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Hierarchical Graph Neural Network Based-on Semi-implicit Variational Inference

Hai-Long Su*, Zhi-Peng Li, Xiao-Bo Zhu, Li-Na Yang, Valeriya Gribova, Vladimir Fedorovich Filaretov, Anthony Cohn, De-Shuang Huang*, *Fellow, IEEE*,

Abstract—Graph neural network(GNN) has obtained outstanding achievements in relational data. However, these data have uncertain properties, for example, spurious edges may be included. Recently, Variational graph autoencoder(VGAE) has been proposed to solve this problem. However, the distributional assumptions in the variational family restrict the variational inference (VI) flexibility and they define variational families using mean-field, which can not capture complex posterior distributional. To solve the above question, in this paper, we proposed a novel GNN model based on semi-implicit variational inference (SIVI), which can embed the node to the latent space to improve VI flexibility and enhance VI expressiveness with mixing distribution. Specifically, to approximate the true posterior, a variational posterior was given utilizing a semi-implicit hierarchical variational framework, which can model complex posterior. Moreover, an iterative decoder is used to better capture graph properties. Besides, due to the hierarchical structure in our model, it can incorporate neighbour information between nodes. Experiments on multiple data sets, our method has achieved state-of-the-art results compared to other similar methods. Particularly, on the citation dataset Citeseer without features, our method outperforms VGAE by nine percentage.

Index Terms—Latent variable, Variational inference, Graph neural network, Semi-implicit model, Hierarchical frame.

I. INTRODUCTION

CONVOLUTIONAL neural networks (CNNs) have made great achievements in the past ten years in the fields of speech [1], image [2, 3] and other fields [4]. However, CNNs can only handle normalized data, e.g., grids, sequences, i.e., Euclidean space data, which has translation invariance. Nevertheless, in reality there is a lot of data that is in non-European space, such as social network data, protein and

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protein (PPI) interaction data, transportation data, etc [8]. To process this data efficiently, graph neural networks (GNNs) have been proposed and also achievements huge progress in many fields, such as social networks [5], recommendation systems [6], protein-protein interaction [7] and action prediction [8][9]. Kipf *et.al* [10] proposed graph convolutional network (GCN), which learns the first-order approximate of nodes and followed a nonlinear activation function to learn graph representation. Considered neighbourhood information between nodes, Hamilton *et.al* [11] proposed GraphSAGE, a inductive representation learning method, which can aggregate neighbourhood information by aggregate function. GraphSAGE is designed to generate low-dimensional vector representations of nodes and is particularly useful for graphs with rich information on node attributes. Different neighbour nodes have different weights, specifying different weights to different nodes in the neighbourhood that can prevent redundant information from being aggregate, hence, Veličković *et.al* [12] proposed graph attention networks(GATs) use masked self-attentional layers. Besides, based on the GNN, some interesting works were also proposed by scholars, such as graph pool [13], place classification [14], and facial expression recognition [15].

Although it's very effective in dealing with relational data, there are some challenges. For example, graphs are very huge in nature, graph structural information is ignored. To eradicate the uncertain problem, Bayesian-based approaches were proposed. For instance, Zhang *et.al* proposed BGCN [16] that incorporate uncertainly graph information via parametric random graph model. However, BGCN relies heavily on the selection of random graphs, which ignore the node features and training labels. To overcome this drawback, Pal *et.al* [17] introduced a non-parametric BGCN, which uses the node features, training labels, and observed graph for posterior inference. Following this idea, many Bayesian-based approaches are proposed [18–20].

Variational autoencoders(VAEs) is a popular method for unsupervised representation learning of high dimensional data [21]. Inspired by VAEs, variational graph autoencoders(VGAEs) [22] was proposed. VGAE uses GCN as an encoder and a simple inner product as the decoder. VGAE assumes the variational posterior is Gaussian distribution, which is restricted variational inference flexibility. And inner product decoder limits the ability to generate models. Recently, semi-implicit variational inference (SIVI) [24] and normalizing flow (NF) [25–27] are proposed, which provide flexibility posterior distribution and effective optimization. Expansion

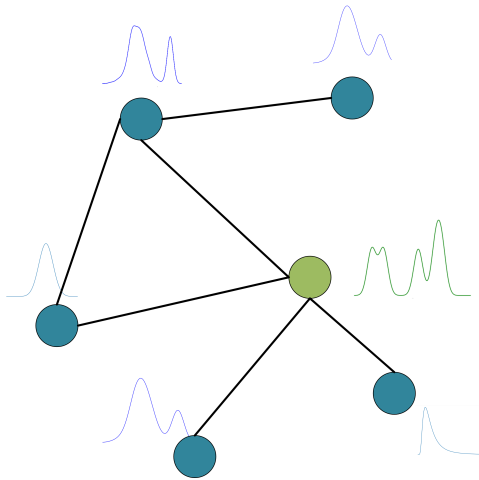


Fig. 1. Node distribution propagation. Our method can diffuse the node's distributions between neighbors.

SIVI to graph domain, SIG-VAE [28] was proposed. SIG-VAE introduced a semi-implicit model to learn the posterior, which can well simulation the complex posterior with heavy tails, skewness, multimodality characteristics.

Through parameterizing VAEs by GNN, Grover *et.al* [29] proposed Graphite, a framework for large graph nodes representation. Graphite uses forward message passing for encoding nodes to latent space and reverse message passing for decoding. Despite the excellent results Graphite obtained, however, in the encoder step, Graphite assumes the posterior is isotropic Gaussian, which results in high variance, and it restricts the expressive ability, hence, it can not model complex posterior.

Inspired by SIVI and SIG-VAE, we proposed a hierarchical graph neural network based on semi-implicit variational inference. Specifically, to handle Graphite's distribution assumptions and inability to model complex distributions, we proposed a hierarchical frame for our model in the encoder step, enhancing the expressiveness of posterior distribution for nodes in the latent space. For do this, we construct stochastic multi-layers hierarchically and inject random noise at every layer. After doing this, the output of GNN is random variables rather than deterministic, hence, uncertainty in the structure of a graph can be measured. More specifically, when obtain node's latent posterior distributions, the distributions of their neighboring nodes are incorporated simultaneously, which is very important for graph representation, as shown in Fig.1. The distributions of aqua green nodes can propagation to olive node. In the decoder step, following Graphite, reverse message passing is used to construct the graph. For model inference, we derive a lower evidence lower bound (ELBO) followed by SIVI [24].

The contributions of our proposed method are as follows:

- 1) We proposed a new GNN model, which can propagation uncertainty between neighbourhood nodes. Contrary to determinant GNN, such as GCN and GAT, our method can metric the uncertainty of graph structure, which is very important for information aggregation.
- 2) We proposed a new encoder framework for variational

posterior learning. Specifically, different from the traditional encoder, which assumes posterior to be Gaussian distribution, we use a hierarchical design for model to learn the parameters of posterior, then, we can obtain a more expressive posterior, which can model complex graph data.

- 3) We use our proposed method to perform link prediction task and conduct extensive experiments on 3 citation datasets(ie, Cora, Cisteer, Pubmed) and 4 different scenario datasets, our method obtains start-of-the-art results compared with the baseline method.

II. PRELIMINARIES

Given a graph $\mathbf{G} = (\mathbf{E}, \mathbf{V})$ where \mathbf{E} and \mathbf{V} refer to edges and nodes of the graph respectively. Additionally, the feature matrix of the graph defined as a m -dimensional signal $\mathbf{X} \in \mathbb{R}^{m \times n}$ associated with each node. In this paper, graph structure is represented by symmetric adjacency matrix $\mathbf{A} \in \mathbb{R}^{n \times n}$ where $n = |\mathbf{V}|$ and $\mathbf{A}_{i,j} = 1$ denote there is an edge between node i and j and $\mathbf{A}_{i,j} = 0$ on the contrary.

A. Graph Neural Networks

Sperduti *et.al* [30] applied neural network to solve directed acyclic graphs, which is regarded as a motivation of GNN. The notion of GNN was declared by Gori *et.al* firstly [31]. After that, Scarselli *et.al* [32] and Gallicchio *et.al* [33] further developed GNN. The intuitive idea behind GNN is message passing between nodes and their neighbourhood by an iterative manner until a fixed point appears or converges. However, this process is inefficient computation, hence many works attempts to overcome this problem [34] [35].

Drawing on the ideas of CNNs, convolutional GNN (ConvGNN) has been proposed. There are two streams of ConvGNN: spectral-based and spatial-based approaches. The first spectral-based method was proposed by Bruna [36], which introduced a graph convolution based on spectral graph theory. Inspired by this method, many works have been raised recently [10, 37, 38]. Spatial-based approaches have been proposed recently by scholars. Compared with spectral-based methods, which involved spectral graph theory, spatial-based approaches operate directly on the nodes of graph. The spatial-based approach maximizes the use of the rich information of the nodes, hence, many researchers focus on this area. For example, GraphSAGE [12] and GAT [11] are proposed benefits from effective node aggregation. In order to capture high-order information of neighbour nodes, multi-hop neighbourhood aggregation is essential. Hence, based on Weisfeiler-Leman(WL) [39], high order GNN has proposed [40, 41], which extension 1-WL test to k-WL test to capture multi-hop information of nodes. Furthermore, many VAE-based [42, 43] GNNs are also proposed. Such as VGAE [22], GraphVAE [44], simple-GVAE [45], CGVAE [46].

B. Semi-implicit Variational Inference

The vanilla variational inference (VI) has the following drawbacks: 1) optimization difficulties when learning posterior

distributions, 2) cannot resolve the distribution with skewness, kurtosis, multimodality, and other characteristics. To overcome the above questions, Semi-implicit Variational Inference(SIVI) [24] has been proposed. SIVI is similar to HVM proposed by Ranganath *et.al* [47]. SIVI assumes that posterior parameters are derived from an implicit distribution rather than being analytic. SIVI also has a hierarchical structure, which can capture complex posterior distribution. More specifically, assuming $\varphi \sim q_\psi(\varphi)$, where ψ denotes distribution parameter can be inferred, the SIVI distribution for Z can be defined as a hierarchical manner: $Z \sim p(Z|\varphi), \varphi \sim q_\psi(\varphi)$. In order to obtain a tractable variational posterior, SIVI derives a lower bound for ELBO [48–51] to optimize the variational parameters.

III. PROPOSED METHOD

When we use probabilistic graphical model for graph representation learning, we are interested in learning a parameterized distribution over adjacency matrix \mathbf{A} and the nodes feature \mathbf{X} is added as conditioning evidence. In this paper, to achieve this goal, we induct a latent variable $\mathbf{Z}_i \in \mathbb{R}^k$ and the node feature $\mathbf{X}_i \in \mathbb{R}^m$ for each node $i \in 1, 2, \dots, n$ along with $\mathbf{A}_{i,j} \in \mathbb{R}$. Without loss of generality, we use concise representation $\mathbf{Z} \in \mathbb{R}^{n \times k}$, $\mathbf{X} \in \mathbb{R}^{n \times m}$, and $\mathbf{A} \in \mathbb{R}^{n \times n}$ for the variables, which are conditional independencies.

Then, through maximizing the marginal likelihood of the observed adjacency matrix \mathbf{A} conditioned on \mathbf{X} , model parameters Θ can be obtained as follows:

$$\operatorname{argmax}_{\Theta} \log p_{\Theta}(\mathbf{A}|\mathbf{X}) = \log \int_{\mathbf{Z}} p_{\Theta}(\mathbf{A}, \mathbf{Z}|\mathbf{X}) d\mathbf{Z} \quad (1)$$

Here, $p(\mathbf{Z}|\mathbf{X})$ is a fixed prior distribution over the latent variable of every node associated with graph \mathbf{G} . However, equation (1) is intractable, hence, a variational posterior $q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})$ is introduced to approximate the true posterior, which can be turned into an optimization problem. Therefore, we can obtain a tractable evidence lower bound (ELBO) to the above objective with parameters Φ :

$$\begin{aligned} \log p_{\Theta}(\mathbf{A}|\mathbf{X}) &= \underbrace{\mathbb{E}_{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \left[\log \frac{p_{\Theta}(\mathbf{A}, \mathbf{Z}|\mathbf{X})}{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \right]}_{\text{ELBO}} \\ &+ \underbrace{\mathbb{E}_{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \left[\log \frac{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})}{p_{\Theta}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \right]}_{\text{KL}(q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})||p_{\Theta}(\mathbf{Z}|\mathbf{A}, \mathbf{X}))} \end{aligned} \quad (2)$$

Since the $\text{KL}(q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})||p_{\Theta}(\mathbf{Z}|\mathbf{A}, \mathbf{X})) \geq 0$, ie, it's non-negative, hence, equation (2) can be rewritten as follows:

$$\log p_{\Theta}(\mathbf{A}|\mathbf{X}) \geq \mathbb{E}_{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \left[\log \frac{p_{\Theta}(\mathbf{A}, \mathbf{Z}|\mathbf{X})}{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \right] \quad (3)$$

when the approximate posterior $q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})$ matches the true posterior $p_{\Theta}(\mathbf{Z}|\mathbf{A}, \mathbf{X})$, the lower bound is tight. Therefore, in order to obtain the best approximate of true posterior, we can maximize the (3) to optimize the parameters. Hence, the question is how to solve the $q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})$ (*encoder*) and $p_{\Theta}(\mathbf{A}|\mathbf{Z}, \mathbf{X})$ (*decoder*).

A. Encoder using semi-implicit variational inference

A typical approach of defining variational posterior is to use the mean-field (MF), then, the posterior can be written as $q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X}) \approx \prod_{i=1}^n q_{\Phi_i}(z_i|\mathbf{A}, \mathbf{X})$, where $q_{\Phi_i}(z_i|\mathbf{A}, \mathbf{X})$ is assumed to be Gaussian distribution. A high variance will occur, however, when the Gaussian distribution mismatching true posterior and it's restricting expressiveness of posterior following from the distributional assumptions [42]. To address the aforementioned problem, instead of using MF in previous work[52], we use a hierarchical semi-implicit framework to approximation the variational posterior. Specifically, in order to diffusion the uncertain between node and it's neighborhoods, first, we inject random noise at every layer of our encoder and concatenating it with nodes attributes, so that the output of GNN are random variables:

$$\mathbf{h}_p = \text{GNN}_p(\mathbf{A}, \text{CON}(\mathbf{X}, \epsilon_p, \mathbf{h}_{p-1})) \quad (4)$$

where $\epsilon_p \sim q_p(\epsilon)$ is Bernoulli distribution of N -dimensional noise. p denotes the number of layers, $\mathbf{h}_0 = \mathbf{X}$, CON denote CONCAT operation. With the above operation, the output of GNN $\mathbf{H} = \text{CON}(\{\mathbf{h}_i\}_{i=1}^p)$ is a random variable. Hence, it has been used to measure the uncertainty of GNN. Then, we use this random \mathbf{H} with \mathbf{X} and \mathbf{A} to learn the parameters of variational posterior:

$$\mu(\mathbf{A}, \mathbf{X}) = \text{GNN}_{\mu}(\text{CON}(\mathbf{X}, \mathbf{h}_p), \mathbf{A}) \quad (5a)$$

$$\sigma(\mathbf{A}, \mathbf{X}) = \text{GNN}_{\sigma}(\text{CON}(\mathbf{X}, \mathbf{h}_p), \mathbf{A}) \quad (5b)$$

CON denote CONCAT operation. Through (5a) and (5b), we can obtain the parameters of variational posterior. Attributed to the equation (4), which can propagate the uncertainly through the different layers, hence, when to learn the parameters μ and σ it can aggregation the neighbourhoods information of nodes, which is very important for aggregation nodes information. Finally, we can obtain the variational posterior as follows:

$$q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X}, \mu, \sigma) = \prod_{i=1}^N q_{\Phi}(\mathbf{z}_i|\mathbf{A}, \mathbf{X}, \mu_i, \sigma_i) \quad (6)$$

where $q_{\Phi}(\mathbf{z}_i|\mathbf{A}, \mathbf{X}, \mu_i, \sigma_i) = \mathcal{N}(\mu_i(\mathbf{A}, \mathbf{X}), \sigma_i(\mathbf{A}, \mathbf{X}))$.

In contrast to Graphite [52], which uses the mean-field approximation to define variational family and assumes the posterior is isotropic Gaussian, which restricts the expression of posterior distribution. Moreover, its output of GNN is determination, which can not propagate uncertainly between the nodes. Our approach can propagate uncertainly between the nodes with the random output of GNN and aggregate the neighborhood information naturally. This can increase the power of expressiveness and capture the complex posterior. Our approach can be seen in Fig.2.

B. Decoder using iterative manner

Following the [52] uses iterative manner for decoding, this paper also uses this approach. Specifically, given the latent variable \mathbf{Z} and feature matrix \mathbf{X} , an intermediate weighted $\tilde{\mathbf{A}}$ can be calculated through the inner-product of \mathbf{Z} .

$$\tilde{\mathbf{A}} = \frac{\mathbf{Z}\mathbf{Z}^T}{\|\mathbf{Z}\|^2} + \mathbf{1}\mathbf{1}^T \quad (7)$$

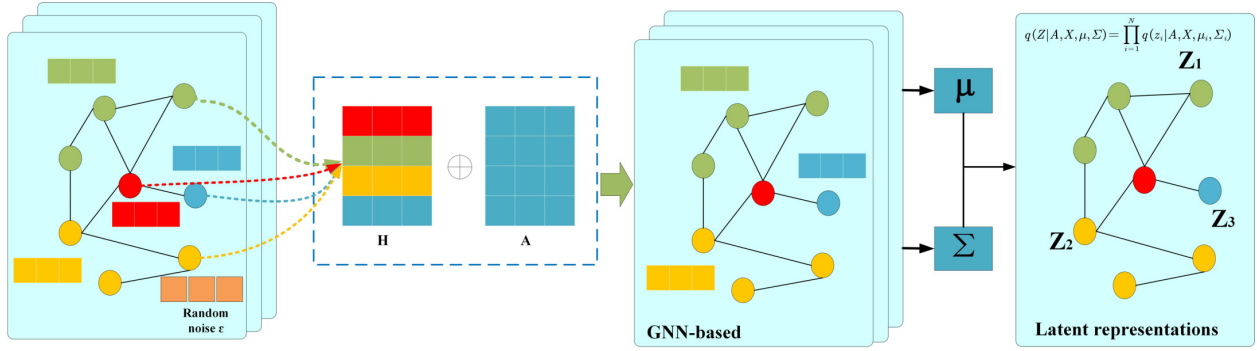


Fig. 2. The flowchart of our encoder. \mathbf{H} and \mathbf{A} denote the output of GNN and adjacency matrix respectively. In the input graph, we add random noise to every layer, which makes output H has randomness. \oplus denote CONCAT operation.

$$\mathbf{Z}^* = \text{GNN}_{\Theta}(\tilde{\mathbf{A}}, [\mathbf{Z}|\mathbf{X}]) \quad (8)$$

Hence, the decoder is given as follows:

$$p_{\Theta}(\mathbf{A}|\mathbf{Z}, \mathbf{X}) = \prod_{i=1}^n \prod_{j=1}^n p_{\Theta}^{(i,j)}(\mathbf{A}_{i,j}|\mathbf{Z}^*) \quad (9)$$

In order to representation learning of large graph more scalable, the matrix right multiplications is adopted for inference. A simplified graph propagation rule is adopted: $\mathbf{H}^{(l)} \leftarrow \zeta_l(\tilde{\mathbf{A}}\mathbf{H}^{(l-1)})$. For finally embedding of nodes, we use $\mathbf{Z}^f = (1 - \lambda)\mathbf{Z} + \lambda\mathbf{Z}^*$.

Towards to link prediction task, We can compute the probability that whether two edges are connected as follows:

$$p(\mathbf{A}_{i,j} = 1|z_i^f, z_j^f) = \delta(\mathbf{Z}^f \mathbf{Z}^{fT}) \quad (10)$$

where λ is hyper-parameter and δ is sigmoid function.

C. Model Inference

The first term on the right side of equation (2) is the ELBO, which is used to inference the model. Following the SIVI [24], we construct a hierarchical function: $\mathbf{Z} \sim q(\mathbf{Z}|\Psi)$, where $\Psi \sim q_{\Phi}(\Psi|\mathbf{X}, \mathbf{A})$. i.e., Ψ drawn from a distribution Therefore, we can rewritten the ELBO as follows:

$$\begin{aligned} \mathbb{L} &= \mathbb{E}_{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \left[\log \frac{p_{\Theta}(\mathbf{A}, \mathbf{Z}|\mathbf{X})}{q_{\Phi}(\mathbf{Z}|\mathbf{A}, \mathbf{X})} \right] \\ &= -\mathbf{KL}(\mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [q(\mathbf{Z}|\Psi)] \| p(\mathbf{Z})) \\ &\quad + \mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [\mathbb{E}_{\mathbf{Z} \sim q(\mathbf{Z}|\Psi)} \log p(\mathbf{A}|\mathbf{Z})]. \end{aligned} \quad (11)$$

Based on the [24]'s first theorem, we have:

$$\begin{aligned} &\mathbf{KL}(\mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [q(\mathbf{Z}|\Psi)] \| p(\mathbf{Z})) \\ &\leq \mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} \mathbf{KL}(q(\mathbf{Z}|\Psi) \| p(\mathbf{Z})). \end{aligned} \quad (12)$$

Hence, equation (11) can be written as follows:

$$\begin{aligned} \mathbb{L} &= -\mathbf{KL}(\mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [q(\mathbf{Z}|\Psi)] \| p(\mathbf{Z})) \\ &\quad + \mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [\mathbb{E}_{\mathbf{Z} \sim q(\mathbf{Z}|\Psi)} \log p(\mathbf{A}|\mathbf{Z})] \\ &\geq \mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [\mathbf{KL}(q(\mathbf{Z}|\Psi) \| p(\mathbf{Z}))] \\ &\quad + \mathbb{E}_{\Psi \sim q_{\Phi}(\Psi|\mathbf{A}, \mathbf{X})} [\mathbb{E}_{\mathbf{Z} \sim q(\mathbf{Z}|\Psi)} \log p(\mathbf{A}|\mathbf{Z})] = \mathbb{L}^* \end{aligned} \quad (13)$$

Directly optimizing \mathbb{L}^* could engender our method degenerates to the vanilla VGAE, duo to could lead to a point mass density as $q_{\Phi}(\Psi|\mathbf{X}, \mathbf{A})$. Therefore, in order to prevent this degeneracy, add a regularization term to \mathbb{L}^* . Assume

that S samples are derived from $q_{\Phi}(\Psi|\mathbf{X}, \mathbf{A})$, which denotes $\{\Psi^{(i)}\}_{i=1}^S$. Hence, we can define \mathcal{L}_S as follows:

$$\mathcal{L}_S = \mathbb{E}_{\Psi, \Psi^{(1)}, \dots, \Psi^{(S)} \sim q_{\Phi}(\Psi|\mathbf{X}, \mathbf{A})} [\mathbf{KL}(q(\mathbf{A}|\Psi) \| \hat{\mathbf{h}}_S(\mathbf{Z}))] \quad (14)$$

where,

$$\hat{\mathbf{h}}_S(\mathbf{Z}) = \frac{\mathbf{q}_{\Phi}(\Psi|\mathbf{A}, \mathbf{X}) + \sum_{s=1}^S \mathbf{q}_{\Phi}(\Psi^{(s)}|\mathbf{A}, \mathbf{X})}{S + 1}$$

Finally, the ELBO can be described as follows:

$$\mathcal{L} = \mathbb{L}^* + \mathcal{L}_S \quad (15)$$

Our method is described in Algorithm 1.

Algorithm 1 The algorithm of our method.

INPUT: \mathbf{A}, \mathbf{X} .

Initializing Θ and Φ .

Sample $\epsilon_p \sim q_p(\epsilon)$ where $q_p(\epsilon)$ is Bernoulli distribution.

for iteration $t = 1, 2, \dots, n$ **do**

Computing \mathbf{h}_P according to equation (4).

Computing μ and σ according to equation (5a) and (5b) respectively.

Computing $q(\mathbf{Z}|\mathbf{X}, \mathbf{A})$ according to equation (6).

Computing $p(\mathbf{A}|\mathbf{X}, \mathbf{Z})$ according to equation (9).

Update Θ and Φ by maximizing \mathcal{L} in (14).

return Θ, Φ .

end for

OUTPUT: Model Parameter Θ .

For link prediction task, computing \mathbf{Z}^* according to equation (8), then, $\mathbf{Z}^f = (1 - \lambda)\mathbf{Z} + \lambda\mathbf{Z}^*$ is used to computing $p(\mathbf{A}_{i,j} = 1|z_i^f, z_j^f) = \delta(\mathbf{Z}^f \mathbf{Z}^{fT})$.

IV. EXPERIMENTS

A. Datasets

We evaluate our method on 3 citation networks, i.e., Cora, Citeseer, and Pubmed with paper as nodes and citations as edges [53]. Furthermore, four different datasets without features are used to validate our method, i.e., NS: a collaborative

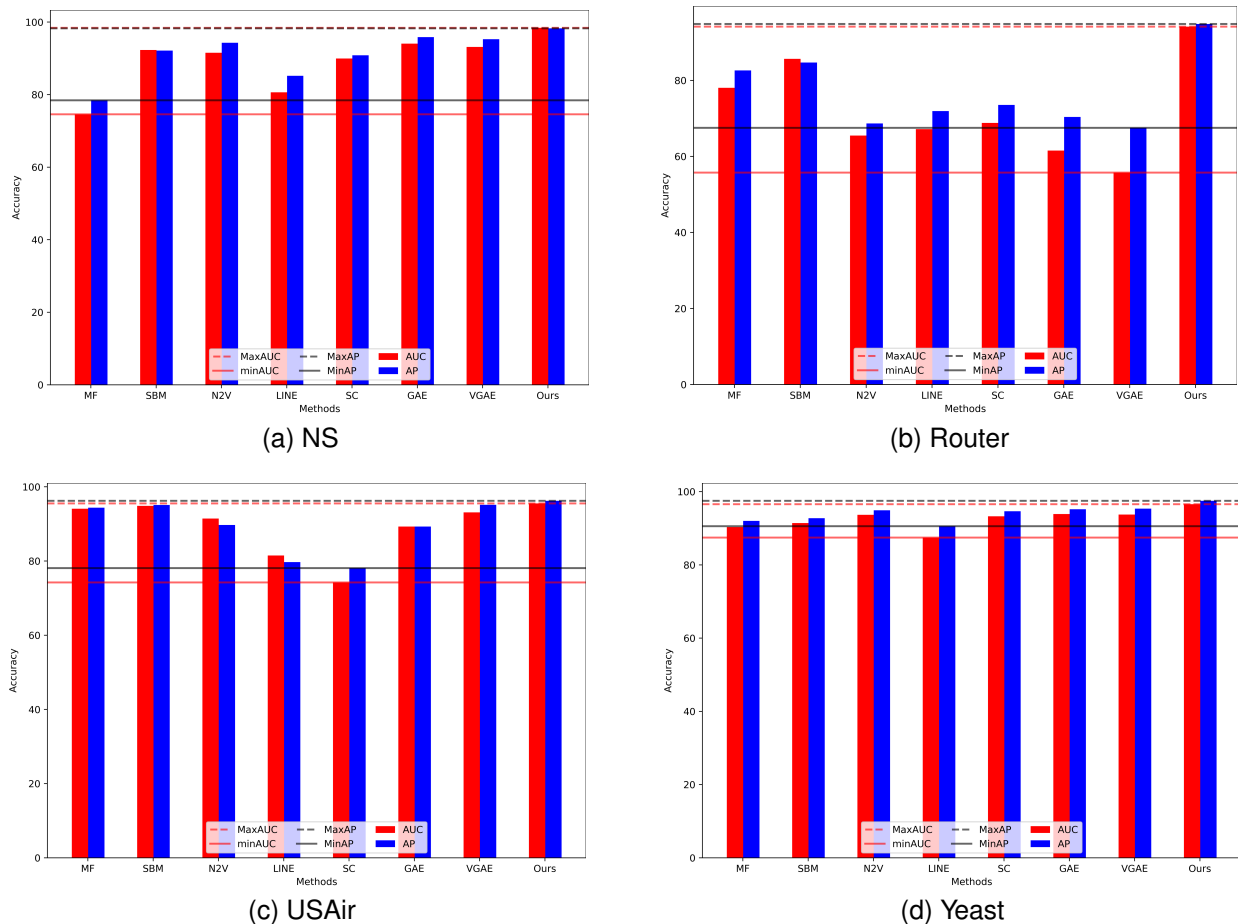


Fig. 3. AUC and AP of link prediction on the four different datasets. (a) NS: Collaboration Network. (b) Router: Internet Network. (c) USAir: Transportation Network. (d) Yeast: Protein Network. The red dashed and solid lines represent the maximum and minimum values of AUC. The black dashed and solid lines represent the maximum and minimum values of AP.

TABLE I
DESCRIPTION OF THE DATASETS DETAILS. '-' INDICATES THAT THE CHARACTERISTIC IS NOT AVAILABLE IN THE CORRESPONDING DATASET

	Nodes	Edges	Node feature	Labels
Cora	2708	5429	1433	7
Citeseer	3327	4732	3703	6
Pubmed	19717	44338	500	3
NS	1589	2742	-	-
USAir	332	2126	-	-
Router	5022	6258	-	-
Yeast	2375	11693	-	-

network of network science researchers, including 1589 nodes and 2742 edges [55], Router: a router internet with 5022 nodes and 6258 edges [56], USAir: US Airlines network, including 332 nodes and 2126 edges [57], Yeast: a protein-protein interaction (PPI) network in yeast, which has 2375 nodes and 11693 edges [58]. The detailed characteristics of the datasets are summarized in TABLE I.

B. Training Configurations

We evaluate the performance of our approach for the link prediction task. All GNN models in the experiments were implemented using GCN [10]. The link prediction task is to

predict whether an edge exists between two nodes [21]. We experiment on two different types of datasets: with and without features. The datasets was split into 5%, 10% and remaining for validation, testing and training respectively as done in [22]. We run our model for 3500 epochs with a learning rate of 0.0005 and training using Adam optimizer. The latent space dimensional is 16. The dimensional of Bernoulli noise ϵ is 64 and 32 for features and featureless respectively.

We organized the experiments on an experimental machine with Intel Xeon(R) CPU E7-8867 v4 @2.00GHz*80, GPU NVIDIA GTX 2080Ti, MEMORY 47.0GiB, Disk 698.4GB. Experimenting with implementation on the GPU version using TensorFlow [54].

C. Baselines And Metric

We compared our approach with some similar methods: Spectral Clustering (SC) [59], DeepWalk (DW) [60], Graphite [52], VGAE [22], GAE [22]. The model was evaluated by Area Under the ROC Curve (AUC) and Average Precision (AP). Particular, SC and DW do not provide the ability to merge node features when learning embeddings, hence they both validate on featureless datasets.

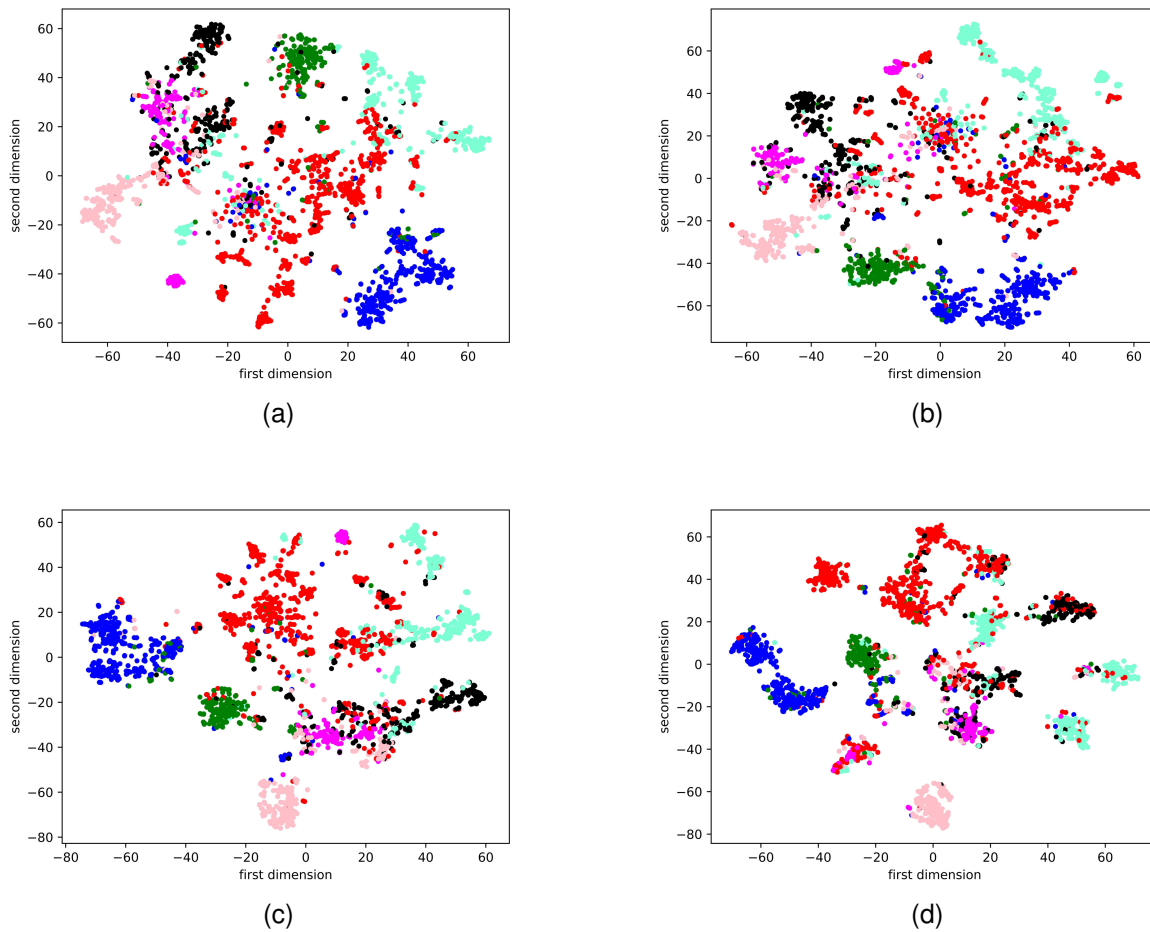


Fig. 4. TSNE embeddings of the latent feature vectors for the Cora dataset. Colors denote labels. (a) VAE. (b) VGAE. (c) Graphite. (d) Ours.

TABLE II
AREA UNDER THE ROC CURVE (AUC) FOR LINK PREDICTION WITH NODE FEATURES. THE BEST RESULTS ARE BOLDED. ‘-’ DENOTE NO NODE FEATURES

	Cora	Citeseer	Pubmed
SC	-	-	-
DW	-	-	-
GAE	91.05	89.99	92.33
VGAE	91.37	90.05	84.64
Graphite	94.50	97.63	95.96
Ours	94.84	98.00	96.89

TABLE III
AVERAGE PRECISION (AP) FOR LINK PREDICTION WITH FEATURES. THE BEST RESULTS ARE BOLDED. ‘-’ DENOTE NO NODE FEATURES.

	Cora	Citeseer	Pubmed
SC	-	-	-
DW	-	-	-
GAE	92.62	91.00	92.61
VGAE	92.26	91.00	86.02
Graphite	95.37	97.41	96.15
Ours	95.37	97.80	97.00

D. Results

1) **With nodes features:** In the first place, we conduct our method on 3 datasets with node features. The results are illustrated in TABLE II and TABLE III. We can discover at the table, our approach outperforms others. Compared with VGAE, which uses simple Gaussian assumptions, our method is more efficient than it in Citeseer and Pubmed datasets with a large margin, which demonstrates our model has powerful expression ability for the complex graph. In particular, our method outperforms VGAE by 10% on the Pubmed dataset.

The reason behind this is our method can capture the complex posterior and propagate uncertainty between nodes and their neighbourhoods, enhancing expressive ability.

2) **Without nodes features:** We execute our method on node featureless datasets. The results are shown in TABLE IV and TABLE V. As we can see in the table, our method achieved excellent results. Compared with GAE and VGAE, our method improves up to nearly 10% with AUC for Citeseer and Pubmed datasets. Besides, we demonstrate our approach on the four different datasets: NS, Router, USAir, Yeast. The results can be shown in Fig. 3. From Fig. 3 we can discover,

TABLE IV
AREA UNDER THE ROC CURVE (AUC) FOR LINK PREDICTION WITHOUT FEATURES. THE BEST RESULTS ARE BOLDED.

	Cora	Citeseer	Pubmed
SC	84.6	80.5	84.2
DW	83.1	80.5	84.4
GAE	85.92	78.34	85.36
VGAE	84.65	79.04	85.13
Graphite	88.58	85.57	95.36
Ours	91.14	91.30	95.01

TABLE V
AVERAGE PRECISION (AP) FOR LINK PREDICTION WITHOUT FEATURES. THE BEST RESULTS ARE BOLDED.

	Cora	Citeseer	Pubmed
SC	88.5	85.0	87.8
DW	85.0	83.6	84.1
GAE	88.74	83.4	88.65
VGAE	87.37	83.12	88.49
Graphite	90.09	85.14	94.32
Ours	92.84	92.89	96.00

our method obtained state-of-the-art results in four datasets. Especially, our method is more effective than LINE and SC in USAir dataset, 20% higher in the AUC and AP. For the Yeast dataset, our approach procures the best result, compared with LINE, our method is nearly seven percentage points higher than the line in AP, ten percentage points higher than the line in AUC. For NS, Router, and Yeast datasets, reference the dash and solid lines, compared to other methods, our method greatly improves the results in terms of both AUC and AP. The results prove that our method is more effective for both sparse and dense graphs.

3) **Visualization Display:** We visualize the embeddings learned by our method using 2-D TSNE for the Cora dataset. As is shown in Fig. 4. As we can see, Our method has a better clustering effect with more compactness between nodes. For example, for the red label, compared with other methods, our method makes clustering more compact. And the division between each class is more obvious.

E. Complexity Analysis

1) **Computational Complexity Analysis:** In our method, the highest complexity is the operation of inner products of potentially dense matrices \mathbf{Z} in Equation (7) (i.e. \mathbf{ZZ}^T). In order to computationally efficient, in our method, we apply a simplified graph propagation rule: $\mathbf{H}^{(l)} \leftarrow \zeta_l(\tilde{\mathbf{A}}\mathbf{H}^{(l-1)})$. Instead of directly compute inner product of \mathbf{Z} , we use associativity property of matrix multiplications. If d_l and d_{l-1} denote dimensional of layers of $\mathbf{H}^{(l)}$ and $\mathbf{H}^{(l-1)}$ respectively, the computational complexity is given by $\mathcal{O}(nkd_{l-1} + nd_{l-1}d_l)$, where k is the dimension of the per-node latent vectors \mathbf{z}_i used to define $\tilde{\mathbf{A}}$, n is the number of nodes.

2) **Time Cost:** For the analysis of the real-world graph dataset Cora on a single NVIDIA GTX 2080Ti GPU node, it

took 12.8, 20.8, and 30.5 seconds for Graphite, VGAE, and Our method with 100 epochs, respectively. For the analysis of the small real-world graph dataset USAir on a same GPU node, it took 4.7, 4.8, and 15.8 seconds for Graphite, VGAE, and Our method with 100 epochs, respectively.

V. CONCLUSION

In this paper, we proposed a novel GNN model based on semi-implicit variational inference. Our method uses a hierarchical frame to construct the model, which can obtain a tractable posterior inference. Specifically, in the encoder step, differs from the traditional method, which assumes the posterior as Gaussian distribution, which restrict expressiveness of posterior, hence, in this paper, we design a hierarchical semi-implicit variational posterior to approximate the true posterior. Contributed to this variational posterior and the hierarchical architecture between GNN layers, our method can capture complex posterior and propagate uncertainly between nodes, which is very essential for information aggregation. We prove our method procure outperformance results on citation networks and four different scenarios datasets. In the future, we will be using more simple approaches to approximate complex posterior, such as normalization flow.

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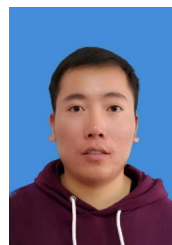
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