

# Linear Algebraic Louvain Method in Python

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**Abstract**—We show that a linear algebraic formulation of the Louvain method for community detection can be derived systematically from the linear algebraic definition of modularity. Using the pygraphblas interface, a high-level Python wrapper for the GraphBLAS C Application Programming Interface (API), we demonstrate that the linear algebraic formulation of the Louvain method can be rapidly implemented.

**Index Terms**—Louvain Method, Community Detection, Graph Algorithms, GraphBLAS

## I. INTRODUCTION

Community detection is a critical component in the analysis of real-world network systems with important applications in social media analytics, biology, recommendation systems, and telecommunications among others [1]. The Louvain method [2] based on modularity optimization [3] is a commonly adopted technique to identify communities.

Current implementations of the Louvain method adopt a traditional vertex-based approach in their algorithmic formulation despite there being a linear algebraic formulation for the computation of modularity [4]. In this paper, we show that by starting with the linear algebraic formulation of modularity, a linear algebraic algorithm for computing the Louvain method can be determined. This formulation can leverage recent community-driven efforts behind linear algebraic graph interfaces, such as the GraphBLAS [5], to rapidly develop an implementation of the Louvain method. Productivity is enhanced through the use of pygraphblas [6], a python interface for the GraphBLAS effort.

## II. BACKGROUND

We assume the goal is to identify communities within a graph  $\mathcal{G} = (V, E)$  where  $V$  is the set of vertices and  $E$  is the set of weighted edges connecting them. The Louvain method is an iterative algorithm where each iteration is composed of two major steps:

- *Step 1)* Vertices are moved between communities such that each move increases the overall modularity of  $\mathcal{G}$ . Step 1 is initialized by creating  $|V|$  communities where each vertex is in its own community. Vertices are sequentially reassigned to new communities as long as modularity can be increased. The choice of communities for a vertex is based on the community that yields the highest modularity increase.

- *Step 2)* A new graph is created from the newly partitioned graph. Identified communities form the set of new vertices, and edges between vertices in different communities are aggregated into a new set of edges. Step 1 is then applied to this newly created graph.

The two steps above are repeated until no further improvement in modularity is obtained.

### A. Definition of Modularity

Modularity is a scalar measure used to quantify the strength of community structures within a graph [3]. It is computed as

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

where  $A_{ij}$  is the edge weight between vertices  $v_i, v_j \in V$ ,  $m$  is the sum of all edge weights,  $k_i$  is the sum of the weights of all edges incident on vertex  $v_i$ ,  $c_i \in C$  is the community to which  $v_i$  is currently assigned, and  $\delta(c_i, c_j) = 1$  if vertices  $v_i$  and  $v_j$  are in the same community (i.e.,  $c_i = c_j$ ) and 0 otherwise.

We will build on the matrix formulation of  $Q$  from [4]:

$$Q = \frac{1}{2m} \Gamma(S^T B S), \quad (1)$$

where  $\Gamma(\cdot)$  computes the trace of a matrix,  $S$  is a  $|V| \times |C|$  matrix that captures the community membership of each vertex, and  $B$  is the  $|V| \times |V|$  modularity matrix where element  $B_{ij}$  is defined as

$$B_{ij} = A_{ij} - \frac{k_i k_j}{2m}.$$

Each column  $s_i$  of  $S$  represents a community of vertices, where a one in the  $j^{\text{th}}$  position of  $s_i$  indicates that vertex  $v_j$  is in community  $c_i$ . We denote vertex  $v_j$  as a column basis vector  $e_j$  of length  $|V|$ , and  $s_i$  is the sum of all vertices that belong to the same community  $c_i$ , i.e.,

$$s_i = \sum_{j: v_j \in c_i} e_j.$$

An empty community is simply the zero vector, i.e.  $s_i = \vec{0}$ .

## III. LINEAR ALGEBRAIC LOUVAIN METHOD

In this section we develop a fully linear algebraic formulation of the Louvain algorithm starting from the linear algebraic definition of modularity.

### A. Modularity of a Community

Starting with Eqn. (1) and expanding on the definition of the trace of a matrix, we express the modularity  $Q$  as

$$\begin{aligned} Q &= \frac{1}{2m} \left( s_0^T B s_0 + s_1^T B s_1 + \dots + s_{|C|-1}^T B s_{|C|-1} \right) \\ &= \frac{1}{2m} \sum_{i=0}^{|C|-1} s_i^T B s_i = \frac{1}{2m} \sum_{i=0}^{|C|-1} Q_i, \end{aligned}$$

where  $Q_i$  is the modularity of community  $c_i$ .

Consider an arbitrary vertex  $v_j$  where  $v_j$  is a vertex (possibly, the only one) in community  $c_i$ . We can define a community vector  $s_i$  in terms of the vertex  $v_j$  and the remaining set of vertices  $s_{i \setminus \{j\}}$  as follows

$$s_i = s_{i \setminus \{j\}} + e_j.$$

Using the above definition, we can derive the following definition of modularity for community  $i$  in terms of the contribution to modularity by introducing vertex  $v_j$  into community  $i$ :

$$\begin{aligned} Q_i &= s_i^T B s_i \\ &= (e_j + (s_i - e_j))^T B (e_j + (s_i - e_j)) \\ &= e_j^T B e_j + s_{i \setminus \{j\}}^T B e_j + e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B s_{i \setminus \{j\}} \\ &= \underbrace{B_{jj} + s_{i \setminus \{j\}}^T B e_j + e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B s_{i \setminus \{j\}}}_{\text{Contribution by vertex } v_j} \end{aligned}$$

### B. Linear Algebraic Louvain Method

We address the two steps of Louvain separately.

- Step 1) Initially, vertices are assigned to their own community and the community matrix  $S$  is the identity matrix,  $I$ .

Each vertex,  $v_j$  is evaluated for the change in modularity when reassigned from community  $c_i$  to community  $c_\ell$ . Mathematically, the change in modularity due to such movement ( $\Delta Q_{i \rightarrow \ell}$ , where  $i \neq \ell$ ) can be computed as:

$$\begin{aligned} \Delta Q_{i \rightarrow \ell} &= (B_{jj} + e_j^T B s_{\ell \setminus \{j\}} + s_{\ell \setminus \{j\}}^T B e_j) \\ &\quad - (B_{jj} + e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B e_j) \\ &= e_j^T B s_{\ell \setminus \{j\}} + s_{\ell \setminus \{j\}}^T B e_j \\ &\quad - (e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B e_j). \end{aligned}$$

Essentially,  $\Delta Q_{i \rightarrow \ell}$  is computed by subtracting the contribution of vertex  $v_j$  to community  $c_i$  from the contribution of vertex  $v_j$  to community  $c_\ell$ .

Let  $q$  be a vector of length  $|V|$  defined as follows:

$$q^T = \left( \Delta Q_{i \rightarrow 0} \mid \dots \mid \Delta Q_{i \rightarrow |V|-1} \right).$$

As the Louvain algorithm only considers moving a vertex to a community that includes at least one neighbor, we can restrict the modularity computation to the communities containing neighbors of  $v_j$  via a mask in the following manner:

$$q_1^T = q^T \circ \underbrace{((e_j^T (A + A^T)))}_{t_q} S \neq 0. \quad (2)$$

The maximum increase in modularity is computed as  $\kappa = \max(q_1)$ , and the community  $c_\ell$  associated with  $\kappa$  can be identified by

$$t = (q_1 == \kappa),$$

where  $t$  is a boolean vector of length  $|V|$  indicating the new community for vertex  $v_j$ . The original community  $c_i$  for vertex  $v_j$  can be computed by extracting the  $j^{\text{th}}$  row of  $S$ . Using the original community and the vector  $t$ , the community matrix  $S$  can be updated as

$$S = S + e_j t^T - e_j e_j^T S.$$

Step 1 is repeated for all vertices until no increase in modularity can be achieved.

- Step 2) A new graph  $\mathcal{G}' = (V', E')$  is constructed, where the set of vertices  $V'$  are the communities identified in Step 1. The new adjacency matrix  $A'$  is computed as

$$A' = S^T A S,$$

where the weight at location  $A'_{ij}$  is the sum of edge weights in  $A$  between vertices in community  $c_i$  and vertices in community  $c_j$ . The diagonal elements  $A'_{ii}$  are the sum of edge weights in  $A$  belonging to the same communities  $c_i$ .

## IV. IMPLEMENTATION CONSIDERATIONS

In this section, we focus on Step 1 of the previously presented algorithm and discuss several optimizations.

- 1) Removing redundant computation. Recall the objective in Step 1 is to find the maximum increase in modularity. Since the expression  $e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B e_j$  is removed from all modularity computation, it does not contribute to the decision of which community should vertex  $v_j$  be reassigned to. Thus, the expression  $e_j^T B s_{i \setminus \{j\}} + s_{i \setminus \{j\}}^T B e_j$  does not need to be computed. Hence, the following computation can be performed instead:

$$\begin{aligned} \Delta \tilde{Q}_{i \rightarrow \ell} &= e_j^T B s_{\ell \setminus \{j\}} + s_{\ell \setminus \{j\}}^T B e_j. \\ &= e_j^T (B + B^T) s_{\ell \setminus \{j\}} \end{aligned} \quad (3)$$

- 2) Bulk computation of  $q$ . Notice that Eqn. (3) can be computed at once for all values of  $\ell$  as:

$$q^T = e_j^T (B + B^T) \bar{S}.$$

where

$$\begin{aligned} \bar{S} &= S - e_j e_j^T S \\ &= (I - e_j e_j^T) S, \end{aligned} \quad (4)$$

which is the community matrix after removing vertex  $v_j$  from its original community.

- 3) Eliminating modularity matrix  $B$ . Modularity matrix  $B$  is a dense  $|V| \times |V|$  matrix. In order to reduce the memory requirements, recall that

$$B = A - \frac{1}{2m} k k^T,$$

where  $k$  is a vector where the  $i^{th}$  value is the node degree of vertex  $v_i$ . Therefore,

$$\begin{aligned} B + B^T &= \left(A - \frac{1}{2m}kk^T\right) + \left(A - \frac{1}{2m}kk^T\right)^T \\ &= (A + A^T) - \frac{1}{m}kk^T. \end{aligned} \quad (5)$$

Instead of storing  $B$ , the above formulation allows us to use a sparse matrix format to store  $G = (A + A^T)$ , and a dense vector  $k$ , thus reducing the memory footprint.

- 4) Handling same change in modularity. Often, moving vertex  $v_j$  into different communities yields the same change in modularity. In such cases, a tie-breaking heuristics has to be applied to favor one community over the rest. In this paper, a community is picked at random.

Applying these optimizations, the final linear algebraic Louvain method algorithm is shown in Fig. 1 (left).

## V. EXPERIMENTS

In this section, we discuss preliminary experimental results in implementing our formulation of the Louvain method discussing both productivity and performance aspects provided by the use of pygraphblas [6], a Python wrapper to the GraphBLAS [5] API for graph algorithms.

TABLE I

COMPARISON BETWEEN TWO PYTHON IMPLEMENTATIONS OF THE LOUVAIN METHOD: NETWORKX (VERTEX-BASED, SEQUENTIAL) AND PYGRAPHBLAS (PRESENT WORK, LINEAR ALGEBRAIC, BOTH SEQUENTIAL, 1T, AND PARALLEL, 4T). THE NUMBER OF COMMUNITIES (GROUND TRUTH) ARE REPORTED IN COLUMN 3. BOTH IMPLEMENTATIONS REPORTED 100% ACCURACY.

V	Graph		NetworkX		pygraphblas			
	E	C	Time (s)		Time (s) w/o Mask		Time (s) w/ Mask	
			1T	4T	1T	4T	1T	4T
50	319	3	0.02	0.06	0.07	0.07	0.07	0.09
100	778	5	0.03	0.07	0.08	0.08	0.08	0.10
500	9.4k	8	0.31	0.41	0.47	0.48	0.58	
1,000	20.1k	11	0.78	0.79	0.89	0.93	1.11	
5,000	101.9k	19	5.59	7.55	8.19	8.63	9.06	
20k	408.8k	32	26.29	103.99	70.03	113.43	77.17	
50k	1.0M	44	88.12	696.27	387.86	711.83	380.24	

### A. Experimental Setup

We implemented our linear algebraic Louvain method using pygraphblas [6], a Python wrapper of the SuiteSparse [7] library v3.2.0 that implements the C GraphBLAS API, and compared with the implementation of the original vertex-based Louvain method for the NetworkX package [8] v.2.4. Our implementation is available at [6].

We ran our experiments on an Intel Core i7-4770K (Haswell) CPU with four cores, 8 MB LLC cache, and 32 GB of memory. We provide both sequential (1T) and parallel (4T,  $G \times B\_CHUNK=1024$ ) results in Table I.

We report time to solutions (mean of seven runs) for implementations using Static Graphs with known truth for the 2017 Partitioning Challenge provided on the Graph Challenge website [9]. All implementations correctly identified communities when compared to the ground truth.

### B. Discussion

In terms of productivity, our pygraphblas implementation can be mapped nearly one-to-one with the linear algebraic formulation of the algorithm, and requires less than 40 lines of code (LOC). In contrast, the vertex-based NetworkX implementation requires  $4 \times$  as many (158) lines of code (LOC). For a graph algorithm developer, the use of the pygraphblas interface greatly simplifies the implementation; resulting in a boost in algorithmic development productivity.

The ability to port between CPU and GPU platforms to take advantage of different hardware while maintaining the same high-level interface (similar to Blanco et. al [10]) is also another advantage of using GraphBLAS.

Performance-wise, due to the use of SuiteSparse as a black-box library that implement the GraphBLAS API, it is unclear how much of the performance is related to the linear algebraic approach or specific implementation decisions. For example, parallelism was performed by the underlying SuiteSparse library, which may choose to use less than the allocated number of threads. Selecting the chunk size used allowed us some limited form of control over this decision. Nonetheless, we highlight key performance-related features of our implementation.

Lines 23-25 in Fig. 1 (right) are essentially the equivalent of the AXPY operation in the Level 1 BLAS [11]. This can be implemented with a single GraphBLAS apply operation in the most recent GraphBLAS 1.3 as follows:

```
k.apply(FP64.TIMES, first=(-k_j/m),
        accum=FP64.PLUS, out=v)
```

However, as this capability is not supported by the version of SuiteSparse used, three separate function calls, each traversing through a vector of length  $|V|$ , have to be used.

It was interesting to note that while the masking operation as described by Eqn. (2) was meant to reduce the computation cost by considering only neighbor communities when determining changes in modularity, the computation of the mask was a significant cost in our implementation. As such, it was generally faster for our implementation to compute the change in modularity for all existing communities. This cost could potentially be amortized with larger graphs.

A considerable fraction of the runtime was spent extracting from or assigning to single matrix columns or rows. While we have implemented the computation of the change in modularity as a bulk operation, the sequential vertex approach remains a large cost in this implementation.

## VI. CONCLUSION

In this paper, we derived a linear algebraic formulation of the Louvain method for community detection starting from the linear algebraic definition of modularity. We also implemented the derived algorithm using the pygraphblas interface to enhance algorithm design productivity.

We believe that alternative modularity-based clustering algorithms (e.g. the synchronized Louvain algorithm [12]), should enable better performance when expressed in terms of bulk linear algebra operations.

```

1
2
3
4
5
6
7  $G = A + A^T$ 
8
9  $k = A \text{ vec}(1)$ 
10  $m = \frac{1}{2} k^T \text{ vec}(1)$ 
11  $S = I$ 
12
13  $v \text{ changed} = \text{True}$ 
14 while  $v \text{ changed}$ :
15      $v \text{ changed} = \text{False}$ 
16     for  $j$  in  $\text{range}(|V|)$ :
17
18          $t_q = (e_j^T G)S$ 
19
20          $t_j^T = e_j^T S$ 
21          $\bar{S} = (I - e_j e_j^T)S$ 
22
23
24
25          $q^T = e_j^T G + (-k_j/m)k^T$ 
26          $q_1^T = (q^T \bar{S}) \circ (t_q \neq 0)$ 
27
28          $\kappa_q = \max(q_1)$ 
29          $t = (q_1 == \kappa_q)$ 
30         while  $t.\text{nvals}() \neq 1$ :
31              $p = \text{random}() \circ t$ 
32              $\kappa_p = \max(p)$ 
33              $t = (p == \kappa_p)$ 
34          $S = \bar{S} + e_j t^T$ 
35
36
37         if  $t \neq t_j$ :
38              $v \text{ changed} = \text{True}$ 
39
40
41
42

```

```

1 def louvain_cluster(graph):
2     rows = graph.nrows
3     S_row = Vector.from_type(BOOL, rows)
4     empty = Vector.from_type(BOOL, rows)
5
6     G = graph.dup()
7     graph.transpose(out=G, accum=FP64.PLUS)
8     G_rows = [ G.extract_row(i) for i in range(rows) ]
9     k = graph.reduce_vector()
10    inv_m = 2.0 / k.reduce_int()
11    S = Matrix.identity(BOOL, rows, format=lib.GxB_BY_COL)
12
13    vertices_changed = len(k)
14    while vertices_changed > 0:
15
16        for j,k_j in k:
17            v = G_rows[j]
18            t_q = v.vxm(S, semiring=BOOL.ANY_PAIR)
19
20            S.extract_row(j, out=S_row)
21            S[j,:] = empty
22
23            q = k.dup()
24            q.assign_scalar(-k_j * inv_m, accum=FP64.TIMES, mask=k)
25            q += v
26            q1 = q.vxm(S, mask=t_q)
27
28            max_q1 = q1.reduce_float(MAX_MONOID)
29            t = q1.select("==", max_q)
30            while len(t) != 1:
31                p = t.apply(random_scaler)
32                max_p = p.reduce_float(MAX_MONOID)
33                t = p.select("==", max_p)
34            S[j,:] = t
35
36            vertices_changed -= 1
37            if len(S_row * t) == 0:
38                vertices_changed = len(k)-1
39            if vertices_changed <= 0:
40                break
41
42    return S

```

Fig. 1. (Left) Formulation of the linear algebraic Louvain method described in Sec. III and (right) its pygraphblas implementation.

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