

# CMPT 843 Project

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# 1 Motivation and Application Background

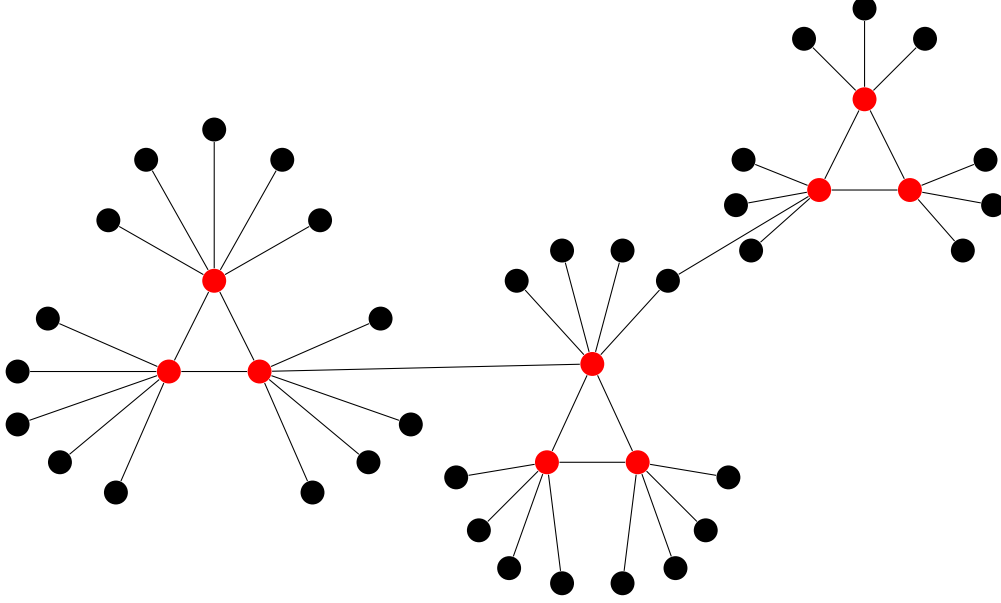
0.5 pages My own application of the method. Also talk about the extra work I did (reproducing the experiments and adding extra measures).

## 2 Background

Triangles can be a more or less important structure of a graph. If many nodes are reachable from those nodes that are forming a triangle this triangle is important to connect the remaining nodes. In this case the triangle is serving as a hub for the nodes around it. Removing the nodes that form the triangle or the edges between them would disconnect a large part of the graph. The importance of one triangle can be calculated as the number of nodes that are adjacent to it. Let this sum be defined as the *connectivity* of a triangle. Figure 1 shows a graph with three triangles with *connectivity* 15, 14 and 10. The average *connectivity* of triangles in this graph is  $\frac{15+14+10}{3} = 13$ . If we want to find the average *connectivity* of triangles in a large scale graph it makes sense to sample a number of triangles and observe the *connectivity* of the sampled triangles. In this report I document the application of the *Neighborhood Sampling* method as introduced in the paper "Counting and Sampling Triangles from a Graph Stream" to estimate the average connectivity of the overall population of triangles in the graph. Here I will limit the experiments to triangles for which the adjacent nodes are distributed evenly across the three nodes of the triangle as in the graph of Figure 1.

## 3 Solution and Analysis

To test how well the presented method is suitable to find the average connectivity of triangles in a graph I will first generate a graph with  $\tau$  triangles. For each triangle I draw a random number  $r$  between 1 and  $a$  and assign each node of the triangle  $r$  adjacent nodes (thus this triangle gets  $3 \times r$  adjacent nodes). All in all the generated graph consists of only triangles, where each triangle gets a random number of adjacent nodes that are equally distributed among the triangle nodes. Next I shuffle the order of the edges to generate a graph stream in which the edges appear in random order. Now I use the *Neighborhood Sampling* method to sample triangles from this stream where I use choose  $e$  estimators. As described in the paper each estimator can either hold a triangle or not. For the case that an estimator is holding a triangle we take this triangle into our sample with probability  $\frac{c}{2\Delta}$  ( $c$  is the number of edges the estimator observed that are adjacent to the first edge of the



**Figure 1:** all nodes in the graph are adjacent to one of three triangles

triangle that arrived in the stream and  $\Delta$  is the maximum degree of a node in the graph). For each triangle that I get in my sample I estimate the total number of adjacent nodes as described in 3.1. Finally I compare the mean and variance of the *connectivity* that I got from my sample with the actual mean and variance of the triangle population.

The sampling method is simple random sampling with replacement. Therefore I expect that I get an unbiased estimator for the overall *connectivity* of the triangles. The size of the samples that the method allows me to draw depends both on the number of estimators and the maximum degree  $\Delta$  in the generated graph. My estimation should get more accurate the higher I choose the number of estimators  $e$  as more estimators return a larger sample. At the same time my estimation should become less accurate, the larger I choose  $a$  because a larger  $a$  results in a larger value  $\Delta$  which decreases the size of the sample I can draw from the estimators.

### 3.1 Estimating the number of adjacent nodes of a triangle

The estimators from which the triangles are being sampled are first sampling a random edge  $r_1$  from the stream, then they sample a random edge  $r_2$  from the substream of edges that arrive after  $r_1$  and are adjacent to it. Finally they sample an edge  $r_3$  that closes the wedge  $r_1 r_2$  and forms the triangle

$r_1 r_2 r_3$ . Each estimator keeps track of the number of edges it observes that are adjacent to  $r_1$  and arrive after it in the stream. This number is kept in a counter  $c$ . Thus  $c$  keeps track of the number of edges that arrive after  $r_1$  and which are adjacent to either of two nodes of the triangle (the two nodes of the triangle that are connected by  $r_1$ ). For each estimator that is returning a triangle into my sample I estimate the total number of adjacent nodes to this triangle as  $((c - 2)\frac{4}{3}) \times \frac{3}{2}$ . The term  $(c - 2)$  is the number of  $r_1$ -adjacent edges that were observed in the stream after  $r_1$  excluding the two edges that belong to the triangle ( $r_2$  and  $r_3$ ). The term  $(c - 2)\frac{4}{3}$  represents the  $(c - 2)$  edges that are adjacent to  $r_1$  and arrived after  $r_1$  in the stream *plus* the expected number of edges that arrived in the stream before  $r_1$  (the expected number of  $r_1$  adjacent edges that arrived before  $r_1$  is  $\frac{1}{3}(c - 2)$ ). Thus the term  $((c - 2)\frac{4}{3})$  represents the number of edges that are adjacent to the  $r_1$  edge of the triangle. The edge  $r_1$  represents two nodes of the triangle. My estimate for the number of edges that are adjacent to the third node of the triangle is  $\frac{1}{2}((c - 2)\frac{4}{3})$ , because I am assuming that the number of adjacent nodes are equally distributed among the three triangle nodes. The total number of adjacent edges to this triangle is then  $((c - 2)\frac{4}{3}) \times \frac{3}{2}$ .

## 4 Experimental Results

### 4.1 Method Implementation

First I implemented the method from the paper and verified that my implementation is correct by reproducing the experiments from the paper with the same datasets which I downloaded from <https://snap.stanford.edu/data>. To verify my implementation I downloaded the amazon, dblp and livejournal graphs and got similar results as the authors of the paper. Further I downloaded a Protein-Protein-Interaction Network from <http://thebiogrid.org/> and ran the method on this dataset. In addition to the number of triangles that was estimated by the method, I noted the number of triangles that I was able to sample from the chosen number of estimators as listed in Table 1.

### 4.2 Testing of the method on simulated graphs

I tested the method for triangle sampling with the following parameters for  $\tau$  (number of generated triangles),  $a$  (maximum number of adjacent nodes that gets assigned to a triangle node) and  $e$  (number of estimators from which we sample the triangles). In Table 2  $\mu$  lists the mean *connectivity* of

$\tau, a \Rightarrow \mu, v$	$e = 10000$	$e = 100000$	$e = 1000000$
500, 10 $\Rightarrow$ 16, 115	$s = 21, \mu_{est} = 16, var_{est} = 119$	$s = 204, \mu_{est} = 14, var_{est} = 99$	$s = 2273, \mu_{est} = 16, var_{est} = 105$
500, 30 $\Rightarrow$ 46, 962	$s = 2, \mu_{est} = 20, var_{est} = 360$	$s = 35, \mu_{est} = 46, var_{est} = 936$	$s = 302, \mu_{est} = 44, var_{est} = 939$
500, 50 $\Rightarrow$ 76, 2633	$s = 1, \mu_{est} = 91, var_{est} = 0$	$s = 11, \mu_{est} = 57, var_{est} = 2129$	$s = 124, \mu_{est} = 79, var_{est} = 2841$
500, 100 $\Rightarrow$ 152, 10576	$s = 0, \mu_{est} = \emptyset, var_{est} = \emptyset$	$s = 2, \mu_{est} = 112, var_{est} = 6210$	$s = 30, \mu_{est} = 158, var_{est} = 9290$
1000, 10 $\Rightarrow$ 16, 116	$s = 18, \mu_{est} = 15, var_{est} = 72$	$s = 196, \mu_{est} = 15, var_{est} = 90$	$s = 2157, \mu_{est} = 15, var_{est} = 100$
1000, 30 $\Rightarrow$ 45, 959	$s = 5, \mu_{est} = 68, var_{est} = 647$	$s = 31, \mu_{est} = 46, var_{est} = 766$	$s = 311, \mu_{est} = 43, var_{est} = 776$
1000, 50 $\Rightarrow$ 76, 2757	$s = 1, \mu_{est} = 19, var_{est} = 0$	$s = 15, \mu_{est} = 78, var_{est} = 1903$	$s = 133, \mu_{est} = 74, var_{est} = 2388$
1000, 100 $\Rightarrow$ 155, 11478	$s = 0, \mu_{est} = \emptyset, var_{est} = \emptyset$	$s = 2, \mu_{est} = 323, var_{est} = 573$	$s = 36, \mu_{est} = 150, var_{est} = 9374$
2000, 10 $\Rightarrow$ 16, 118	$s = 27, \mu_{est} = 19, var_{est} = 75$	$s = 223, \mu_{est} = 16, var_{est} = 106$	$s = 2078, \mu_{est} = 15, var_{est} = 99$
2000, 30 $\Rightarrow$ 46, 996	$s = 2, \mu_{est} = 55, var_{est} = 1437$	$s = 29, \mu_{est} = 39, var_{est} = 940$	$s = 323, \mu_{est} = 42, var_{est} = 762$
2000, 50 $\Rightarrow$ 76, 2766	$s = 1, \mu_{est} = 7, var_{est} = 0$	$s = 16, \mu_{est} = 82, var_{est} = 1710$	$s = 131, \mu_{est} = 79, var_{est} = 2614$
2000, 100 $\Rightarrow$ 150, 10605	$s = 0, \mu_{est} = \emptyset, var_{est} = \emptyset$	$s = 2, \mu_{est} = 69, var_{est} = 255$	$s = 28, \mu_{est} = 155, var_{est} = 7854$

**Table 1:** This table lists my experiment results. It can be read as follows: The first column lists the propoerties of the generated graphs. The column entry 1000, 50  $\Rightarrow$  76, 2757 means I generated a graph that consists of 1000 triangles where for each triangle  $t$  I draw a number  $r$  between 1 and 50 and assigned each node of  $t$   $r$  adjacent nodes. The mean *connectivity* of the triangles in this graph was 76 and the variance of the *connectivity* was 2757. Then I used  $10^3, 10^4$  and  $10^5$  estimators to sample triangles from this graph. For  $10^4$  estimators for example I could draw  $s = 15$  triangles into my sample. For each of these triangles I estimated the total number of adjacent nodes as explained in 3.1. Finally I computed the mean and variance *connectivity* of my sample as  $\mu_{est}$  and  $var_{est}$ .

the generated triangles and  $v$  lists the *connectivity*-variance of the generated triangles for the chosen parameters  $\tau$  and  $a$ . In the table cells,  $s$  lists the sample size that I got from the  $e$  estimators and  $\mu_{est}, var_{est}$  list the mean and variance of *connectivity* I got from my sample.

## 5 Conclusion and Discussion

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