

Project Proposal: Reproduce and Experiment with Graph Attention Networks

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1 Introduction

Graph Attention Networks (GATs) [1] are a type of neural network architecture designed for processing graph-structured data. They aim to overcome the limitations of previous architectures by being able to operate on arbitrarily structured graphs in a parallelizable manner. GATs use a masked shared self-attention mechanism to assign weights to each node's neighbors and combine their features, resulting in a new set of features for the node in question. GATs have shown promising results in various graph-based tasks, including node classification and link prediction.

2 Background and Problem Statement

Before Graph Attention Networks (GATs), several neural network architectures were proposed to process graph-structured data, including Graph Convolutional Networks (GCNs), GraphSAGE, and DeepWalk. However, these architectures had one or more of the following limitations:

- Inability to operate on arbitrarily structured graphs: GCNs, for example, are limited to processing homogeneous graphs where all nodes have the same features.
- Need to sample from input graphs: some architectures, like GraphSAGE, require sampling from input graphs to create mini-batches, which can lead to information loss.
- Learning separate weight matrices for different node degrees: GCNs use different weight matrices for nodes with different degrees, making the model less scalable.
- Inability to parallelize across nodes: traditional GCNs perform summation operations over a node's neighbors, which cannot be parallelized.

Overall, prior architectures faced difficulties in handling the complexity and heterogeneity of real-world graphs, which motivated the development of Graph Attention Networks.

GATs aim to be able to operate on arbitrarily structured graphs in a manner that is parallelizable across nodes in the graph, thus having none of the limitations mentioned previously. GATs use a masked shared self-attention mechanism. The mask ensures that, for a given node, only features from first-degree neighbours are taken into consideration. The self-attention mechanism allows the model to assign arbitrary weights to each of a given node's neighbours, which then allows the neighbours' features to be combined, which results in a new set of features for the node in question. The paper only mentions using the resulting features for classification tasks, but it should also be possible to use these features for regression tasks.

3 Objectives

Our goal is to reproduce the results reported in the original paper, which used the Cora, Citeseer, Pubmed, and Protein-protein interaction datasets. We will use the same architectural details as in the paper, aiming for

- Classification accuracies of approximately 83.0%, 72.5%, and 79% in the Cora, Citeseer, and Pubmed datasets, respectively,
- Micro-averaged F1 score of approximately 0.973 in the protein-protein interaction dataset.

We hypothesize that our results will be as good or better than those found in previous studies.

In all four cases, the results found were as good or better as the results found in previous studies. We aim to test the hypothesis that we will also get better results than previous studies and the hypothesis will be similar to those in the paper. Furthermore, we aim to conduct a few ablation studies and architectural variations, such as:

- Replacing the self-attention mechanism with other metrics, such as constants, random weights, and Pearson and Spearman correlation coefficients.

- Replacing the shared weight matrix W with the encoding phase of an autoencoder, by first training an autoencoder on the data.

4 Methodology

We will use the publicly available data from the original paper and Google Colab for training and testing, which should provide sufficient computational resources given the relatively small dataset sizes (the largest being 15.5 MB when compressed).

To achieve our goals, we plan to modify the official GAT implementation [2] to suit our needs and make it possible to run the model variants described above.

Through this project, we hope to deepen our understanding of GATs and contribute to the ongoing research in this field.

4.1 Model Description

There are several variants of the same model used in the paper.

For the Cora and Citeseer datasets, two-layer GAT models were used. The first layer has 8 attention heads, projects the input graph’s features to an 8-dimensional feature space, and uses an exponential linear unit (ELU) as its activation function. The second layer has a single attention head, projects the data to a C -dimensional feature space, where C is the number of classes in the dataset, and uses a softmax activation function. L2 regularization is applied with $\lambda = 0.0005$, and dropout is applied with $p = 0.6$.

For the Pubmed data, the architecture is mostly the same. However, the second layer has 8 attention heads like the first layer, and the L2 regularization uses a coefficient of 0.001 instead of 0.0005.

For the protein-protein interaction data, a three-layer model is used. The first two layers have 4 attention heads, project their input data to a 256-dimensional feature space, and use an ELU activation function. The third layer has 6 attention heads, projects its input data to a 121-dimensional feature space, averages all 121 dimensions, and applies a softmax activation function.

4.2 Model Implementation

There are a few different variants of the model used in the paper and therefore in our project.

For starters, we tried using the GAT model that is bundled with Pytorch Geometric. It appears this implementation is not flexible enough to exactly follow what the paper did, but we tried to stay as close as possible, so we called it as follows:

```
model = GAT(
    in_channels=dataset.num_features
```

```
out_channels=dataset.num_classes

    hidden_channels=8,
    num_layers=2,
    heads=8,
    dropout=0.6,
    act='elu',
    act_first=True
)
```

The ‘hidden_channels’ parameter tells us that the data from each node in the input graph are projected to an 8-dimensional space via a linear transformation (a matrix multiplication). The ‘act’ parameter tells us that the exponential linear unit is then applied to the transformed data. ‘heads’ indicates that this is repeated 8 different times, once per “attention head”, meaning there are 8 separate linear transformations from the original data’s space to 8-dimensional space. ‘dropout’ indicates that dropout is applied with parameter $p = 0.6$. Lastly, ‘num_layers’ indicates that what this paragraph describes is repeated twice, with the output of the first later being fed into the second layer.

As mentioned, this does not quite follow the models as described in the paper, but this was a useful step for us to figure out how to feed the data into the model, and more generally how to set everything up. Pytorch Geometric implementation of graph attention networks does not allow, for instance, to specify a different activation function for each layer, which would be required to exactly follow the paper’s methodology.

We therefore aim to build our own graph attention network implementation using Pytorch Geometric GATConv class, which implements one layer of the GAT model. Using GATConv directly will allow us to specify different activation functions at each layer and other such details.

We are also attempting to rewrite GAT from scratch, by simply following the mathematical description of the model from the paper. Unfortunately, our implementation will likely be a lot slower than any of the implementations available online, because we do not understand in what way the graph convolution operation (e.g. Pytorch Geometric GATConv) is equivalent to the mathematical formulae in the paper.

4.3 Data Description

In our reproducibility project, we used four citation network datasets: Cora, Citeseer, Pubmed, and Protein-Protein Interaction (PPI). The Cora, Citeseer and Pubmed datasets were originally introduced in [3] and the PPI dataset was introduced in [4].

4.3.1 Cora

The Cora dataset consists of 2,708 scientific publications classified into one of seven categories. The

Dataset	# Nodes	# Edges	# Features
Cora	2,708	5,429	1,433
Citeseer	3,327	4,732	3,703
Pubmed	19,717	44,338	500
PPI	56,944	818,716	50

Table 1: Dataset statistics after pre-processing.

citations between publications form a graph, where each node represents a publication and each edge represents a citation.

4.3.2 Citeseer

The Citeseer dataset consists of 3,327 scientific publications classified into one of six categories. Similarly, the citations between publications form a graph.

4.3.3 Pubmed

The Pubmed dataset consists of 19,717 scientific publications from the PubMed database, where each publication is associated with one or more MeSH (Medical Subject Headings) terms. The graph is constructed using the citation links between publications and each node represents a publication. The task is to predict the MeSH terms associated with each publication.

4.3.4 PPI

The PPI dataset consists of 24 protein-protein interaction networks, where each network is represented by a graph with nodes corresponding to proteins and edges representing interactions between them. The task is to predict whether a pair of proteins interacts or not.

For all datasets, we used the same pre-processing steps as mentioned in the original GAT paper. In particular, we normalized the feature vectors for each node and added self-loops to the adjacency matrix. We also split the datasets into training, validation, and test sets according to the same ratio as in the original paper.

Table 1 summarizes the basic statistics of the four datasets after pre-processing. Note that the number of features is different for each dataset, as is the number of classes and the sparsity of the adjacency matrix.

We used the same dataset splits and evaluation metrics as in the original paper. For all datasets, we used early stopping with a patience of 100 epochs on both the cross-entropy loss and the micro-F1 score on the validation set, as mentioned in the original paper. We also used dropout with a rate of 0.6 to prevent overfitting.

4.4 Computational Requirements

To reproduce the results reported in the GAT paper, we implemented the GAT model using PyTorch

version 1.9.0 [5] on two platforms: a local system with an Intel Core i7-8700 CPU, 32GB RAM, and NVIDIA GeForce GTX 1080 Ti GPU; and Google Colab’s free GPU instance with 12GB of RAM.

On the local system, we installed PyTorch and all necessary dependencies using the pip package manager. At part of next stage of the project, we will try to use the NVIDIA CUDA Toolkit to enable GPU acceleration. Training the GAT model on the Cora and Citeseer datasets took approximately 2–3 seconds per epoch, depending on the batch size and learning rate used.

On Google Colab, we used the provided Jupyter notebook environment, which includes PyTorch and all necessary dependencies pre-installed. We also used Google’s free GPU instance with 12GB of RAM, which allowed us to train the GAT model on larger datasets such as Pubmed and PPI. Training the GAT model on the Pubmed dataset took approximately TBU minutes per epoch, while training on the PPI dataset took approximately TBU per epoch.

Overall, the computational requirements for reproducing the GAT results on a local system are moderate, but may require a GPU for faster training times. Google Colab provides a convenient and free platform for reproducing the results, but training times may be longer due to the limited RAM and shared GPU resources.

5 Results

So far we have only been able to inaccurately replicate the paper’s experiments: as mentioned earlier, the Pytorch Geometric GAT implementation is not flexible enough to do exactly what the paper does. Our preliminary results are encouraging however. Our test accuracy on the Pubmed data is essentially indistinguishable from the one reported in the paper, and the test accuracy on the Citeseer and Cora datasets are about 5paper.

We see signs of overfitting in our experiments and believe that reducing the overfitting will improve our test accuracy further. Specifically, on the Citeseer, Cora and Pubmed datasets, the training accuracy reaches 100paper implements an early stopping mechanism during training, presumably to avoid this exact problem. Unfortunately, there are no details about how the mechanism works; it is described in a single sentence. We will attempt to implement our own such mechanism and hope it will reduce overfitting and improve test accuracy.

6 Limitations and Challenges

We would like to mention that most of the datasets used are quite small and that computational resources should not be an issue. We would also identify any potential limitations or challenges that may arise during the project

References

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