Notes on Graph Representation Learning

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1 Representing Graphs

Formally, a graph is defined as follows.

$$\mathcal{G} = (\mathcal{V}, \mathcal{E}). \tag{1}$$

In which, \mathcal{V} and \mathcal{E} are the nodes and edges of the graphs. Edges going from $u \in \mathcal{V}$ to $v \in \mathcal{V}$ are represented as $(u, v) \in \mathcal{E}$.

Graphs can be represented using an adjacency matrix $\mathbf{A} \in \mathbb{R}^{|\mathcal{V}|x|\mathcal{V}|}$, in which A[u,v] represent $(u,v) \in \mathcal{E}$. Given output unit of every node $\mathbf{u} \in \mathbb{R}^{|\mathcal{V}|}$, $\mathbf{s} = \mathbf{A}\mathbf{u}$ is the combined weighed signal from every adjacent node. Moreover, $\mathbf{A}^i[u,v]$ represents the number of *i*-length paths that connects u and v.

Nodes within a graph may have features or attributes associated with them. They can be represented with matrix $\mathbf{X} \in \mathbb{R}^{|V| \times m}$, for nodes with m features.

1.1 Graph Laplacians

Graph Laplacians have many useful mathematical properties.

1.1.1 Unnormalized Laplacian

$$\mathbf{L} = \mathbf{D} - \mathbf{A},\tag{2}$$

where \mathbf{D} is matrix with node degrees (see Equation 3) on its diagonals. The Laplacian summarizes many important properties of the graph.

2 Types of Graphs

- **Simple graphs** are graphs with at most one edge between each pair of nodes, and all edges are undirected.
- **Heterogeneous graphs**: Nodes have types, and edges satisfy constraints according to those types.
- Multiplex graphs: Graph is decomposed in a set of k *layers*. Every node is assumed to belong to every layer, and each layer corresponds to a unique relation.

3 Machine Learning on Graphs

The following lists the types of tasks that may be done on graphs.

• Node classification: Classify a node based on its features and relations to other nodes. Ideas such as homophily, which is the tendency for nodes to share attributes with their neighbors; structural equivalence, which is the idea that nodes with similar local neighborhood structures will have similar labels; and heterophily, which presumes that nodes will be preferentially connected to nodes with different labels, have been used to assist node classification.

- Relation prediction: Also known as link prediction, graph completion, relational inference, is the task of inferring edges between nodes in a graph.
- Clustering and community detection: Finding subgroups that may be useful.
- Graph classification, regression, and clustering: Graphs are treated as data points, which is to be classified, regressed, or clustered.

4 Graph Statistics and Features

The following lists node-level statistics and features.

• Node degree: The number of edges incident to a node.

$$d[u] = \sum_{v \in V} \mathbf{A}[u, v], \tag{3}$$

$$\mathbf{d} = \mathbf{A}\mathbf{1}_{|V|},\tag{4}$$

in which, $\mathbf{1}_{|V|} = (1, ..., 1) \in \mathbb{R}^{|V|}$.

• Node centrality: Eigenvector centrality is a measure of node importance. The eigenvector \mathbf{e} corresponding to the largest eigenvalue λ contains the importance of each node on its indices.

$$\lambda \mathbf{e} = \mathbf{A}\mathbf{e} \tag{5}$$

$$\mathbf{e}[u] = \frac{1}{\lambda} \sum_{v \in V} \mathbf{A}[u, v] \mathbf{e}[v]. \tag{6}$$

• Clustering coefficient: Measures the degree of clustering for a node. It is the ratio between the number of adjacent node pairs that are also adjacent to each other by the total combination of node pairs for u.

$$c[u] = \frac{|(v_1, v_2) \in \mathcal{E} : v_1, v_2 \in \mathcal{N}(u)|}{\binom{d[u]}{2}}.$$
 (7)

An alternative way of viewing the clustering coefficient is that it counts the number of closed triangles that is connected to a particular node.

The following lists graph-level features and kernels.

- Bag of nodes: An aggregated view of node-level representations. Such as histograms of degrees, centralities, and clustering coefficients of the nodes in the graph.
- Weisfeiler-Lehman (WL): Is a kernel and an algorithm that iteratively aggregates the neighborhood to find statistics beyond the immediate neighbor of a node. The algorithm is as follows:

- 1. Assign an initial label to each node. In many cases, this is the degree of each node; $l_0(v) = d[v]$.
- 2. Next, a new label is iteratively assigned with an aggregation function.

$$l_i(v) = AGGR(\{l_{i-1}(u) \ \forall u \in \mathcal{N}(v)\})$$
 (8)

- 3. After K iterations we have $l_K(v)$, which is a K-hop summary of every node. Which can be used to compare nodes at a higher level.
- **Graphlets**: Graphlets are subgraph structures that may commonly exhibit within larger graphs. The number of these different structures can be treated as graph-level features.

The following lists statistics on node-to-node relationships within graphs.

• Similarity matrix: $\mathbf{S} \in \mathbb{R}^{|V| \times |V|}$ is a matrix containing the number of shared neighbors of each node pairs.

$$\mathbf{S}[u,v] = |\mathcal{N}(u) \cap \mathcal{N}(v)|. \tag{9}$$

• Sorenson index: Similarity matrix with normalization.

$$\mathbf{S}_{\text{Sorenson}}[u, v] = \frac{2\mathbf{S}[u, v]}{d[u] + d[v]}.$$
 (10)

 \bullet Salton index: Similarity matrix with normalization.

$$\mathbf{S}_{\text{Salton}}[u, v] = \frac{2\mathbf{S}[u, v]}{\sqrt{d[u]d[v]}}.$$
 (11)

• Jaccard index: Similarity matrix with normalization.

$$\mathbf{S}_{\text{Jaccard}}[u, v] = \frac{\mathbf{S}[u, v]}{|\mathcal{N}(u) \cup \mathcal{N}(v)|}.$$
 (12)

• Resource allocation (RA) index: Counts the inverse degrees of common neighbors,

$$\mathbf{S}_{\mathrm{RA}}[v_1, v_2] = \sum_{u \in \mathcal{N}(v_1) \cap \mathcal{N}(v_2)} \frac{1}{d[u]}$$
 (13)

• Adamic-Adar (AA) index: Similar to RA index but using the inverse log.

$$\mathbf{S}_{\mathrm{AA}}[v_1, v_2] = \sum_{u \in \mathcal{N}(v_1) \cap \mathcal{N}(v_2)} \frac{1}{\log(d[u])}$$
(14)

• **Katz index**: Counts the number of paths of all lengths between a pair of nodes.

$$\mathbf{S}_{\mathrm{Katz}}[u,v] = \sum_{i=1}^{\infty} \beta^{i} \mathbf{A}^{i}[u,v], \tag{15}$$

where $\beta \in \mathbb{R}^+$ is a user-defined parameter controlling how much weight is given to short versus long paths.

• LHN similarity: A normalized version of the Katz index, which reduces a high-degree bias in the Katz index.

$$\mathbf{S}_{\text{LNH}}[u, v] = \mathbf{I}[u, v] + \frac{2m}{d[u]d[v]} \sum_{i=0}^{\infty} \beta^{i} \lambda_{1}^{1-i} \mathbf{A}^{i}[u, v], \quad (16)$$

in which λ_1 is the largest eigenvalue of **A**.

• Random walk similarity: Similarity between two nodes is proportional to how likely we are to reach each node from random walk starting from the other node.

$$\mathbf{P} = \mathbf{A}\mathbf{D}^{-1}.\tag{17}$$

Here $\mathbf{P}[\mathbf{u}, \mathbf{v}]$ is a stochastic adjacency matrix, with adjacency probability scaled proportional to the node's inverse degree.

$$\mathbf{q}_{u} = c\mathbf{P}\mathbf{q}_{u} + (1-c)\mathbf{e}_{u}.\tag{18}$$

This implicit equation models the probability of reaching each node with a random walk policy. The c term determines the probability that the random walk restarts at u and \mathbf{e}_u is a one-hot indicator vector for node u. The solution to this recurrence is as follows.

$$\mathbf{q}_u = (1 - c)(\mathbf{I} - c\mathbf{P})^{-1}\mathbf{e}_u. \tag{19}$$

Finally, random walk similarity formulated as follows.

$$\mathbf{S}_{\mathrm{RW}}[u,v] = \mathbf{q}_{u}[v] + \mathbf{q}_{v}[u]. \tag{20}$$