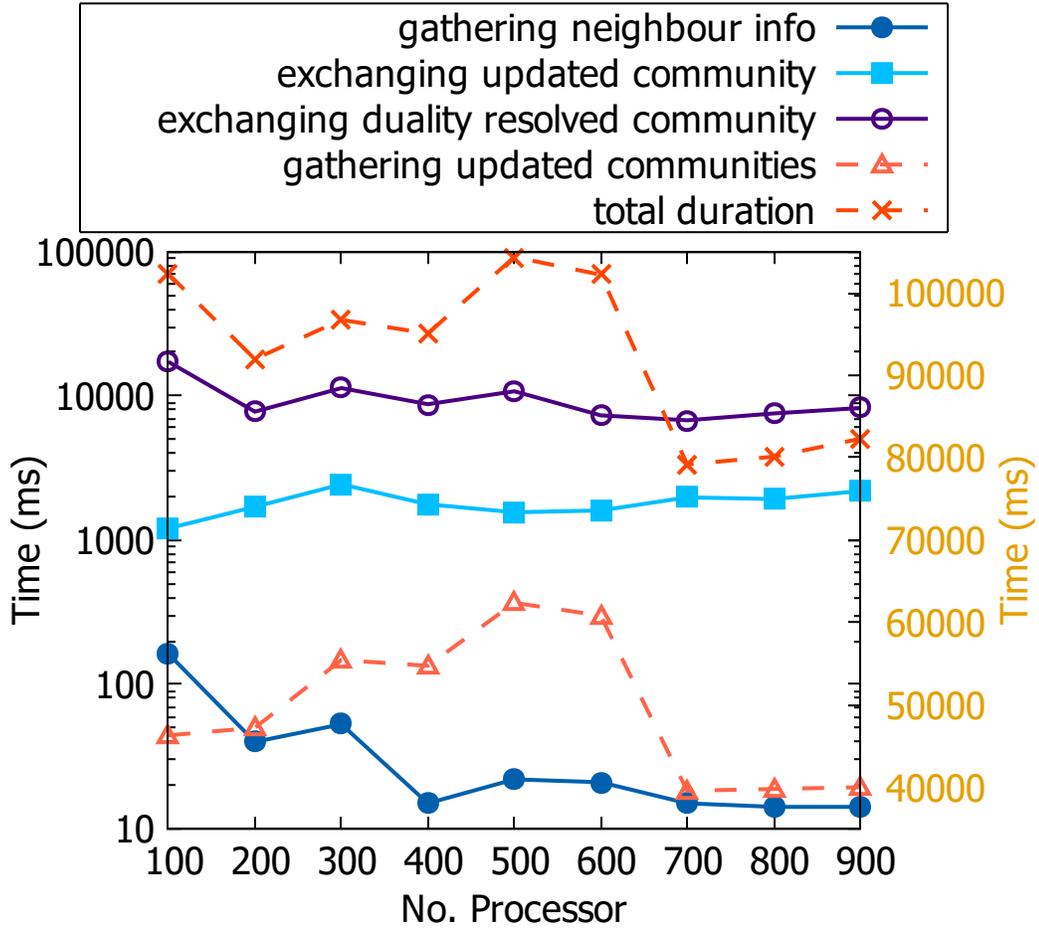


Figure 6.3: Runtime analysis of RoadNet-PA graph with DPLAL algorithm for varied number of processors. We show a breakdown of execution times for different modules or functions in the algorithm. Time for *gathering updated communities* and *total duration* are plotted w.r.t the right  $y$ -axis.

time for *gathering neighbor info* decreases with growing network size whereas both time for *gathering updated communities* and *exchanging duality resolved community* increase. Communication time for *exchanging updated community* increases up-to a certain point and then starts decreasing afterwards. For larger networks ( $> 80K$ ), total runtime increases proportionately with growing network size. As smaller graphs do not scale to 800 processors, these do not follow the trend, but it can be inferred that these will behave the same way for their optimum number of

processors.

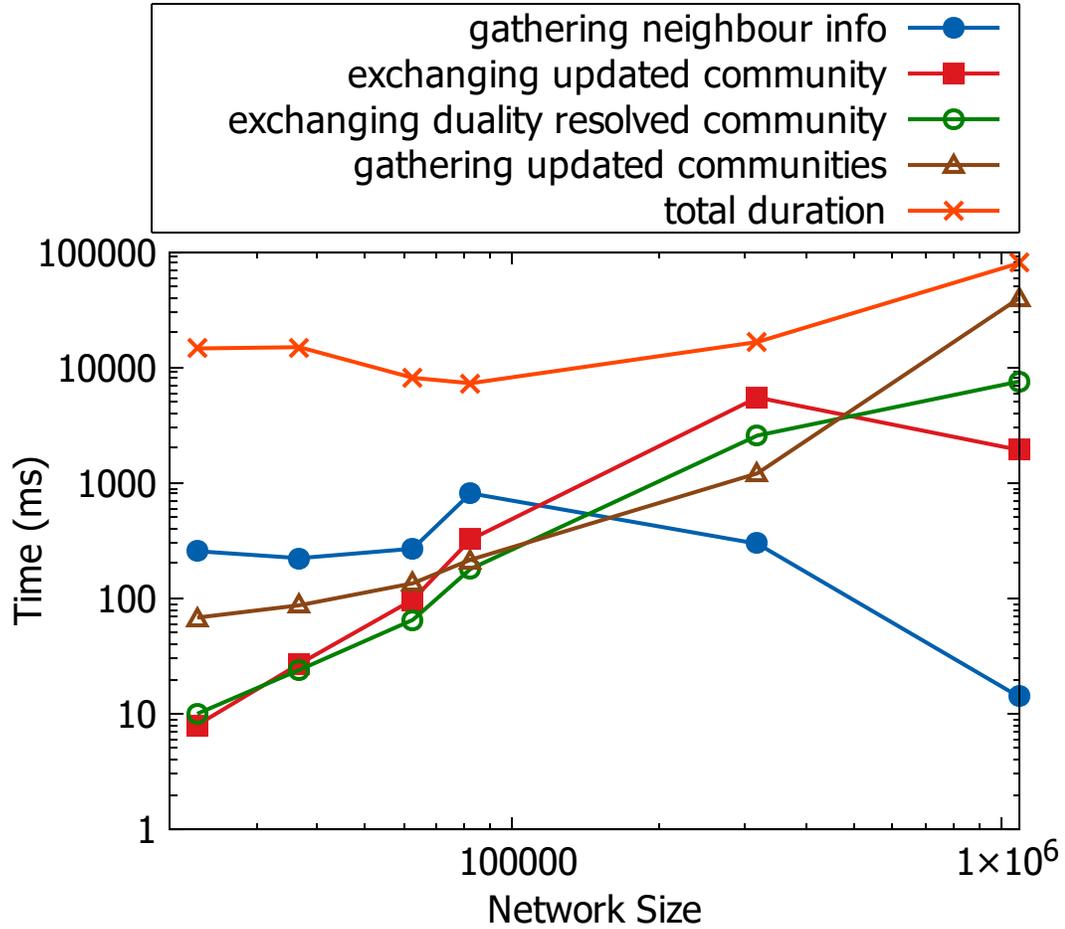


Figure 6.4: Increase in runtime of DPLAL algorithm with an increase in the sizes of the graphs keeping the number of processors fixed.

## 6.6 METIS partitioning approaches

We also compare the METIS partitioning techniques, between edge-cut and communication volume minimization, to find out the efficient approach for our algorithm. Fig. 6.5 shows the runtime comparison between edge-cut and communication volume minimization techniques. We find that the communication volume minimization approach always takes similar or higher time than that of edge-cut

partitioning. So, in our subsequent experimentation, we have used edge-cut partitioning approach.

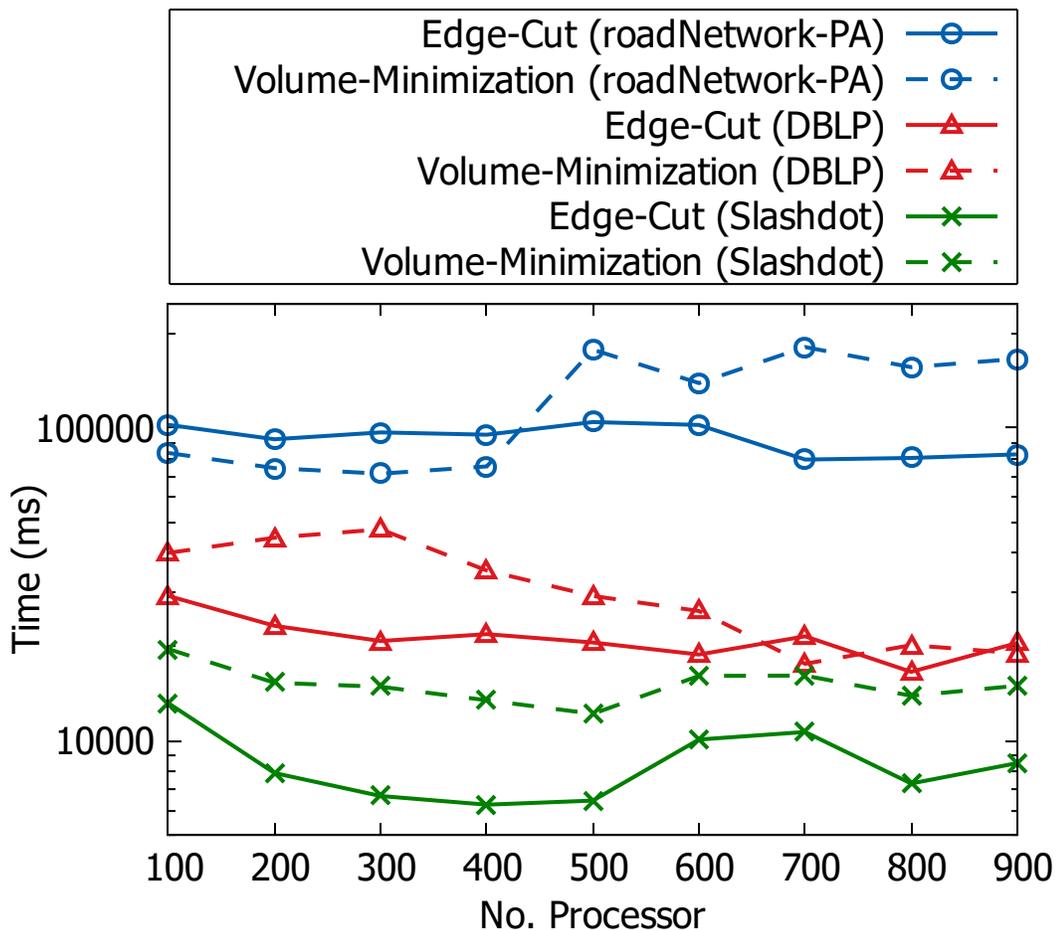


Figure 6.5: Comparison of METIS partitioning approaches (edge-cut versus communication volume minimization) for several networks. The edge-cut approach achieves better runtime efficiency for the above real-world networks.

## 6.7 Performance Analysis

We present a comparative analysis of our algorithms, its sequential version, and another existing distributed memory algorithm.

### 6.7.1 Comparison with Other Parallel Algorithms

We compare the performance of DPLAL with another distributed memory parallel implementation of Louvain method given in Charith et al. [1]. For a network with 500,000 nodes, Charith et al. achieved a maximum speedup of 6 whereas with DPLAL for a network with 317,080 nodes we get a speedup of 12 using 800 processors. The largest network processed by them has 8M nodes and achieved a speedup of 4. Our largest network achieves a comparable speedup (4-fold speedup with 1M nodes). The work in [1] did not report runtime results so we could not compare our runtime with theirs directly. Their work reported scalability to only 16 processors whereas our algorithm is able to scale to almost a thousand of processors.

### 6.7.2 Comparison with Sequential Algorithm

We have compared our algorithms with the sequential version [15] to analyze the accuracy of our implementations. Deviation of the number of communities between sequential and our implementations is represented in Table 6.1. The deviation is negligible compared to network size. The number of communities is not constant and they vary because of the randomization introduced in the Louvain algorithm. Table 6.1 gives an approximation of the communities.

Although shared memory based parallel Louvain has the least deviation, the speedup is not remarkable. Whereas, DPLAL shows a moderate deviation but its speedup is 3 times of that of shared parallel Louvain algorithm.

Table 6.1: Deviation of the number of communities for different parallel Louvain Algorithms from the sequential algorithm.

| Algorithm   | Network   |          |           |          |
|-------------|-----------|----------|-----------|----------|
|             | com-DBLP  |          | wiki-Vote |          |
|             | Comm. No. | Dev. (%) | Comm. No. | Dev. (%) |
| Sequential  | 109,104   | -        | 1,213     | -        |
| Shared      | 109,102   | .0006    | 1,213     | 0        |
| Distributed | 109,441   | 0.106    | 1,216     | 0.042    |
| Hybrid      | 104,668   | 1.39     | 1,163     | 0.71     |
| DPLAL       | 109,063   | 0.0129   | 1,210     | 0.042    |

# Chapter 7

## Conclusion

Our parallel algorithms for Louvain method demonstrate good speedup on several types of real-world graphs. As instance, for DBLP graph with 0.3 million nodes, we get speedups of around 4, 1.5 and 2 for shared memory, distributed memory, and hybrid implementations, respectively. Among these three algorithms, shared memory parallel algorithm gives better speedup than others. However, shared memory system has limited number of physical cores and might not be able to process very large networks. A large network often requires distributed processing and each computing node stores and works with a part of the entire network. As we plan to work with networks with billions of nodes and edges, we work towards the improvement of the scalability of our algorithms by reducing the communication overhead. We have identified the problems for each implementation and come up with an optimized implementation **DPLAL**. With our improved algorithm DPLAL, community detection in DBLP network achieves a 12-fold speedup. Our largest network, roadNetwork-PA has 4-fold speedup for same number of processors. With increasing network size, number of processor also increases. We will work with larger networks increasing the number of processors in our future work. The optimum number of processor largely depends on the network size. We will

also experiment with other load-balancing schemes to find an efficient load balancing scheme to make DPLAL more scalable. We also want to eliminate the effect of small communities that create misconception to understand the community structure and its properties. Further, we will explore the effect of node ordering (e.g., degree based ordering, random ordering) on the performance of parallel Louvain algorithms.

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## Vita

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