# Comparing Shallow and Deep Graph Models for Brain Network Analysis

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#### Introduction

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- Analyze different approaches for classifying brain networks
  - kernelized SVM<sup>1</sup>
  - message passing GNNs<sup>2</sup>
  - graph kernel GNNs<sup>3</sup>
- Suggest several methods to motivate further research in brain network analysis

<sup>&</sup>lt;sup>1</sup>Hofmann, Schölkopf, and Smola, "Kernel methods in machine learning", 2008

<sup>&</sup>lt;sup>2</sup>Cui et al., BrainGB: A Benchmark for Brain Network Analysis with Graph Neural Networks, 2022

<sup>&</sup>lt;sup>3</sup>Feng et al., "KerGNNs: Interpretable Graph Neural Networks with Graph Kernels", 2022

### Classification Task

- The standard graph classification task considers the problem of classifying graphs into two or more categories
- In this project, we perform binary classification on neuroimaging data to distinguish between negative and positive diagnoses

#### **Datasets**

- We are working with 2 datasets, one classifying HIV and the other classifying bipolar disorder
- Each dataset consists of:
  - diffusion tensor imaging (DTI) scans
  - functional magnetic resonance imaging (fMRI) scans
  - classification labels: positive diagnosis, negative diagnosis

#### **Datasets**

Preliminaries

- The DTI and fMRI brain scans of each patient i are represented as weighted adjacency matrices  $\mathbf{W}_i \in \mathbb{R}^{M \times M}$ 
  - The fMRI scans are considered to be more robust than DTI scans, so our experiments prioritize working with them
  - The fMRI datasets have been cleaned for us and consist of 70 (HIV) and 97 (bipolar disorder) patients
- Nodes in the brain network represent regions of interest (ROIs), and edge links between nodes indicate the strength of the connection between ROIs

# Threshold Rounding

Preliminaries

- We implement a rounding scheme to remove edge weights and sparsify the adjacency matrices
- We have:  $A_{ij} = \begin{cases} 1 & \text{if } A_{ij} \geq \alpha \\ 0 & \text{otherwise} \end{cases}$ , where  $A_{ij}$  is the ij-th entry of the adjacency matrix A and  $\alpha \in [0,1]$  is our rounding threshold

## Threshold Rounding

Preliminaries

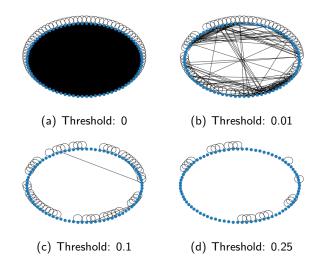


Figure 1: Effect of threshold rounding on network density.

## Graph Kernels

- Popular in graph-based learning because they can be computed implicitly (inner product)
- We compute graph kernel matrices using the *GraKel* Python package and plug them into SVM to perform classification
- Consider WL, WLOA, shortest path, and graphlet sampling kernels in experiments

## Graph Kernels

- Weisfeiler-Lehman subtree kernel is built on the Weisfeiler-Lehman graph isomorphism test<sup>4</sup> and is essentially a relabeling procedure
  - Computationally inexpensive, taking O(hm) time, where h is the number of iterations and m is the number of edges.
- WL optimal assignment kernel uses valid assignment theory to improve the performance of the WL subtree kernel<sup>5</sup>
  - Computed in linear time, taking O(|X| + |Y|) time, where X and Y are elements of  $[\mathcal{X}]^n$ .  $[\mathcal{X}]^n$  denotes the set of all *n*-element subsets of the set  $\mathcal{X}$ .

<sup>&</sup>lt;sup>4</sup>Weisfeiler and Lehman, "The reduction of a graph to canonical form and the algebra which appears therein", 1968

<sup>&</sup>lt;sup>5</sup>Kriege, Giscard, and Wilson, "On Valid Optimal Assignment Kernels and Applications to Graph Classification", 2016

## Graph Kernels

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- **Shortest path** kernel decomposes graphs into shortest paths and compares pairs of them<sup>6</sup>
  - Computationally expensive when number of n nodes is large, taking  $O(n^4)$  time
- **Graphlet sampling** kernel decomposes graphs into graphlets of k nodes and compares the number of matching graphlets between two graphs<sup>7</sup>
  - Computationally intractable for large k, taking  $O(n^k)$  time
  - Experiments show k=5 generally performs the best

<sup>&</sup>lt;sup>6</sup>Borgwardt and Kriegel, "Shortest-path kernels on graphs", 2005

<sup>&</sup>lt;sup>7</sup>Przulj, "Biological network comparison using graphlet degree distribution", 2007

## Support Vector Machines

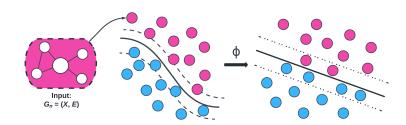


Figure 2: Overview of kernel SVM.

## **Graph Neural Networks**

- GNNs combine node features and graph structures to perform prediction tasks
- General framework:
  - computing the representation of each node
  - applying a pooling strategy to obtain the graph representation
  - multilayer perceptron (MLP) can be applied to make predictions

- We implement MPGNNs using the BrainGB Python package and focus on two types of MPGNNs:
  - Graph attention network (GAT) is a type of convolutional neural network that operates on graphs
  - Graph convolutional network (GCN) is a special case of GATs with attention fully determined by graph structure alone, without node features
- Conduct experiments using settings based on extensive studies from Cui et al (2022)

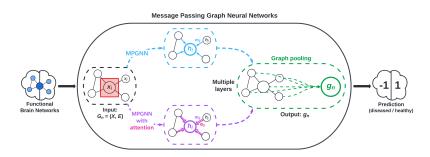


Figure 3: BrainGB framework. Adapted from Fig. 1 in Cui et al (2022). The node representation of node  $x_i$  is  $h_i$ , the message from node  $x_j$  to  $x_i$  is  $m_{ij}$ , and the attention weight from node  $x_j$  to  $x_i$  is  $a_{ij}$ .

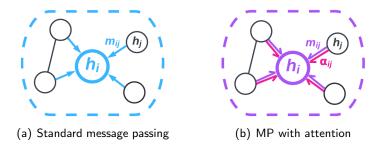


Figure 4: The message passing schemes in the BrainGB framework.

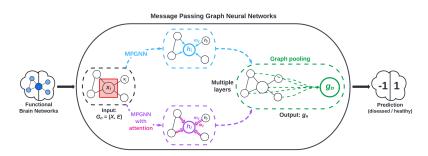


Figure 5: BrainGB framework. Adapted from Fig. 1 in Cui et al (2022). The output  $g_n$  is the pooled information that will be passed through a MLP to make the prediction.

#### Kernel GNNs

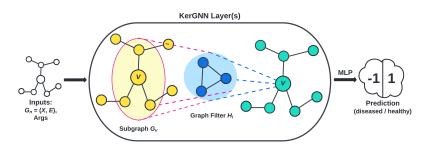


Figure 6: KerGNN framework. Adapted from Fig. 3 in Feng et al (2022).

#### Kernel GNNs

| Number of epochs      | 100; 150; 200; 250; 300; 350; 400; 450; 500               |  |
|-----------------------|---|--|
| Learning rate         | $10^{-2}$ ; $10^{-3}$ ; $10^{-4}$ ; $10^{-5}$ ; $10^{-6}$ |  |
| Dropout rate          | 0.1; 0.2; 0.3; 0.4; 0.5; 0.6; 0.7; 0.8; 0.9               |  |
| Nodes in graph filter | 2; 4; 6; 8; 10; 12; 14; 16; 18; 20                        |  |
| Subgraph size         | 5; 10; 15; 20   |  |
| k-hop neighborhood    | 1; 2; 3   |  |
| Max step of RW        | 1; 2; 3; 4; 5   |  |

Table 1: Hyperparameter Search Range

| Data | Method          | Accuracy          | F1                | AUC               |
|------|-----------------|-------------------|-------------------|-------------------|
| HIV  | WL-0.21         | $0.67_{\pm 0.17}$ | _                 | _                 |
|      | WLOA-0.21       | $0.65_{\pm 0.17}$ |                   | _                 |
|      | SP-0.01         | $0.66_{\pm 0.20}$ |                   | _                 |
|      | GS-0.03         | $0.66_{\pm0.18}$  |                   |                   |
|      | GCN-concat      | $0.64_{\pm 0.15}$ | $0.59_{\pm 0.20}$ | $0.77_{\pm 0.20}$ |
|      | GAT-concat      | $0.73_{\pm 0.16}$ | $0.71_{\pm 0.17}$ | $0.81_{\pm 0.19}$ |
|      | GCN-edge concat | $0.71_{\pm 0.11}$ | $0.69_{\pm 0.12}$ | $0.77_{\pm 0.17}$ |
|      | GAT-edge concat | $0.69_{\pm 0.18}$ | $0.67_{\pm 0.19}$ | $0.73_{\pm 0.24}$ |
|      | KerGNN          | $0.64_{\pm 0.19}$ | _                 | _                 |
| ВР   | WL-0.4          | $0.63_{\pm 0.19}$ |                   | _                 |
|      | WLOA-0.42       | $0.66_{\pm0.12}$  |                   | _                 |
|      | SP-0.02         | $0.64_{\pm0.12}$  |                   | _                 |
|      | GS-0.04         | $0.62_{\pm 0.15}$ |                   |                   |
|      | GCN-concat      | $0.53_{\pm 0.13}$ | $0.51_{\pm 0.14}$ | $0.54_{\pm 0.16}$ |
|      | GAT-concat      | $0.53_{\pm 0.13}$ | $0.50_{\pm 0.13}$ | $0.57_{\pm 0.19}$ |
|      | GCN-edge concat | $0.63_{\pm 0.12}$ | $0.61_{\pm 0.13}$ | $0.61_{\pm 0.17}$ |
|      | GAT-edge concat | $0.52_{\pm 0.17}$ | $0.51_{\pm 0.16}$ | $0.59_{\pm 0.19}$ |
|      | KerGNN          | $0.68_{\pm 0.16}$ | _                 | _                 |

#### Discussion

- Limited data (70 and 97 patients in each dataset)
- GNNs are usually shallow; deep GNNs are still an active area of research
- For brain networks, what kinds of graph structures are effective beyond the pairwise connections are still unknown

#### Discussion

Preliminaries

- Cui et al (2021)<sup>8</sup> notes HIV affects 2 sub-networks, while bipolar disorder only affects 1 sub-network
  - This may make accurate classification difficult
- Li et al (2020)<sup>9</sup> found utilizing multimodal neuroimaging (fMRI and MRI) improves SVM classification performance

<sup>&</sup>lt;sup>8</sup>Cui et al., "BrainNNExplainer: an interpretable graph neural network framework for brain network based disease analysis", 2021

<sup>&</sup>lt;sup>9</sup>Li et al., "Identification of bipolar disorder using a combination of multimodality magnetic resonance imaging and machine learning techniques", 2020

#### Future Work

- There are many graph kernels and GNNs that we hope are useful in the area of brain network analysis
- Some of these include: graph kernel neural networks<sup>10</sup> (GKNN), graph stochastic attention<sup>11</sup> (GSAT), *k*-dimensional GNNs<sup>12</sup> (*k*-GNN), message passing graph kernels<sup>13</sup> (MPGK), and motif convolutional networks<sup>14</sup> (MCN)

<sup>&</sup>lt;sup>10</sup>Cosmo et al., Graph Kernel Neural Networks, 2021

<sup>&</sup>lt;sup>11</sup>Miao, Liu, and Li, *Interpretable and Generalizable Graph Learning via Stochastic Attention Mechanism*, 2022

<sup>&</sup>lt;sup>12</sup>Morris et al., Weisfeiler and Leman Go Neural: Higher-order Graph Neural Networks, 2018

<sup>&</sup>lt;sup>13</sup>Nikolentzos and Vazirgiannis, *Message Passing Graph Kernels*, 2018
<sup>14</sup>Lee et al., *Higher-order Graph Convolutional Networks*, 2018

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