# Deep Graph Infomax







Given a graph G with N nodes:

```
Each node has a node feature vector x_i \in \mathbb{R}^F, and we denote X = \{x_1, x_2, ..., x_N\}
Adjacency matrix A \in \{0,1\}^{N \times N}
```

In the *l*-th layer of Graph Neural Network (GNN)<sup>[1]</sup>, to update **node** representation: [Encoder]

$$a_{i}^{(l)} = AGGRE^{(l)}\left(\left\{h_{j}^{(l-1)}: j \in \mathcal{N}(i)\right\}\right), \qquad h_{i}^{(l)} = COMBINE^{(l)}\left(h_{i}^{(l-1)}, a_{i}^{(l)}\right)$$

For example:

In GCN<sup>[2]</sup>, two operations are integrated:

$$h_{i}^{(l)} = ReLU\left(W \cdot MEAN\left\{h_{j}^{(l-1)}, \forall j \in \mathcal{N}(i) \cup \{i\}\right\}\right)$$

In GraphSAGE<sup>[3]</sup>:

$$a_i^{(l)} = MAX\left(\left\{ReLU\left(W_1 \cdot h_j^{(l-1)}\right), \forall j \in \mathcal{N}(i)\right\}\right), \qquad h_i^{(l)} = W_2 \cdot \left[h_i^{(l-1)}, a_i^{(l)}\right]$$

 $MAX(\cdot)$  here is element-wise max pooling.



Given a graph *G* with *N* nodes:

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```

After the *L*-th layer, we can use a **readout** function for **graph** representation (summary):

$$s = \mathcal{R}\left(\left\{h_i^{(L)} \middle| i \in G\right\}\right)$$

For example:

$$s = \frac{1}{N} \sum_{i \in G} h_i^{(L)}$$



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Basic assumption: some types of transformation retain the important information<sup>[4]</sup>.



For *n*-th graph, two transformed samples *i* and *j*:

$$l_n = -\log \frac{\exp(sim(z_{n,i}, z_{n,j})/\tau)}{\sum_{n'=1,n'\neq n}^{N} \exp(sim(z_{n,i}, z_{n',j})/\tau)}$$



Given a graph with *N* nodes:

Node feature  $x_i \in \mathbb{R}^F$ , and feature matrix  $X = \{x_1, x_2, ..., x_N\} \in \mathbb{R}^{N \times F}$ , adjacency matrix  $A \in \{0, 1\}^{N \times N}$ 

[Goal] An encoder  $\mathcal{E}: \mathbb{R}^{N \times F} \times \mathbb{R}^{N \times N} \to \mathbb{R}^{N \times F'}$  that encodes **node** with **high-level** representation:  $H = \mathcal{E}(X, A) = \{h_1, h_2, ..., h_N\}$ , where  $h_i \in \mathbb{R}^{F'}$ .

(Example: GCN or GraphSAGE can be considered as an encoder.)

A readout function  $\mathcal{R}: \mathbb{R}^{N \times F} \to \mathbb{R}^{F}$  summarizes the nodes representations to a **graph-level** representation:  $s = \mathcal{R}(\mathcal{E}(X, A))$ 

(Example: MEAN or SUM can be considered as a readout function.)

## **Deep Graph Infomax**



Original node feature matrix  $X = \{x_1, x_2, ..., x_N\}$ , adjacency matrix AAn encoder  $\mathcal{E}$  and new node representations:  $H = \mathcal{E}(X, A)$ A readout function  $\mathcal{R}$  results in a summary:  $s = \mathcal{R}(\mathcal{E}(X, A))$ 

A discriminator  $\mathcal{D}: \mathbb{R}^F \times \mathbb{R}^F \to \mathbb{R}$  assigns probability to node:  $\mathcal{D}(h_i, s)$ [To decide if the node should be contained within the summary] [Example: cosine similarity.  $\mathcal{D}(h_i, s) = \sigma(h_i^T W s)$  in this paper]

Negative samples for  $\mathcal{D}$ :  $\tilde{h}_i$  from an alternative graph  $\tilde{G} = (\tilde{X}, \tilde{A})$ 

 $\tilde{G} = (\tilde{X}, \tilde{A})$  can come from another graph in the multiple-graphs dataset. Or can be obtained by a corruption function  $\mathcal{C}: \mathbb{R}^{N \times F} \times \mathbb{R}^{N \times N} \to \mathbb{R}^{M \times F} \times \mathbb{R}^{M \times M}: (\tilde{X}, \tilde{A}) = \mathcal{C}(X, A)$ [Example:  $\tilde{A} = A$  and random row-wise shuffling X to get  $\tilde{X}$ ]

[C generates negative samples, while T in Contrastive Learning generates positive samples.]



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Metric: Mutual Information (MI), evaluated by the JSD MI estimator<sup>[5]</sup>:

$$\max MI(X, A; h_i) \approx \max \left[ \log (\mathcal{D}(h_i; X, A)) + \log \left( 1 - \mathcal{D}(\tilde{h}_j; X, A) \right) \right]$$

Approximate (X, A) by  $s = \mathcal{R}(\mathcal{E}(X, A))$ 

Objective: Noise Contrastive objective with BCE loss

$$\mathcal{L} = \frac{1}{N+M} \left( \sum_{i=1}^{N} \mathbb{E}_{(X,A)} \left[ \log \mathcal{D}(h_i, s) \right] + \sum_{j=1}^{M} \mathbb{E}_{(\tilde{X}, \tilde{A})} \left[ \log \left( 1 - \mathcal{D}(\tilde{h}_j, s) \right) \right] \right)$$

### **Deep Graph Infomax**



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- 1. Sample a negative example by  $(\tilde{X}, \tilde{A}) \sim C(X, A)$
- 2. Obtain  $h_i$ ,  $\tilde{h}_j$  by the encoder,  $H = \mathcal{E}(X, A)$  and  $\tilde{H} = \mathcal{E}(\tilde{X}, \tilde{A})$
- 3. Readout the input graph:  $s = \mathcal{R}(H)$
- 4. Update parameters of  $\mathcal{E}$ ,  $\mathcal{R}$  and  $\mathcal{D}$  by gradient descent to maximize  $\mathcal{L}$ .





Q1: We approximate (X, A) by  $s = \mathcal{R}(\mathcal{E}(X, A))$ , in which case they are the closest to each other? A1:  $\mathcal{R}$  is injective.

Given a probability distribution of graphs, p(X), we can draw  $\{X^{(k)}\}_{k=1}^{|X|}$  from p(X) uniformly, e.g.,  $p(X^{(k)}) = p(X^{(k)'})$ .  $s^{(k)} = \mathcal{R}(X^{(k)})$  is the summary of *k*-th graph with marginal distribution p(s).

The optimal classifier between the p(X, s) and p(X)p(s) has an error rate upper bounded by  $Err^* = \frac{1}{2}\sum_{k=1}^{|X|} p(s^{(k)})^2$ , and the upper bound is achieved when  $\mathcal{R}$  is injective.



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Define  $Q^{(k)} = \{ \mathbf{X}^{(j)} | \mathcal{R}(\mathbf{X}^{(j)}) = s^{(k)} \}$  contains all graphs being mapped to  $s^{(k)}$ . Sample  $(\mathbf{X}^{(k)}, s^{(k)})$  drawn from the product of marginals with probability  $p(\mathbf{X})p(s)$ :

$$p(s^{(k)})\sum_{s} p(\mathbf{X}^{(k)}, s) = p(s^{(k)})p(\mathbf{X}^{(k)}, s^{(k)}) = p(s^{(k)})p(\mathbf{X}^{(k)}|s^{(k)})p(s^{(k)}) = p(s^{(k)})^{2} \frac{p(\mathbf{X}^{(k)})}{\sum_{\mathbf{X}' \in \mathcal{Q}^{(k)}} p(\mathbf{X}')}$$

$$\begin{split} p\big(\boldsymbol{X}^{(k)}, \boldsymbol{s}^{(j)}\big) &= 0 \text{ when } j \neq k \text{ since } \mathcal{R} \text{ is deterministic.} \\ p\big(\boldsymbol{X}^{(k)}, \boldsymbol{s}^{(k)}\big) &= p\big(\boldsymbol{s}^{(k)}\big)p\big(\boldsymbol{X}^{(k)}|\boldsymbol{s}^{(k)}\big) \text{ from the definition of conditional probability.} \\ p\big(\boldsymbol{X}^{(k)}|\boldsymbol{s}^{(k)}\big) &= \frac{p(\boldsymbol{X}^{(k)})}{\sum_{\boldsymbol{X}' \in \mathcal{Q}^{(k)}} p(\boldsymbol{X}')} \text{ from the definition of } \mathcal{Q}^{(k)}. \end{split}$$



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Define  $Q^{(k)} = \{ \mathbf{X}^{(j)} | \mathcal{R}(\mathbf{X}^{(j)}) = s^{(k)} \}$  contains all graphs being mapped to  $s^{(k)}$ . Sample  $(\mathbf{X}^{(k)}, s^{(k)})$  drawn from the product of marginals with probability  $p(\mathbf{X})p(s)$ :

$$\frac{p(\mathbf{X}^{(k)})}{\sum_{\mathbf{X}' \in \mathcal{Q}^{(k)}} p(\mathbf{X}')} p(s^{(k)})^2 = \rho^{(k)} p(s^{(k)})^2 \le p(s^{(k)})^2$$

We have  $\rho^{(k)} \leq 1$ , since  $X^{(k)} \in Q^{(k)}$ . When  $Q^{(k)} = \{X^{(k)}\}$ , ( $\mathcal{R}$  is injective),  $\rho^{(k)} = 1$ .



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Define  $Q^{(k)} = \{ \mathbf{X}^{(j)} | \mathcal{R}(\mathbf{X}^{(j)}) = s^{(k)} \}$  contains all graphs being mapped to  $s^{(k)}$ . Sample  $(\mathbf{X}^{(k)}, s^{(k)})$  drawn from the joint  $p(\mathbf{X}, s)$ :

$$p(\mathbf{X}^{(k)}, s^{(k)}) = p(\mathbf{X}^{(k)}|s^{(k)})p(s^{(k)}) = \rho^{(k)}p(s^{(k)})$$

Compare it with previous results:

 $\rho^{(k)}p(s^{(k)}) \ge \rho^{(k)}p(s^{(k)})^2$ 

The optimal classifier always classifies samples to the joint for a lower error

$$Err \leq \frac{1}{2} \sum_{k=1}^{|X|} \rho^{(k)} p(s^{(k)})^2 \leq \frac{1}{2} \sum_{k=1}^{|X|} p(s^{(k)})^2$$

When  $\mathcal{R}$  is injective for all  $\mathbf{X}^{(k)}$ ,  $Err^* = \frac{1}{2} \sum_{k=1}^{|\mathbf{X}|} p(s^{(k)})^2$ 



Q2: If we assume that  $\mathcal{R}$  is injective and  $|s^*| = |\mathbf{X}|$ , what else can we know?

[|s| is the number of allowable states in s, and  $s^*$  is the optimal summary w.r.t the classification error.] A2:  $s^* = argmax_s MI(X; s)$ . In other words, minimizing the classification error is equivalent to maximizing the mutual information.

#### Proof

MI is invariant under invertible transforms.

Since  $\mathcal{R}$  is injective and  $|s^*| = |X|$ , we always can find an inverse function  $\mathcal{R}^{-1}$ .

For any *s*:

$$MI(\mathbf{X}; s) \leq H(\mathbf{X}) = MI(\mathbf{X}; \mathbf{X}) = MI(\mathbf{X}; \mathcal{R}(\mathbf{X})) = MI(\mathbf{X}; s^*)$$

The definition of MI:  $MI(\mathbf{X}; s) = H(\mathbf{X}) - H(\mathbf{X}|s)$ 



Datasets:

- Transductive: Cora, Citeseer, Pubmed;
- Inductive: Reddit (large graph), PPI (multiple graphs).

In Cora, Citeseer, Pubmed (transductive): Encoder  $\mathcal{E}$  is one-layer GCN:

$$\mathcal{E}(X,A) = \sigma(\widehat{D}^{-\frac{1}{2}}\widehat{A}\widehat{D}^{-\frac{1}{2}}X\Theta)$$

where  $\hat{A} = A + I_N$  and  $\sigma$  is parametric ReLU.

Corruption C set  $\tilde{A} = A$  and  $\tilde{X}$  as randomly row-wise shuffling of X.



Datasets:

- Transductive: Cora, Citeseer, Pubmed;
- Inductive: Reddit (large graph), PPI (multiple graphs).

In Reddit (large graph):

Encoder  $\mathcal{E}$  is three-layer mean-pooling model:

 $MP(X,A) = \widehat{D}^{-1}\widehat{A}X\Theta,$  $\widetilde{MP}(X,A) = \sigma(X\Theta'||MP(X,A)),$  $\mathcal{E}(X,A) = \widetilde{MP}_3(\widetilde{MP}_2(\widetilde{MP}_1(X,A),A),A)$ 

where  $\hat{A} = A + I_N$  and  $\sigma$  is parametric ReLU.

Corruption C set  $\tilde{A} = A$  and  $\tilde{X}$  as randomly row-wise shuffling of X.



Datasets:

- Transductive: Cora, Citeseer, Pubmed;
- Inductive: Reddit (large graph), PPI (multiple graphs).

In Reddit (multiple graphs):

Encoder  $\mathcal{E}$  is three-layer mean-pooling model with skip connections:

$$MP(X, A) = \widehat{D}^{-1}\widehat{A}X\Theta,$$
  

$$H_1 = \sigma(MP_1(X, A)),$$
  

$$H_2 = \sigma(MP_2(H_1 + XW_{skip}, A)),$$
  

$$\mathcal{E}(X, A) = \sigma(MP_3(H_2 + H_1 + XW_{skip}, A))$$

where  $W_{skip}$  is a learnable matrix, and  $\sigma$  is parametric ReLU.

Corruption  $\mathcal{C}$  is to select other graphs.



For all experiments:

Readout function  $\mathcal{R}$ :

$$\mathcal{R}(H) = \sigma\left(\frac{1}{N}\sum_{i=1}^{N}h_i\right)$$

 $\sigma$  is the sigmoid function.

Discriminator  $\mathcal{D}$ :

$$\mathcal{D}(h_i, s) = \sigma(h_i^T W s)$$

 $\sigma$  is the sigmoid function.

All models are initialized using Glorot initialization.

## **Experiments – Results**



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Transductive						
Available data	Method	Cora	Citeseer	Pubmed		
X	Raw features	$47.9\pm0.4\%$	$49.3\pm0.2\%$	$69.1\pm0.3\%$		
$\mathbf{A}, \mathbf{Y}$	LP (Zhu et al., 2003)	68.0%	45.3%	63.0%		
Α	DeepWalk (Perozzi et al., 2014)	67.2%	43.2%	65.3%		
$\mathbf{X}, \mathbf{A}$	DeepWalk + features	$70.7\pm0.6\%$	$51.4\pm0.5\%$	$74.3\pm0.9\%$		
$\mathbf{X}, \mathbf{A}$	Random-Init (ours)	$69.3 \pm 1.4\%$	$61.9 \pm 1.6\%$	$69.6 \pm 1.9\%$		
$\mathbf{X}, \mathbf{A}$	<b>DGI</b> (ours)	$\textbf{82.3}\pm0.6\%$	$\textbf{71.8} \pm 0.7\%$	$\textbf{76.8} \pm 0.6\%$		
$\mathbf{X}, \mathbf{A}, \mathbf{Y}$	GCN (Kipf & Welling, 2016a)	81.5%	70.3%	79.0%		
$\mathbf{X}, \mathbf{A}, \mathbf{Y}$	Planetoid (Yang et al., 2016)	75.7%	64.7%	77.2%		

		Inductive				
	Available data	Method	Reddit	PPI		
MICro-F1 In Inductive	X	Raw features	0.585	0.422		
$\sum_{i} TP_i$	$\mathbf{A}$	DeepWalk (Perozzi et al., 2014)	0.324	_		
$Precision_{mi} = \frac{\Sigma_i + \tau_i}{\Sigma_i [TP_i + FP_i]}$	$\mathbf{X}, \mathbf{A}$	DeepWalk + features	0.691	—		
	$\mathbf{X}, \mathbf{A}$	GraphSAGE-GCN (Hamilton et al., 2017a)	0.908	0.465		
$Pocall - \sum_{i} TP_i$	$\mathbf{X}, \mathbf{A}$	GraphSAGE-mean (Hamilton et al., 2017a)	0.897	0.486		
$\frac{1}{\sum_{i} [TP_i + FN_i]}$	$\mathbf{X}, \mathbf{A}$	GraphSAGE-LSTM (Hamilton et al., 2017a)	0.907	0.482		
	$\mathbf{X}, \mathbf{A}$	GraphSAGE-pool (Hamilton et al., 2017a)	0.892	0.502		
$F1_{mi} = 2 \frac{Precision_{mi} \times Recall_{mi}}{Precision_{mi} + Recall_{mi}}$	<b>X</b> , <b>A</b> <b>X</b> , <b>A</b>	Random-Init (ours) DGI (ours)	$\begin{array}{c} 0.933 \pm 0.001 \\ \textbf{0.940} \pm 0.001 \end{array}$	$\begin{array}{c} 0.626 \pm 0.002 \\ \textbf{0.638} \pm 0.002 \end{array}$		
	$\overline{ \begin{array}{c} \mathbf{X}, \mathbf{A}, \mathbf{Y} \\ \mathbf{X}, \mathbf{A}, \mathbf{Y} \end{array} }$	FastGCN (Chen et al., 2018) Avg. pooling (Zhang et al., 2018)	$\begin{array}{c} 0.937 \\ 0.958 \pm 0.001 \end{array}$			

Accuracy in transductive



t-SNE embeddings of the nodes in the Cora dataset from:

the raw features (left),

features from a randomly initialized DGI model (middle),

and a learned DGI model (right).





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## Thank you

