ComPAS: Community Preserving Sampling for Streaming Graphs

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Ref: S. Sikdar, T. Chakraborty, S. Sarkar, N. Ganguly, A. Mukherjee: ComPAS: Community Preserving Sampling for Streaming Graphs. AAMAS 2018





Streaming Graphs

- Sequence of edges ordered in time
- Graph G is the aggregation of all the edges over time
- Typical examples include citation network, email log, facebook posts







Streaming Graph Sampling





3





Streaming Graph Sampling with Community

- Given a streaming graph G, the objective is obtain a sample G_s such that the properties of G are maintained in G_s
- Existing algorithms are designed for preserving simple structural properties
- We propose ComPAS which is capable of retaining the underlying community structure
- Applications Obtaining stratified samples in online learning















Proposed Algorithm: ComPAS

- Maximize modularity
- Identify high fidelity nodes over time
- Allow merging, splitting and creation of new communities





Proposed Algorithm: ComPAS

Parameters:

- sample size (n)
- alpha $(0 < \alpha < 1)$
- Buffer (H) consisting of two variables
 - H_c Number of times a node is encountered
 - H_p Current parent

Node	Count	Parent
i	1	d
j	3	1
k	1	m
1	4	j
m	3	e
n	1	k







Dynamics of ComPAS

 Keep adding edges into the sample as long as a certain number of nodes are inserted (*α* * n)



• Once the threshold is reached a pre-selected community detection algorithm is executed on the sample to obtain initial community structure.







Role of Buffer

- From this point on whenever a new node is encountered it is pushed to buffer
- Estimate the importance of a node
- More recurrent node is perhaps more important

Node	Count	Parent
i	1	d
j	3	1
k	1	m
1	4	j
m	3	e
n	1	k







10





Genesis of Six Modules

- Both vertices in the sample
- Both vertices in buffer
- One in sample and one in buffer
- One in sample and one is new
- One in buffer and one is new
- Both are new
- Constraints
 - A new node cannot be directly added to the sample
 - Only nodes from buffer are eligible to enter the sample
 - If sample size is reached node must be deleted to make way





Both in Sample

This can be further divided into two sub cases -

• The edge is intra-community



Add the edge to the sample

• The edge is inter-community



- u may leave its current community and join v's
- v may leave its current community and join u's
- u and v leave their current communities and form new one





• edge (j,k)

Node	Count	Parent	Node	Count	Parent
i	1	d	i	1	d
j	3	1	j	4	1
k	1	m	k	2	m
1	4	j	1	4	j
m	3	e	m	3	e
n	1	k	n	1	k





One in Sample one in Buffer

• edge (u,k)

Node	Count	Parent		Node	Node Count
i	1	d		i	i 1
j	4	1		j	j 4
k	2	m		k	k 3
1	4	j		1	1 4
m	3	e		m	m 3
n	1	k		n	n l





Dynamics of ComPAS

Both vertices in the sample
Both vertices in buffer
One in sample and one in buffer
One in sample and one is new
One in buffer and one is new
Both are new





Dynamics of ComPAS

Both vertices in the sample
Both vertices in buffer
One in sample and one in buffer
One in sample and one is new
One in buffer and one is new
Both are new

At least one node is new





Entry of a new Node

- In the subsequent cases at least one node is new
- This node triggers rearrangement -
 - Remove node from buffer to make way for new node Preferentially (based on $H_c(x)$) remove a node x from buffer with additional constraint that P(x) in sample
 - Remove node from sample to make way for x Node with lowest degree and clustering coefficient is removed from sample







Deletion of a Node from Sample

- New node (v) is encountered
- Buffer is full
- Sample size has been reached
 - 1. Preferentially select u from buffer and add it to sample
 - 2. Assign u the community of its parent P(u)
 - 3. Remove a node w with the lowest degree and clustering coefficient from sample
 - 4. Add v to buffer (cannot be directly added to the sample)





edge: (u,v)

- u is in sample and v is new
 - v is inserted into buffer which might trigger rearrangement of the buffer and sample
- u is in buffer and v is new
 - Increase $H_c(u)$ by 1
 - Insert v into buffer
- Both u and v are new
 - Insert both u and v into buffer







20





Evaluation

- Experiments performed on 4 real-world and 1 synthetic datasets
- Two ways of evaluation
 - Quality of the community structure
 - Content of the communities
- Baselines -
 - Streaming node (SN), streaming edge (SE), streaming BFS (SBFS) and Partially induced edge sampling (PIES)
 - Novel Green Algorithm (sample obtained on aggregated graph)





Evaluation

- Quality of community structure
 - Based on 13 topological measures proposed by Yang and Leskovec
 - Structural properties like average degree, internal density ... (calculated for each community)
- We compare using D-statistics -
 - Consider a property X
 - Calculate distribution of X across communities in the ground-truth (f(X)) and the obtained sample g(X)
 - Calculate D-statistics between f(X) and g(X)







Evaluation

- Content of the community structure
- Similarity measured through -
 - Purity
 - Normalized Mutual Information (NMI)
 - Adjusted Rand Index (ARI)

ComPAS outperforms all other streaming graph sampling algorithm





Future directions

- Theoretical guarantees on the quality of the sample
- Complexity of the algorithm
- Allow deletion of edges over time





Thank You

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Ref: Sandipan Sikdar, Tanmoy Chakraborty, Soumya Sarkar, Niloy Ganguly, Animesh Mukherjee: ComPAS: Community Preserving Sampling for Streaming Graphs. AAMAS 2018



