Structural Patterns and Generative Models of Real-world Hypergraphs (User Guide)

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1 General Information

This is a guide for using the code of the paper 'Structural Patterns and Generative Models of Real-world Hypergraphs" ³.

2 Introduction

The paper introduces a decomposition tool, which facilitates convenient analysis of hypergraphs, and proposes a generator model, *HyperPA*, which generates hypergraphs having real-world patterns. This github repository contains:

- Decomposition: produces decomposed graphs of a hypergraph.
- Generator: consists of our proposed model, HyperPA, together with 2 baseline models.

3 Decomposing a hypergraph

- The decomposition code requires Java be installed in the system.
- Specify the directory containing the input hypergraph at Line 118 String directory = "";
- The files containing the decomposed graphs will be in the same directory.

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• Run main.java to generate the edge-level, triangle-level, 4clique-level decomposed graphs.

3.1 Input File Format of decomposition

main.java assumes that in the input hypergraph G=(V,E), each node $v\in V$ has its unique integer id . In the input file, each line represents a single hyperedge. Each hyperedge $e\in E$ consists of an arbitrary number of nodes, seperated by spaces. This is the format of the datasets under the Datasets repository. Examples on a sample sample_hypergraph.txt, in this case, contains only 2 hyperedges: (1,2,3),(2,3), formated as follows:

E.g. sample_hypergraph.txt

1	2	3	
2	3		

Note that as described in the main paper, only hyperedges of sizes less than or equal to 7 are considered for these 3 levels of decomposition.

3.2 Output File Format of decomposition

For each decompostion level, there are 2 output files: one contains the corresponding decomposed graph with translated node ids, the other one contains labels of the translated node ids. In the decomposed graph file, each line is a clique of nodes, separated by spaces. In the label file, each line provides a label for which subset of nodes in the original hypergraph corresponds to which node id in the decomposed graph file. For example, for the edge-level decomposed graph of sample_hypergraph.txt:

E.g. sample_hypergraph-edge_level-node-labels.txt

```
\begin{bmatrix} 1 & 1, 2 \\ 2 & 1, 3 \\ 3 & 2, 3 \end{bmatrix}
```

According to this label file, subset of nodes 1, 2 corresponds to the node id 1 in the decomposed graph file, subset 1, 3 corresponds to 2, 2, 3 corresponds to 3, respectively. In the edge-level decomposed graph, the hyperedge (1,2,3) results a clique of 3 nodes (1,2)-(1,3)-(2,3), and the hyperedge (2,3) results in a single node (2,3). Therefore, the format of the output decomposed graph is:

E.g. sample_hypergraph-edge_level-decomposed.txt

1	2	3	
3			

4 Generating a Hypergraph

Under the Generator directory, there are 3 code files SS.py, hyper_preferential_attachment.py, preferential_attachment.py corresponding to the 3 models mentioned in the main paper: Subset Sampling, HyperPA, and NaivePA, respectively.

4.1 How to Generate

Run the corresponding code file to generate the hypergraph based on the generator.

4.2 Parameters

Each generator depends on the following input files and parameters

- Parameter name: Path to the output file.
- Parameter *num_nodes*: Number of nodes for the output hypergraph.
- Input file $size_distribution$: directory of this input file must be specified in the method $learn_size_distribution()$. This method learns the distribution of hyperedge sizes, from which the size of each new hyperedge will be sampled. The input file must follow the same format as the files under the "size distribution" repository: the number in line the *i*-th is the relative number of hyperedges of size *i* that will be in the hypergraph. The probability of having each particular hyperedge size is computed based on these numbers.

For example, if the size distribution file is as follow:

E.g. sample_size_distribution.txt



then in the generated hypergraph, the proportions of hyperedges of sizes 1, 2, 3, 4 will be approximately 20%, 30%, 25%, 25%, respectively.

• Inputfile simplex per node: directory of this input file must be specified in the method learn_number_simplices_per_node(). This method learns the distribution of number of hyperedges per new node, from which the number of new hyperedges introduced by each new node will be sampled. The input file must follow the same format as the files under the "simplex per node" repository: the number in the line i-th is the relative number of nodes that introduce i new hyperedges when they arrive.

For example, if the simplex_per_node file is as follow:

E.g. sample_simplex_per_node.txt

10		
20		
30		
30		
20 30 30 10		

then in the generated hypergraph, for each new node, there is a 10% chance this node brings 1 new hyperedge to the hypergraph, 20% chance for 2 hyperedges, 30% for 3 hyperedges, 30% for 4 hyperedges, and 10% for 5 hyperedges.

5 Format of Output File

In the output file, each line represents a hyperedge where nodes are separated by spaces. For example, if the generated hypergraph has 3 hyperedges: (1,2), (1,3,4), (2,3,4,5), the output file will be as follow:

 $E.g.\ sample_generated_hypergraph.txt$