Pranav Nerurkar Madhav Chandane ® SUNIL BHIRUD ®

EXPLORING CONVOLUTIONAL AUTO-ENCODERS FOR REPRESENTATION LEARNING ON NETWORKS

Abstract

A multitude of important real-world or synthetic systems possess network structures. Extending learning techniques such as neural networks to process such non-Euclidean data is therefore an important direction for machine learning research. However, this domain has received comparatively low levels of attention until very recently. There is no straight-forward application of machine learning to network data, as machine learning tools are designed for i.i.d data, simple Euclidean data, or grids. To address this challenge, the technical focus of this dissertation is on the use of graph neural networks for network representation learning (NRL); i.e., learning the vector representations of nodes in networks. Learning the vector embeddings of graph-structured data is similar to embedding complex data into low-dimensional geometries. After the embedding process is completed, the drawbacks associated with graph-structured data are overcome. The current inquiry proposes two deep-learning auto-encoder-based approaches for generating node embeddings. The drawbacks in such existing auto-encoder approaches as shallow architectures and excessive parameters are tackled in the proposed architectures by using fully convolutional layers. Extensive experiments are performed on publicly available benchmark network datasets to highlight the validity of this approach.

Keywords

network representation learning, deep learning, graph convolutional neural networks

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1. Introduction

"We will never understand complex systems unless we develop a deep understanding of the networks behind them" – Albert Laszlo Barabasi [2]. In the scientific literature, different models have been developed to generate efficient representations and visualizations of data for its analysis [7]. However, the advantage of networks in the representation of data is that they provide a general language for describing and modeling complex systems [8]. Hence, networks have become ubiquitous across various scientific disciplines and are being used to represent many real or artificial systems [12]; for instance, internet, transportation systems, social networking websites, biological networks, etc. [18,19]. In modern times, the task of network-representation models has become increasingly difficult, as they must incorporate additional information of systems such as attribute data (side information), heterogeneous entities, high dimensionality, etc. To resolve these issues, network representation learning (NRL) frameworks are used [10,11]. The function of NRL frameworks is to learn a complex non-linear mapping function that embeds networks from their original (high) dimension to a latent (low) dimension.

Four families of the NRL techniques presented in the literature are adjacency-preserving methods [1,21,23,24,26,30], multi-hop distance-preserving methods, neighborhood overlap-preserving methods, and random walk occurrence-preserving methods [17, 22, 25, 28]. Given a set of nodes $x_1, ..., x_n$ and its pairwise relationships $r(x_i, x_j)$, these NRL techniques seek to map objects x_i to vectors $x_i \in \mathbb{R}^d$ such that $||x_i - x_j|| \sim r(x_i, x_j)$. These pairwise relationships correspond to the notion of "similarity" between the nodes in the network, such as adjacency, neighborhood overlap, random walk co-occurrence, second- or higher-order proximity, etc. [3, 9]. An important step of these NRL techniques is to feature engineering as a suitable "similarity" measure. This is the key drawback of such techniques.

Deep learning has impacted a wide range of domains such as image processing, machine translation, signal processing, etc. The availability of data, computation power, and fast optimization techniques have made it possible to stack multiple hidden layers that learn features in data-driven way – directly from raw data, thus eliminating the need for feature engineering [32]. However, there is no straight-forward application of deep learning to network data, as these tools are designed for i.i.d data, simple Euclidean data, or grids. Until very recently, this domain has received comparatively low levels of attention [27]. As a multitude of important real-world or synthetic systems possess a network structure: citation networks, internet, protein-protein interactions, brain connector data, and social networking websites. Extending machine-learning techniques to process non-Euclidean data is an important research direction [10].

In the literature, deep-learning architectures are based on graph convolutional neural (GCN) networks or graph auto-encoder (GAE) networks. GCN and its variants focus on information aggregation, whereas GAE models focus on inferring a latent geometry in the data. Most existing architectures have two to three stacked

layers. Stacking additional layers results in a model's performance dropping dramatically [29,33]. The success of deep learning lies in deep neural architectures; therefore, the current inquiry proposes two auto-encoder architectures that modify existing techniques like graph auto-encoders (GAE) and graph variational auto-encoders (GVAE) by replacing the fully connected layers with 1D convolutional ones.

The focus is to increase the depth of these architectures while maintaining computational and storage efficiency, a fixed number of parameters (independent of input network size), localization, and applicability to inductive problems. The rest of the paper is organized as follows: Section 2 provides the background of the research and literature review; Section 3 describes the mathematical model of the proposed approaches; and Section 4 describes the methodology of the experiments. The paper is concluded in Section 5.

2. Related Work

2.1. Background of Graph Convolutional Neural (GCN) Networks [13]

With V as the node set and E as the dyad set, graph G(V, E) has binary adjacency matrix $A \in \mathbb{R}^{|V|*|V|}$ and node feature matrix $X \in \mathbb{R}^{|V|*m}$. The node feature matrix is formed by horizontally stacking the m-dimension feature vector of each node. GCN has the following steps:

- 1. Convolutional filter \hat{A} is constructed from new adjacency matrix $\tilde{A} = A + I$ and new degree matrix $\tilde{D} = D + I$ by operation $\hat{A} = \tilde{D}^{\frac{-1}{2}} \tilde{A} \tilde{D}^{\frac{-1}{2}}$
- 2. A propagation rule for the graph convolution layer is defined as $H^{(1)} = \sigma(\hat{A}H^{(0)}\theta^{(0)})$. $H^{(1)}$ is the matrix of activations of the first layer, $H^{(0)} = X$, $\theta^{(0)}$ is a trainable weight matrix of layer 0, and σ is a non-linear activation function.
- 3. The convoluted representations of vertices $\hat{A}H^{(0)}$ are fed into a standard fully connected layer as shown in Figure 1.

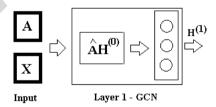


Figure 1. Graph Convolution Neural Network

2.2. Deep Learning Network Representative Learning Frameworks

Several attempts have been made to overcome the drawbacks of GCN-based architectures for representation learning. Kipf et al. proposed a variant of an existing

GCN-based architecture that uses two layers of feature aggregation so that higherorder dependencies can be captured in the vector embedding of the node. The inputs are adjacency matrix A and node feature matrix X. In the absence of the node feature matrix, the input can be a matrix with one-hot encoded representations of the nodes. The output of the model is the R^d vector embeddings of the nodes of the input graph [13]. One of the negative effects of graph convolutions is that they pushthe encoded representations of neighboring nodes closer to each other [16]. Theoretically, this would mean that the nodes of a graph would converge to a single point in latent space with multiple convolutions.

The framework proposed by M. Defferrard et al. samples a node neighborhood to form sub-graphs using a pooling operation [4]. Feature aggregation is performed on the sub-graphs for generating the final hidden representations of the nodes. The representations are calculated on the sub-graphs instead of on the original graph. This makes the framework more suitable for graph classification than representation learning. The graph auto-encoder with GCN proposed by T. Kipf et al. takes adjacency matrix A of the entire graph and node feature matrix X as its input. It outputs reconstructed adjacency matrix \hat{A} and node-embedding matrix Z. The loss function during learning is reconstruction loss, which is calculated as $L = E_{q(Z|X,A)}[log_p(A|Z)]$. An inner product function is used as decoder $\hat{A} = \sigma(ZZ^T)$. As the encoder function minimizes the loss due to reconstruction error, the representations of the nodes preserve first-order proximity. Global graph features are not considered in such node representations.

The graph spatial-temporal networks with GCN proposed by B. Yu $et\ al.$ [31] use GCN to learn the representations of the nodes to capture the spatial dependency of the nodes and 1-D CNN to capture the temporal dependency. The framework takes the adjacency matrix at time t and node feature matrix at time t as its input. As temporal information is not available in the case of static graphs, this framework is unsuitable for NRL on static graphs.

The encoder of the graph variational auto-encoder proposed by T. Kipf et al. [14] infers latent variables Z by sampling from a normal distribution. The μ, σ^2 of this normal distribution is generated through a two-layer GCN as $\mu = GCN_{\mu}(X,A), \sigma^2 = GCN_{\sigma}(X,A), GCN(X,A) = \hat{A}ReLU(\hat{A}X\theta^{(0)})\theta^{(1)}$. Then, for each input x_i , $q(Z_i|X,A) = N(Z_i|\mu_i, diag(\sigma_i^2))$. The decoder uses the same method as the inner product function of GAEs to obtain reconstructed adjacency matrix $A' = \sigma(Z.Z^T)$. The loss function has two parts: first, the construction loss to ensure a reconstructed adjacency matrix is similar to the input one. The second part is KL-divergence to ensure that the distribution of the latent variable is close to a normal distribution. The graph-VAE model depends on the GCN; hence, the model's performance is stagnant with multiple stacked layers due to the fully connected layers.

$$L = E_{q(Z|X,A)}[logp(X|Z)] - KL(q(Z|X,A)||p(Z)) \tag{1} \label{eq:logp}$$

where $p(Z) = \prod_i p(z_i) = \prod_i N(z_i|0,I)$

Drawbacks of existing architectures:

- To integrate global graph structural information, a much deeper GCN architecture would be required
- GCN-based architectures cause a dramatic drop in model performance when stacked together [16].
- Excessive parameter due to fully connected layers in encoder

3. Deep convolutional graph auto-encoder

Figure 2 provides the architecture of the encoder of the GAE with the use of the 1D convolutional (FC) layer in place of the fully connected (dense) layer. These FC encoder blocks are stacked to obtain the encoder of the GAE. The inner product decoder is kept the same.

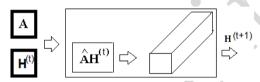


Figure 2. Architecture of convolutional graph auto-encoder

Similar to the 1D convolutional graph auto-encoder, an FC encoder block is used to substitute the fully connected layer in the GVAE as shown in Figure 3.

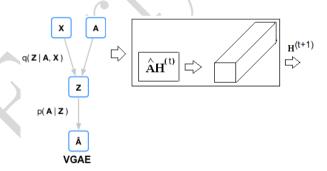


Figure 3. Architecture of graph variational auto-encoder with fully convolutional layer

3.1. Deep convolutional Variational Auto-encoders

The two models proposed in Sections 3 and 3.1 are trained to learn the embeddings using the procedure given in Algorithm 1.

The FC auto-encoder architectures are given in Table 1 for the BlogCatalog, CiteSeer, Cora, Flickr, Protein, and Wiki datasets.

Algorithm 1: Model Training and Fitting

Data: X, AResult: Z

- 1 Initialization of encoder weights θ and bias b;
- 2 Configure model to minimize Loss in Eq. 2.2;
- 3 Add activation function: rectified linear unit to encoder;
- 4 Create inner product decoder layer function;
- 5 Split data as 75:25 into train and test sets;
- 6 Set epochs, batch size;
- 7 Optimize the loss function L using gradient descent;

| Layer name | \mathbf{BlogC} | Cora | CiteSeer | Flickr | Protein | Wiki |
|-------------|------------------|-----------|----------|------------|-----------|-----------|
| X | 5198*8189 | 2709*1434 | 914*3703 | 7577*12047 | 3891*50 | 4778*40 |
| A | 5198*5198 | 2709*2709 | 914*914 | 7577*7577 | 3891*3891 | 4778*4778 |
| conv_encode | 1x64 | 1x64 | 1x64 | 1x64 | 1x4 | 1x4 |
| stride | 1 | 1 | 1 | 1 | 1 | 1 |
| padding | valid | valid | valid | valid | valid | valid |
| conv_encode | 1x32 | 1*64 | 1x64 | 1x64 | 1*3 | 1x3 |
| stride | 1 | 3 | 2 | 1 | 1 | 1 |
| padding | valid | valid | valid | valid | valid | valid |
| conv_encode | 1x32 | 1*64 | 1*32 | 1x32 | _ | _ |
| stride | 1 | 2 | 2 | 1 | _ | _ |
| padding | valid | valid | valid | valid | _ | _ |
| conv_encode | 1x32 | 1x32 | 1*32 | 1x32 | _ | _ |
| stride | 1 | 1 | 1 | 1 | _ | _ |
| padding | valid | valid | valid | valid | _ | _ |
| conv_encode | 1x32 | 1*64 | _ | 1x32 | _ | _ |
| stride | 1 | 1 | 1 | 1 | _ | _ |
| padding | valid | valid | valid | valid | _ | _ |
| conv_encode | 1*32 | 1*64 | _ | 1x32 | _ | _ |
| stride | 1 | 2 | _ | 2 | _ | _ |
| padding | valid | valid | _ | valid | _ | _ |
| conv_encode | _ | 1*32 | _ | _ | _ | _ |
| stride | _ | 2 | _ | _ | _ | _ |
| padding | _ | valid | _ | _ | _ | _ |
| conv_encode | _ | 1*32 | _ | _ | _ | _ |
| stride | _ | 1 | _ | _ | _ | _ |
| padding | _ | valid | _ | _ | _ | - |

The auto-encoder of the BlogCatalog dataset has six FC encoder blocks, the encoder for the CiteSeer dataset has four FC encoder blocks, the encoder for Cora has eight FC encoder blocks, the encoder of the Flickr dataset has five FC encoder

blocks, the encoder for the Protein dataset has two FC encoder blocks, and the encoder for Wiki has two FC encoder blocks.

3.2. Relationship of Auto-encoders with Laplacian smoothing

Consider a curve with points p_i, p_{i-1}, p_{i+1} .. as shown in Figure 4. When laplacian smoothing is applied to the graph, it results in each point getting closer to the weighted average of its neighbors as given in Eqs. 2 and 3.

$$p_i \leftarrow p_i + \frac{1}{2}L(p_i) \tag{2}$$

$$L(p_i) = \frac{p_{i+1} + P_{i-1}}{2} = p_i \tag{3}$$

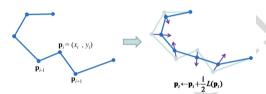


Figure 4. Example of laplacian smoothing where curve is shown before and after smoothing operation is applied. Image extracted from [16]

For a graph G = (V, E) with adjacency matrix A such that $a_{ij} = 1, |i - j| = 1$, and degree matrix $D = diag(d_1, d_2, ..., d_n)$, the smoothing operation can be rewritten as Eqs. 4 and 5:

$$P \leftarrow (I - \frac{1}{2}L_{rw})P \tag{4}$$

where:

$$P = \begin{bmatrix} x_1 & y_1 \\ x_2 & y_2 \\ \vdots & \vdots \\ x_n & y_n \end{bmatrix} \quad L_{\text{rw}} = \begin{bmatrix} 1 & -1 \\ -\frac{1}{2} & 1 & -\frac{1}{2} \\ & \ddots & \ddots & \ddots \\ & & -\frac{1}{2} & 1 & -\frac{1}{2} \\ & & & -1 & 1 \end{bmatrix}$$
 (5)

Matrix L=D-A is the normalized graph laplacian and has two versions $-L_{sym}=D^{-1/2}LD^{-1/2}$ and $L_{rw}=D^{-1}L$. The laplacian-smoothing operation is $P\leftarrow (I-\gamma L_{rw})P$, and $0\leq \gamma \leq 1$ controls the strength of the smoothing. Setting $\gamma=0$, the laplacian smoothing becomes equivalent to the non-linear mapping function (identity function) being learned by the auto-encoders. The laplacian smoothing computes the local average of each vertex as its new representation. Since vertices in

,

the same cluster tend to be densely connected, the smoothing makes their features similar, which makes the subsequent representation learning task-amenable.

4. Experiments

All of the datasets are publicly available at [15], and the details of the construction of the features for each data set are mentioned at the source.

4.1. Datasets

CiteSeer – there are 3312 scientific publications as nodes connected by the directed edge to a node if they cite it. There are 4732 links in the network. The feature vector of each node has 3704 features [20].

Cora – this is a citation network of 2708 nodes and 5429 links. The feature vector of each node has 1434 features [11].

BlogCatalog – this is a friendship network of 10,312 and 333,983 links. The feature vector of each node has 8189 features [10].

Flickr-Net – a "follower-following" network of 7575 users with 239,738 links. The feature vector of each node has 12,047 features [10].

Protein-Net – an interaction network of 3890 proteins with 37,845 edge links. The feature vector of each node has 50 features [10].

Wiki-Net – a citation network of Wikipedia articles with 4777 nodes and 54,810 links. The feature vector of each node has 40 features [10].

4.2. Performance metrics:

4.2.1. Separation index:

computed based on the distances for each point to the closest point not in the same cluster. The separation index is then the mean. A lower value indicates that the clusters are well separated.

4.2.2. Widest within-cluster gap:

the largest link in a within-cluster minimum spanning tree is computed. A larger value indicates good clustering.

4.2.3. Average silhouette width:

a measure of how similar an object is to its own cluster as compared to other clusters. The silhouette ranges from -1 to +1, where a high value indicates that the object is well-matched to its own cluster and poorly matched to neighboring clusters.

4.2.4. Average distance between clusters:

For two clusters r, s, the average distance between clusters should be large enough to indicate well-separated clusters as given by Eq. 6.

$$L(r,s) = \frac{1}{n_r * n_s} \sum_{i=1}^{n_r} \sum_{i=1}^{n_s} D(x_{ri}, x_{sj})$$
 (6)

4.2.5. Dunn index:

A lower Dunn index indicates better clustering. $\delta(C_i, C_j)$ is the inter-cluster distance metric between clusters C_i and C_j , Δ_k is the maximum within cluster variance as given by Equation 7.

$$DI_m = \frac{\min_{1 \le i < j \le m} \delta(C_i, C_j)}{\max_{1 \le k \le m} \Delta_k}$$
 (7)

4.3. Results and discussions

As can be seen in Figure 5, the loss decreases exponentially with the number of epochs in first two models, with a steeper decline in the VAE-FC model. Comparatively, the sparse auto-encoder has a high bias and cannot effectively learn the representations for the data-points in the training set.

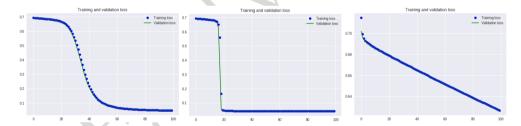


Figure 5. Training and cross-validation loss (binary cross-entropy loss) on CiteSeer dataset for graph auto-encoder with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

The loss seen in Figure 6 decreases exponentially with the number of epochs in first two models, with a steeper decline in the VAE-FC model. Comparatively, the sparse auto-encoder has a high bias and cannot effectively learn the representations for the data-points in the training set. The application of deep auto-encoders on the graph is equivalent to the application of multiple laplacian-smoothing operations on the graph. Each hidden layer computes a new latent representation of the data-point that is at a greater proximity to its neighbors. The resultant effect of the operations is that the encoder learns the representations that preserve a Euclidean proximity between the data-points in the latent space.

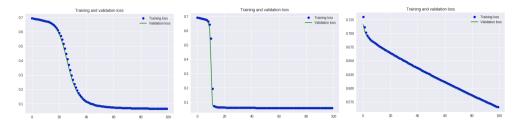


Figure 6. Training and cross-validation loss (binary cross-entropy loss) on Cora dataset for graph auto-encoder with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

The loss seen in Figure 7 decreases exponentially with the number of epochs in the first two models, with a steeper decline in the VAE-FC model. Comparatively, the sparse auto-encoder has a high bias and cannot effectively learn the representations for the data-points in the training set. The VAE-FC model computes the representations of the data-points with added constraints of minimizing the Kullback-Leibler divergence between the encoder's distribution over the representation's $q_{\theta}(z|x)$ and the distribution of the the latent representations of the data-point's $p(z) \sim \mathcal{N}(0,1)$. This leads to the representation vectors being normally distributed and organized into spherical clusters.

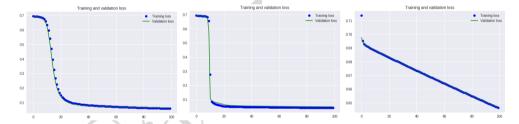


Figure 7. Training and cross-validation loss (binary cross-entropy loss) on BlogCatalog dataset for graph auto-encoder with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

The plot of the training and cross-validation loss vs. the epochs in Figure 8 for a sparsity constraint encoder reveals a high-bias and low-variance model. The sparsity constraints are responsible for the regularization of the model; this further exacerbates the high bias problem. The model fails to effectively capture the representations for the data-points of the network. To reduce the bias, the GAE-FC and VAE-FC models introduce additional parameters and more hidden layers. Both of these models are able to generalize to unseen data-points. The loss decreases exponentially with the epochs.

The feature vector of the Protein social network has \mathbb{R}^{50} . The small feature set allows for effective representation learning with sparsity constraint auto-encoders.

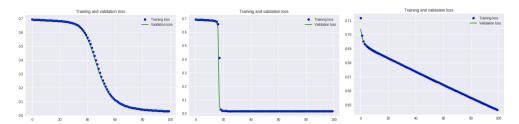


Figure 8. Training and cross-validation loss (binary cross-entropy loss) on Flickr dataset for graph auto-encoders with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

The loss with the epochs reduces exponentially for all three models (as can be seen in Fig. 9). The advantage of VAE-FC over the other models is the constraint of minimizing the Kullback-Leibler divergence between the encoder's distribution over representations $q_{\theta}(z|x)$ and the distribution of the latent representations of the datapoint's $p(z) \sim \mathcal{N}(0,1)$. This leads to the representation vectors being normally distributed and organized into spherical clusters.

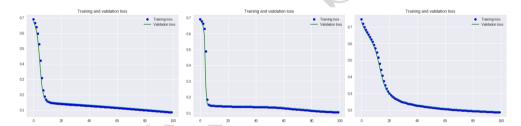


Figure 9. Training and cross-validation loss (binary cross-entropy loss) on Protein dataset for graph auto-encoders with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

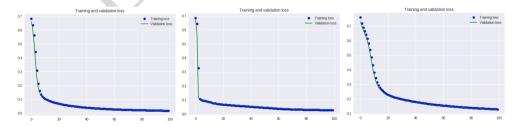


Figure 10. Training and cross-validation loss (binary cross-entropy loss) on Wikipedia dataset for graph auto-encoders with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6]

The feature vector of the Protein social network has \mathbb{R}^{40} . The three models learn representations of the data-points, and the training and cross-validation loss are minimal (as can be seen in Fig. 10). The encoders are able to generalize training parameters over unseen samples in the cross-validation data.

Table 2 describes the trainable parameters of the three neural encoder architectures. Sparsity constraint encoders have a lower count of parameters and are more suitable for smaller datasets. For larger datasets, the GAE-FC and VAE-FC deepencoder models would be preferred.

| Description | Sparse Enc | GAE-FC | VAE-FC |
|-------------------|------------|-----------|-----------|
| Parameters | 532,317 | 2,125,469 | 6,205,263 |
| Training Time per | 8 | 13 | 25 |
| epoch (in sec) | | | |

The three encoders were utilized for exploratory data analysis, and the performance of these models was evaluated on three datasets (see Figs. 11, 12). Hierarchical clustering (AGNES) was used to cluster the representation vectors. Internal cluster validation measures reveal that GAE-FC and VAE-FC obtain better results as compared to the simple encoder or sparsity constraint auto-encoders.

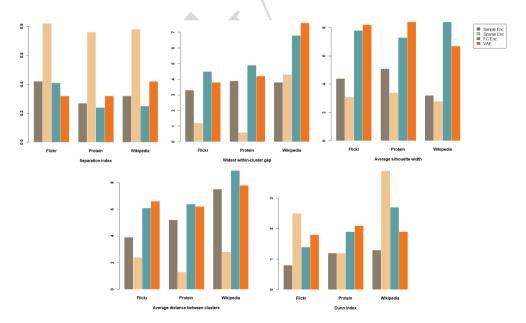


Figure 11. Comparison of performance of graph auto-encoders with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5,6] on datasets

The representations learned by the auto-encoders were clustered using unsupervised clustering – AGNES. It was observed that the graph auto-encoders with fully convolutional layers (GAE-FC) and the deep variational auto-encoder with fully convolutional layer (VAE-FC) learned the latent variable model for its input data performed better.

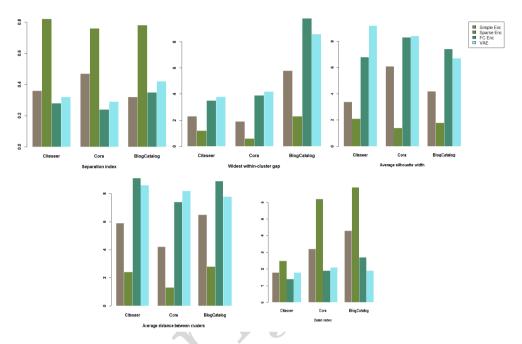


Figure 12. Comparison of performance of graph auto-encoders with fully convolutional layer (GAE-FC), deep variational auto-encoder with fully convolutional layer (VAE-FC), and sparsity constraint auto-encoder [5, 6] on datasets

5. Conclusion

With the advent of the digital transformation of our society, the generation of data is occurring at an exponential rate. Network models for the representation of data have become popular in scientific domains due to their ease in interpretation as well as analysis. In spite of such considerable advantages in research, networks or graphs are also associated with disadvantages such as the lack of independent observations and the existence of coupling between data-points. Representation learning is proposed as a suitable solution for such drawbacks. However, the existing techniques for representation learning are based on heuristics and, hence, cannot capture complex non-linear patterns in the data-points and encode them into representations.

The neural network-based models in the literature focus on semi-supervised learning on graphs. However, as labeled datasets are costly and unavailable in real-world

systems, the current investigation was motivated to utilize unsupervised neural architectures such as encoders. It was observed that graph auto-encoders with a fully convolutional layer (GAE-FC) and a deep variational auto-encoder with a fully convolutional layer (VAE-FC) performed significantly better at representation learning when compared to the sparsity constraint architecture. Such architectures were also able to learn representations for data-points not observed in the training phase. This feature makes them more suitable for deployment on real-world systems than heuristic-based shallow techniques. The operation of aggregation and transformation performed in the hidden layer of a neural network is equivalent to a laplacian-smoothing operation. Hence, deep variational auto-encoders with fully convolutional layer (VAE-FC) architectures learn representations such as data-points associated with similar characteristics in the high-dimension space are located in the proximity of the latent space. The experiments performed on the network datasets experimentally validated this theoretical insight.

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Affiliations

Pranav Nerurkar

Computer Engineering and IT Department, VJTI, Mumbai, India, ORCID ID: https://orcid.org/0000-0002-9100-6437

Madhav Chandane

ORCID ID: https://orcid.org/0000-0002-2872-4647

Sunil Bhirud

ORCID ID: https://orcid.org/0000-0002-9100-6437

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