

Graph Classification via Reference Distribution Learning: Theory and Practice

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Motivation: Current global pooling methods suffer from information loss.

- Most of the global pooling methods are naive, often employing methods such as simple summation or averaging. These pooling methods collect only the first-order (statistics) information, leading to a loss of structural or semantic information.
- More sophisticated pooling operations retain more meaningful information, but still carry the inherent risk of information loss.

Model and Optimization

- GRDL is composed of two parts.

- f_G is a backbone GNN to transform each graph G_i with adjacency matrix $A_i \in \mathbb{R}^{n_i \times n_i}$ and node feature $X_i \in \mathbb{R}^{n_i \times d_0}$ to a node embedding matrix $H_i \in \mathbb{R}^{n_i \times d}$, which encodes the graph's information

$$H_i = f_G(G_i) = f_G(A_i, X_i),$$

- A reference layer f_D computes the similarity between each graph embedding H_i and reference distributions $\{D_1, \dots, D_K\}$

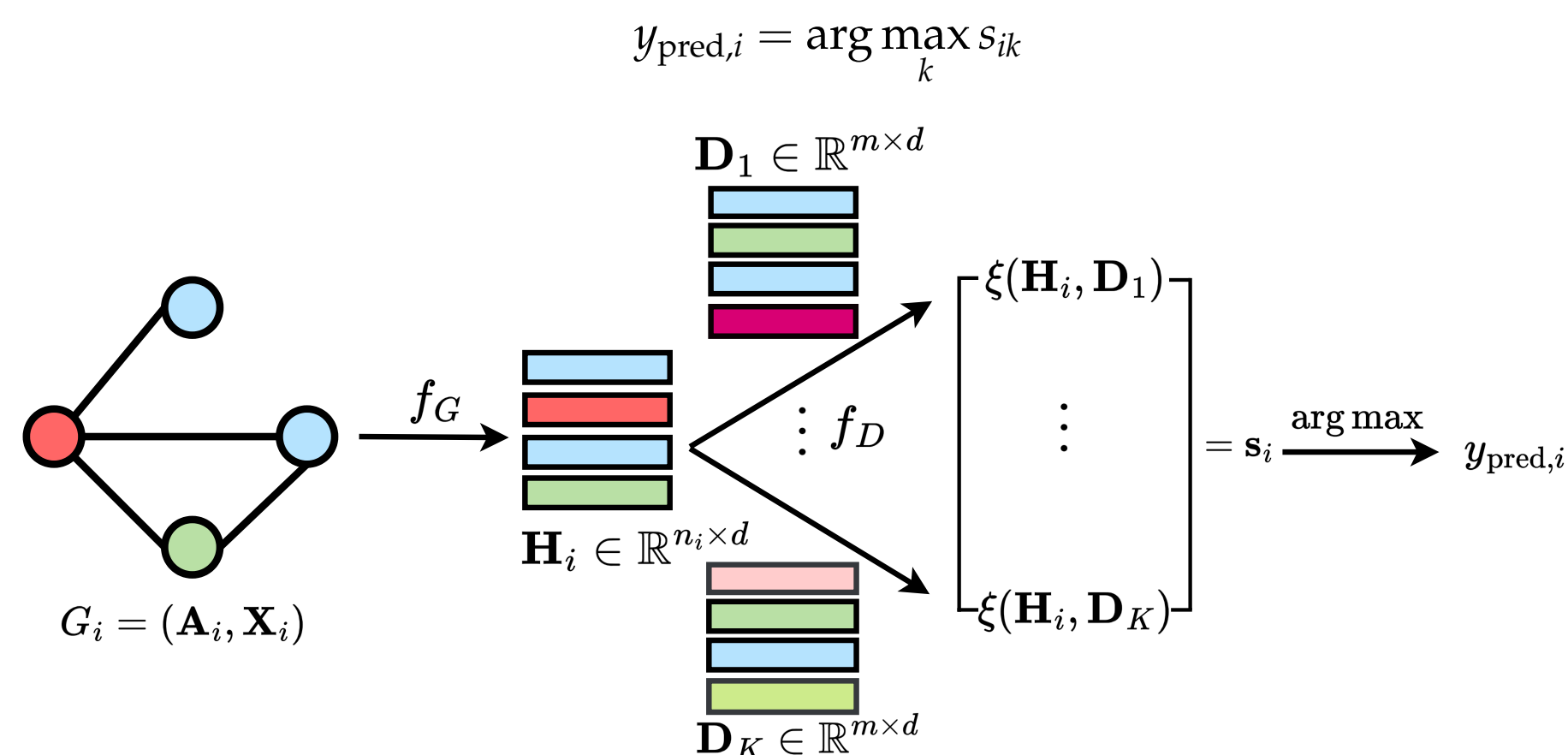
$$f_D(H_i) = [s_{i1}, s_{i2}, \dots, s_{iK}] = [\xi(H_i, D_1), \xi(H_i, D_2), \dots, \xi(H_i, D_K)]^\top \in \mathbb{R}^K$$

where $\xi(\cdot, \cdot)$ is a similarity measure between two distributions, and is chosen to be the negative squared maximum mean discrepancy (MMD) in our experiment:

$$\xi(H, D) = \frac{2}{mn} \sum_{i=1}^n \sum_{j=1}^m k(h_i, d_j) - \frac{1}{n^2} \sum_{i=1}^n \sum_{i'=1}^n k(h_i, h_{i'}) - \frac{1}{m^2} \sum_{j=1}^m \sum_{j'=1}^m k(d_j, d_{j'}),$$

$k(\cdot, \cdot)$ is chosen to be the Gaussian kernel $k(x, x') = \exp(-\theta \|x - x'\|_2^2)$.

- The graph is assigned the label of the reference that exhibits the highest similarity



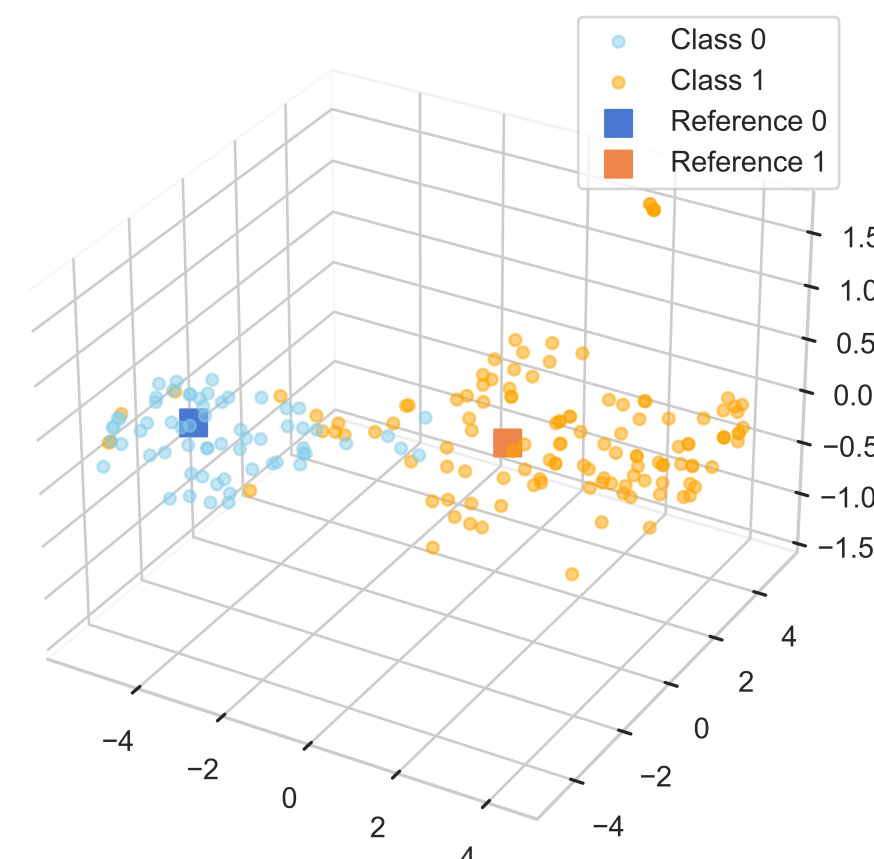
- Optimization problem of GRDL:

$$\min_{f_G, f_D} -\frac{1}{N} \sum_{i=1}^N \sum_{k=1}^K y_{ik} \log \frac{\exp(s_{ik})}{\sum_{j=1}^K \exp(s_{ij})} + \lambda \sum_k \sum_{k' \neq k} \xi(D_k, D_{k'})$$

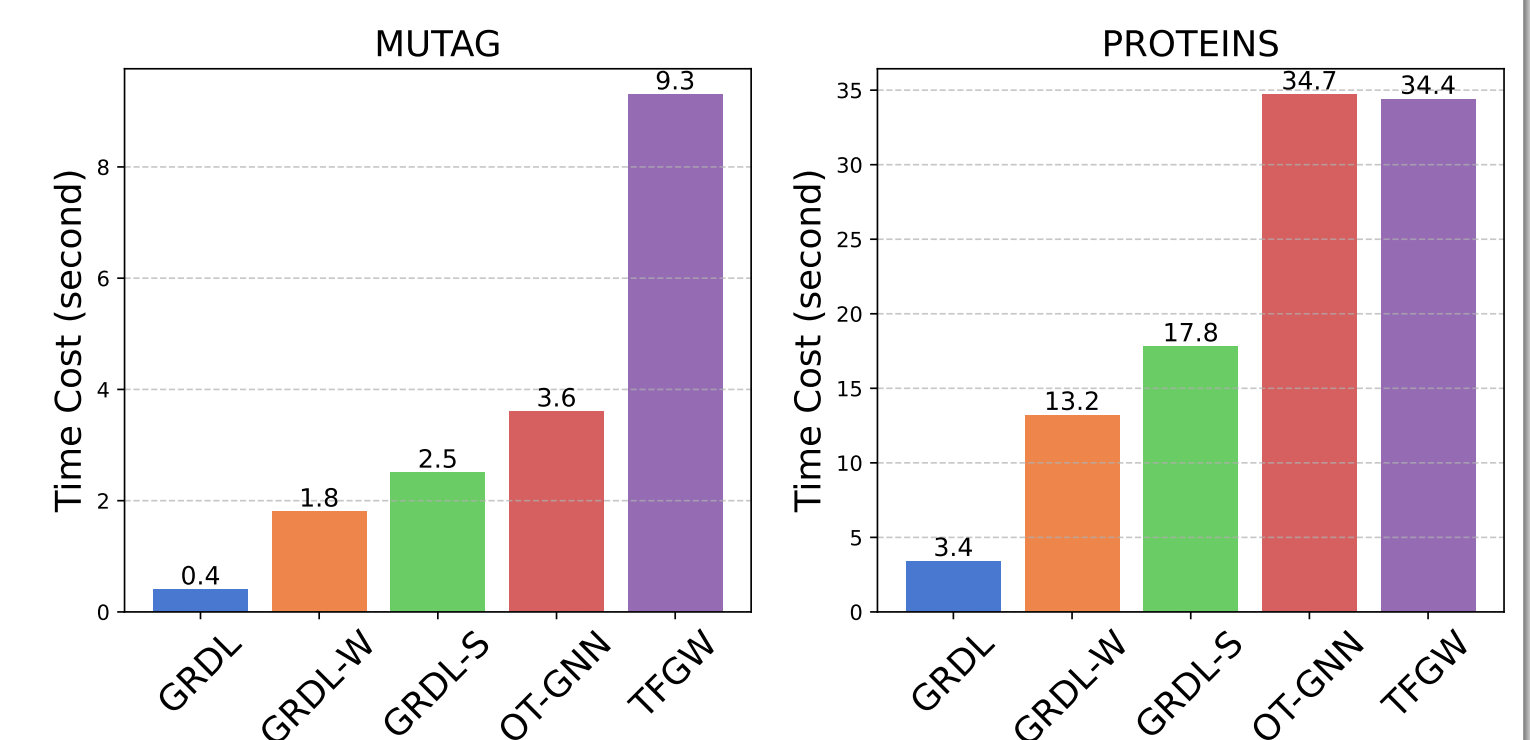
Experiments on Graph Dataset & Visualizations

| METHOD | DATASET | | | | | | | | AVERAGE |
|------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-------------|
| | MUTAG | PROTEINS | NCI1 | IMDB-B | IMDB-M | PTC-MR | BZR | COLLAB | |
| PATCHY-SAN | 92.6±4.2 | 75.1±3.3 | 76.9±2.3 | 62.9±3.9 | 45.9±2.5 | 60.0±4.8 | 85.6±3.7 | 73.1±2.7 | 71.5 |
| GIN | 89.4±5.6 | 76.2±2.8 | 82.2±0.8 | 64.3±3.1 | 50.9±1.7 | 64.6±7.0 | 82.6±3.5 | 79.3±1.7 | 73.6 |
| DroPgin | 90.4±7.0 | 76.9±4.3 | 81.9±2.5 | 66.3±4.5 | 51.6±3.2 | 66.3±8.6 | 77.8±2.6 | 80.1±2.8 | 73.9 |
| DIFFPOOL | 89.4±4.6 | 76.2±1.4 | 80.9±0.7 | 61.1±3.0 | 45.8±1.4 | 60.0±5.2 | 79.8±3.6 | 80.8±1.6 | 71.8 |
| SEP | 89.4±6.1 | 76.4±0.4 | 78.4±0.6 | 74.1±0.6 | 51.5±0.7 | 68.5±5.2 | 86.9±0.8 | 81.3±0.2 | 75.8 |
| GMT | 89.9±4.2 | 75.1±0.6 | 79.9±0.4 | 73.5±0.8 | 50.7±0.8 | 70.2±6.2 | 85.6±0.8 | 80.7±0.5 | 75.7 |
| MinCutPool | 90.6±4.6 | 74.7±0.5 | 74.3±0.9 | 72.7±0.8 | 51.0±0.7 | 68.3±4.4 | 87.2±1.0 | 80.9±0.3 | 75.0 |
| ASAP | 87.4±5.7 | 73.9±0.6 | 71.5±0.4 | 72.8±0.5 | 50.8±0.8 | 64.6±6.8 | 85.3±1.3 | 78.6±0.5 | 73.1 |
| WITopoPool | 89.4±5.4 | 80.0±3.2 | 79.9±1.3 | 72.6±1.8 | 52.9±0.8 | 64.6±6.8 | 87.8±2.4 | 80.1±1.6 | 75.9 |
| OT-GNN | 91.6±4.6 | 76.6±4.0 | 82.9±2.1 | 67.5±3.5 | 52.1±3.0 | 68.0±7.5 | 85.9±3.3 | 80.7±2.9 | 75.7 |
| WEGL | 91.0±3.4 | 73.7±1.9 | 75.5±1.4 | 66.4±2.1 | 50.3±1.0 | 66.2±6.9 | 84.4±4.6 | 79.6±0.5 | 73.4 |
| FGW - ADJ | 82.6±7.2 | 72.4±4.7 | 74.4±2.1 | 70.8±3.6 | 48.9±3.9 | 55.3±8.0 | 86.9±1.0 | 80.6±1.5 | 71.5 |
| FGW - SP | 84.4±7.3 | 74.3±3.3 | 72.8±1.5 | 65.0±4.7 | 47.8±3.8 | 55.5±7.0 | 86.9±1.0 | 77.8±2.4 | 70.6 |
| WL | 87.4±5.4 | 74.4±2.6 | 85.6±1.2 | 67.5±4.0 | 48.4±4.2 | 56.0±3.9 | 81.3±0.6 | 78.5±1.7 | 72.4 |
| WWL | 86.3±7.9 | 73.1±1.4 | 85.7±0.8 | 71.6±3.8 | 52.6±3.0 | 52.6±6.8 | 87.6±0.6 | 81.4±2.1 | 73.9 |
| SAT | 92.6±4.3 | 77.7±3.2 | 82.5±0.8 | 70.0±1.3 | 47.3±3.2 | 68.3±4.9 | 91.7±2.1 | 80.6±0.6 | 76.1 |
| GRAPHORMER | 89.6±6.2 | 76.3±2.7 | 78.6±2.1 | 70.3±0.9 | 48.9±2.0 | 71.4±5.2 | 85.3±2.3 | 80.3±1.3 | 75.1 |
| GRDL | 92.1±5.9 | 82.6±1.2 | 80.4±0.8 | 74.8±2.0 | 52.9±1.8 | 68.3±5.4 | 92.0±1.1 | 79.8±0.9 | 77.9 |
| GRDL-W | 90.8±4.6 | 82.1±0.9 | 80.9±0.8 | 72.2±3.1 | 53.1±0.9 | 68.5±3.2 | 90.6±1.5 | 80.4±1.1 | 77.3 |
| GRDL-S | 90.6±5.7 | 81.1±1.4 | 81.2±1.5 | 72.4±3.3 | 52.5±1.1 | 64.2±3.2 | 91.6±1.3 | 78.6±1.3 | 76.5 |

Table: Classification accuracy (%). Bold text indicates the top 3 mean accuracy.



(a) MUTAG node embedding and reference distributions.



(b) Time cost comparison.

Theoretical Results

Generalization bound and its implications (see our paper).