

# GraphNorm:

A Principled Approach to Accelerating  
Graph Neural Network Training

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<https://github.com/lcj2408/GraphNorm>



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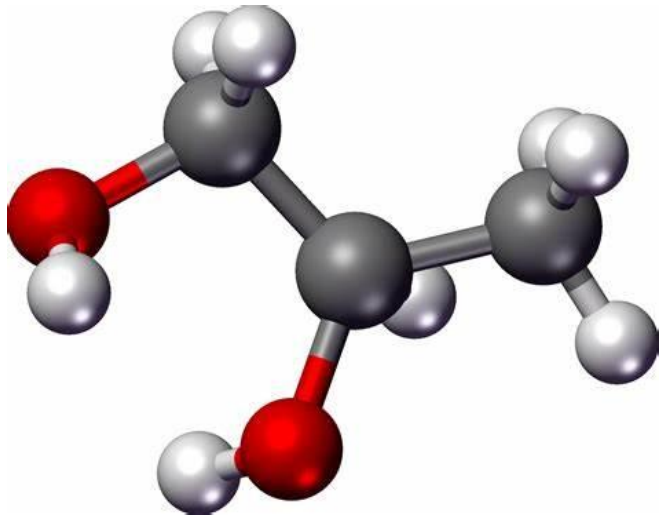


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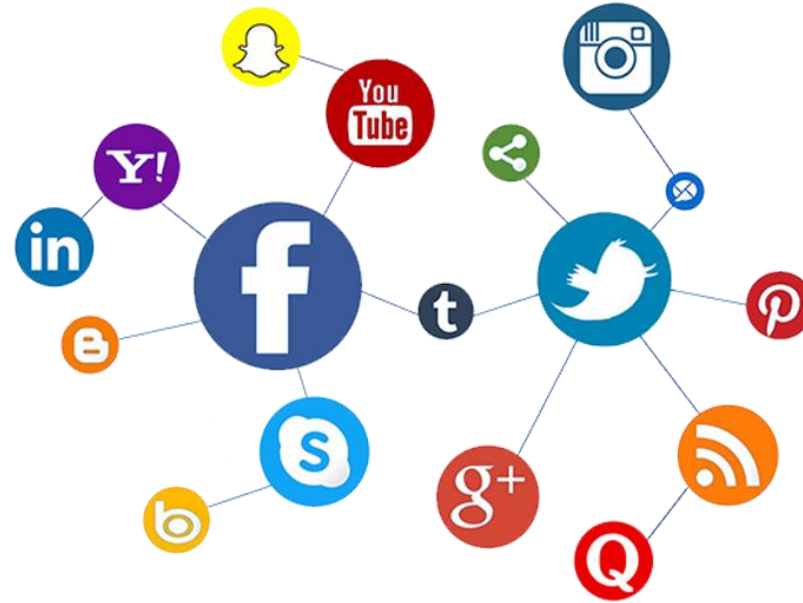
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# Learning with graph – a general form of data



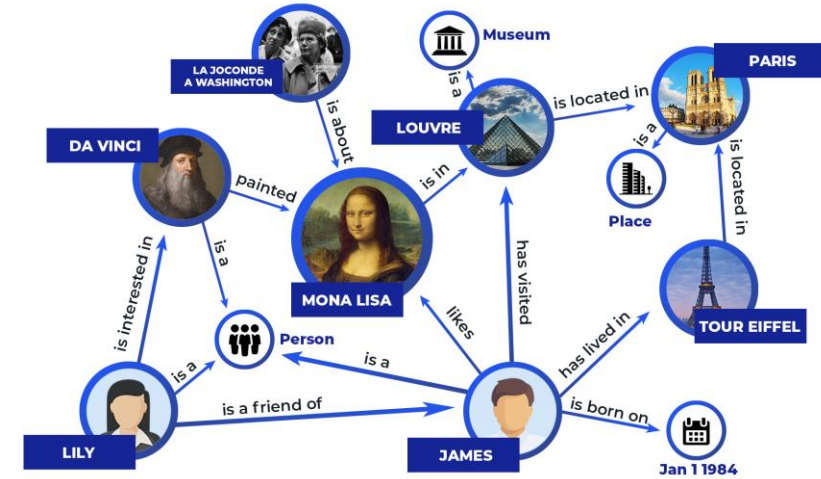
## Drug Discovery

(phys.org)



## Social Networks

(acwits)



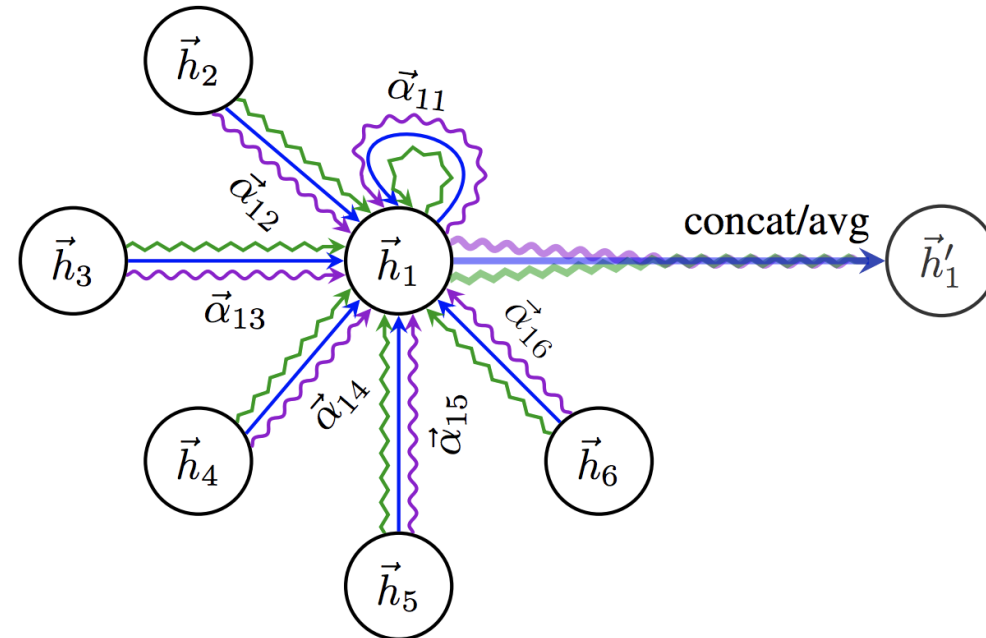
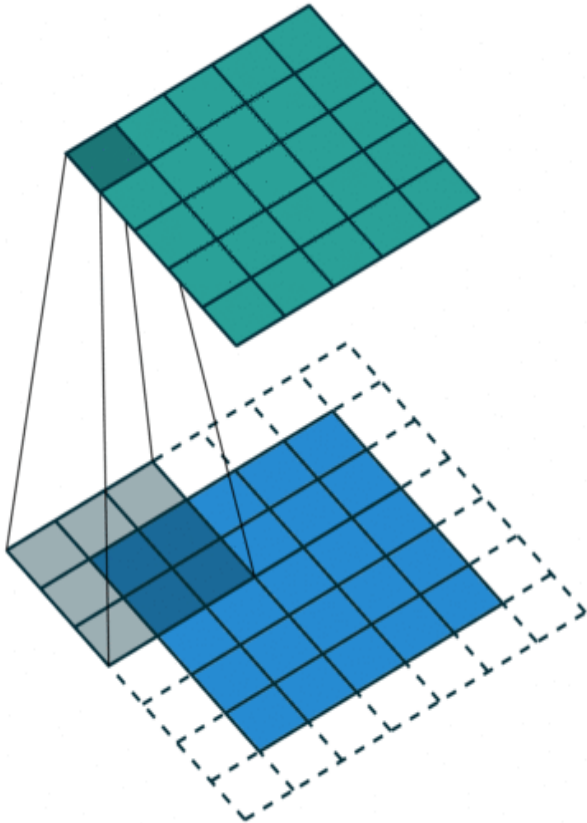
## Knowledge Graph

(yashueth.blog)

# Graph Neural Networks

**Neighborhood Aggregation** a.k.a. Message Passing or Graph Convolution

Aggregate neighbor features with permutation invariance functions



# Graph Neural Networks

## Neighborhood Aggregation

$$h_i^{(k)} = \text{AGGREGATE}^{(k)} \left( h_i^{(k-1)}, \{h_j^{(k-1)} : v_j \in \mathcal{N}(v_i)\} \right)$$

### Example -- GIN

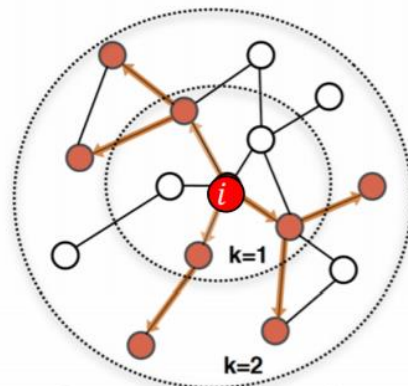
Feature of node  $v_j$  in layer  $k-1$ .

$$h_i^{(k)} = \text{MLP}^{(k)} \left( (1 + \epsilon^{(k)}) \cdot h_i^{(k-1)} + \sum_{j \in \mathcal{N}(v_i)} h_j^{(k-1)} \right)$$

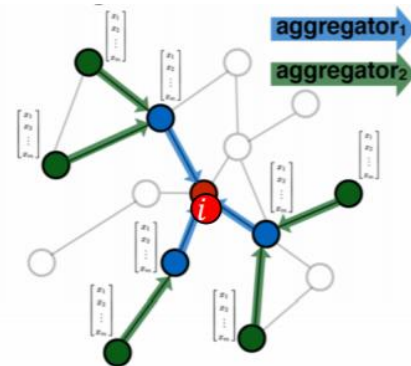
$\epsilon^{(k)}$  is a learnable parameter

## Readout Function

$$h_G = \text{READOUT}(\{h_i^{(K)} \mid v_i \in V\})$$



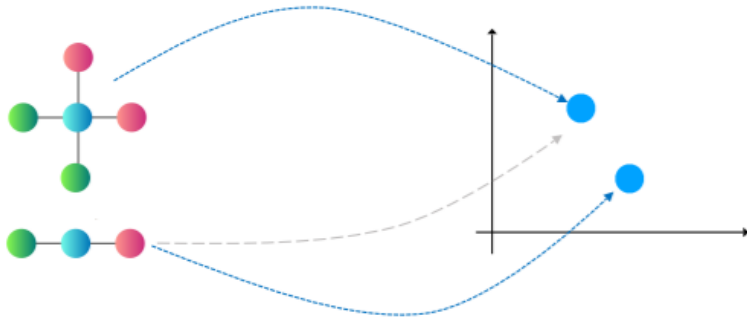
Determine node computation graph



Propagate and transform information

# Investigations on Graph Neural Network

## Expressive Power



Which graphs can a GNN distinguish?

## Reasoning

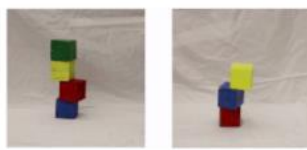
Reasoning tasks as dynamic programming (DP):



graph algorithms

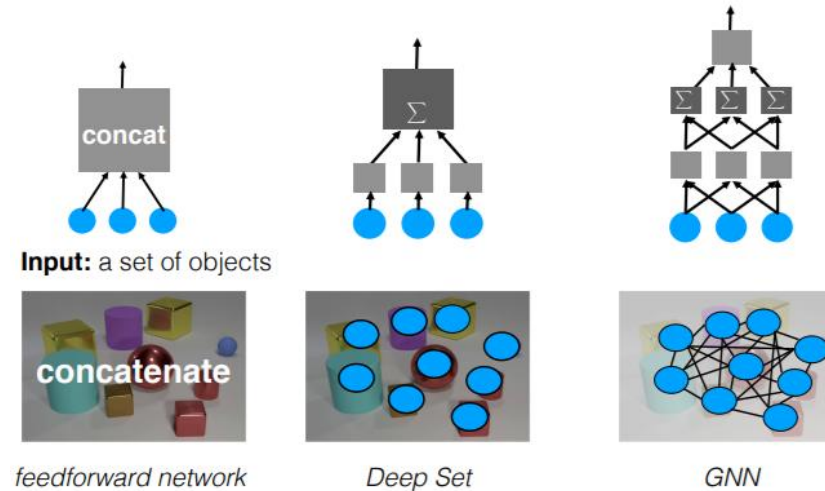


visual question answering

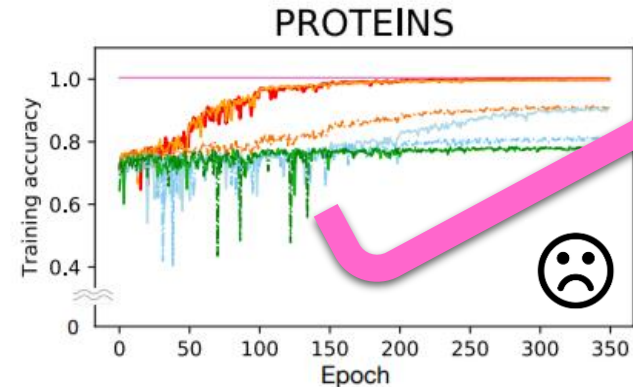


Intuitive physics

## Generalization



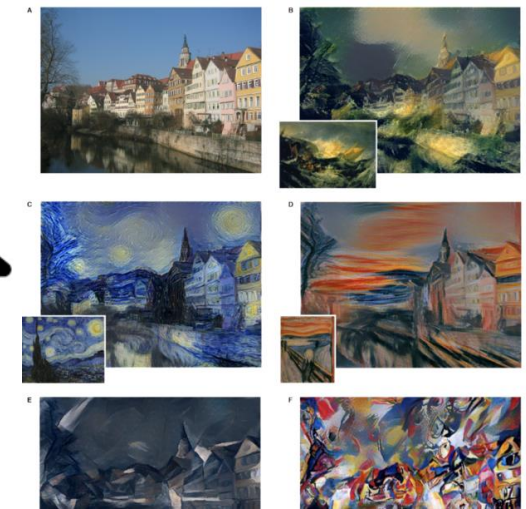
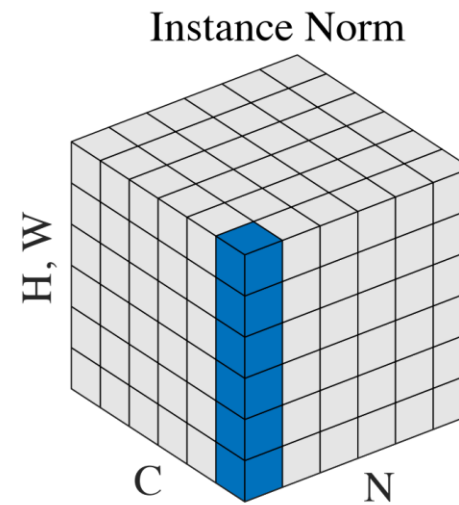
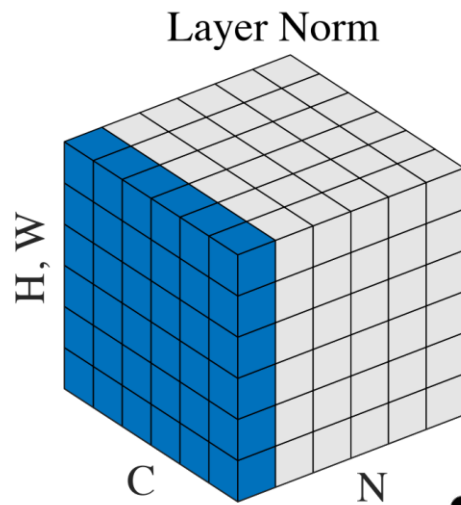
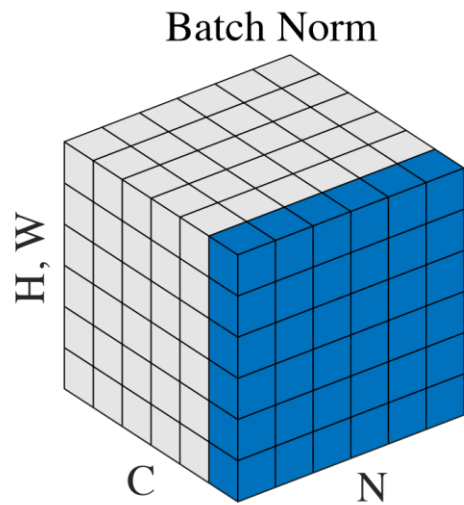
## Optimization?



- Training instability
- Slow convergence

**Even with Normalization methods.**

# Normalization for Neural Networks



Style Transfer

IMAGENET

Image Classification



NLP Tasks

*For GNNs, BN and LN are simply adopted without further investigations.*



What **normalization** methods are **effective** for **Graph Neural Networks**?



# This paper



Adapting and evaluating existing normalization methods to GNNs.



Explaining the effectiveness of InstanceNorm over BatchNorm.



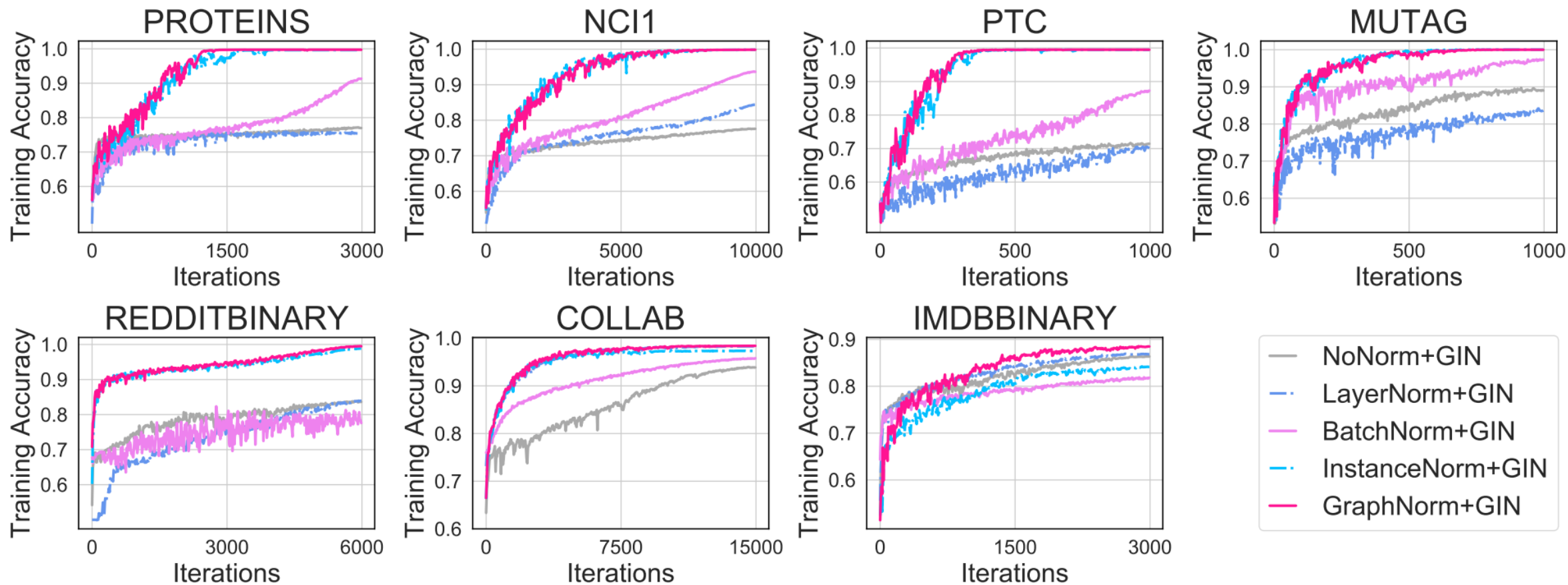
Identifying an expressiveness degradation of InstanceNorm.



Proposing GraphNorm which addresses the issue and converges faster.



# Evaluation of existing normalization methods



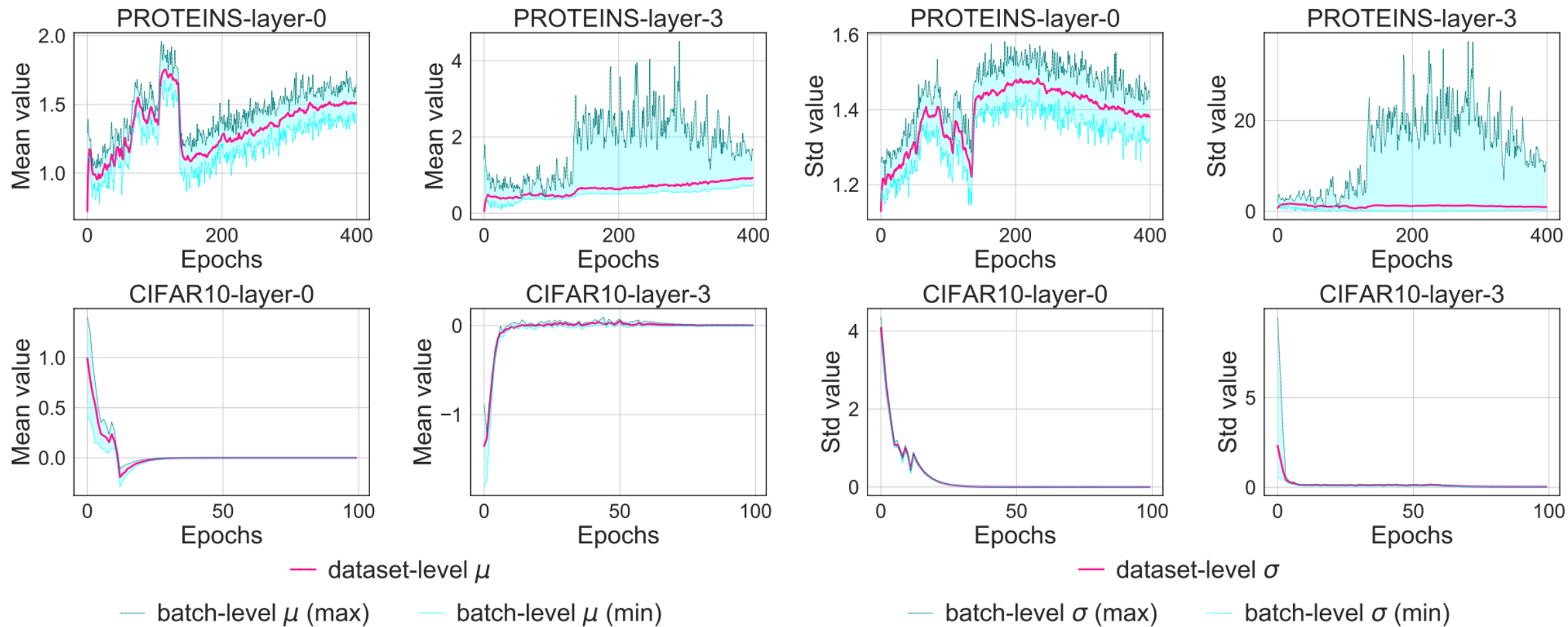
# Preconditioning effect of InstanceNorm

**Theorem 3.1** (Shift Serves as a Preconditioner of  $Q$ ). *Let  $Q, N$  be defined as in Eq. (6),  $0 \leq \lambda_1 \leq \dots \leq \lambda_n$  be the singular values of  $Q$ . We have  $\mu_n = 0$  is one of the singular values of  $QN$ , and let other singular values of  $QN$  be  $0 \leq \mu_1 \leq \mu_2 \leq \dots \leq \mu_{n-1}$ . Then we have*

$$\lambda_1 \leq \mu_1 \leq \lambda_2 \leq \dots \leq \lambda_{n-1} \leq \mu_{n-1} \leq \lambda_n, \quad (7)$$

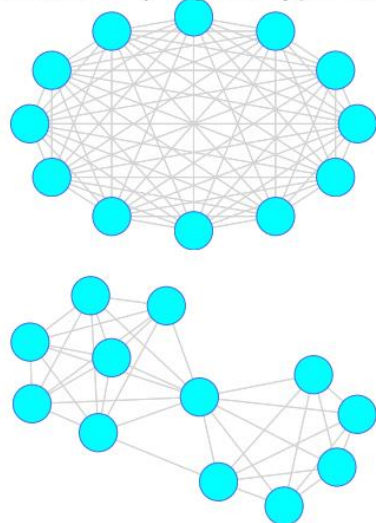
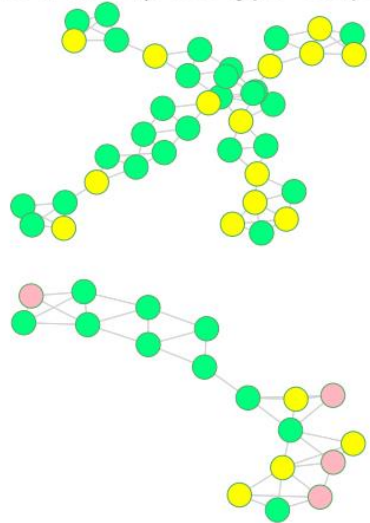
*where  $\lambda_i = \mu_i$  or  $\lambda_i = \mu_{i-1}$  only if there exists one of the right singular vectors  $\alpha_i$  of  $Q$  associated with  $\lambda_i$  satisfying  $\mathbf{1}^\top \alpha_i = 0$ .*

# Heavy batch noise in graphs



# Expressiveness degradation of InstanceNorm

PROTEINS (Tree-type Graphs) IMDBBINARY (Regular-type Graphs)




**Proposition 4.1.** *For a  $r$ -regular graph with one-hot encodings as its features described above, we have for GIN, Norm  $(W^{(1)}H^{(0)}Q_{\text{GIN}}) = S(W^{(1)}H^{(0)}Q_{\text{GIN}})N = 0$ , i.e., the output of normalization layer is a zero matrix without any information of the graph structure.*

**Proposition 4.2.** *For a complete graph ( $r = n - 1$ ), we have for GIN,  $Q_{\text{GIN}}N = \xi^{(k)}N$ , i.e., graph structural information in  $Q$  will be removed after multiplying  $N$ .*

# Proposed method: GraphNorm

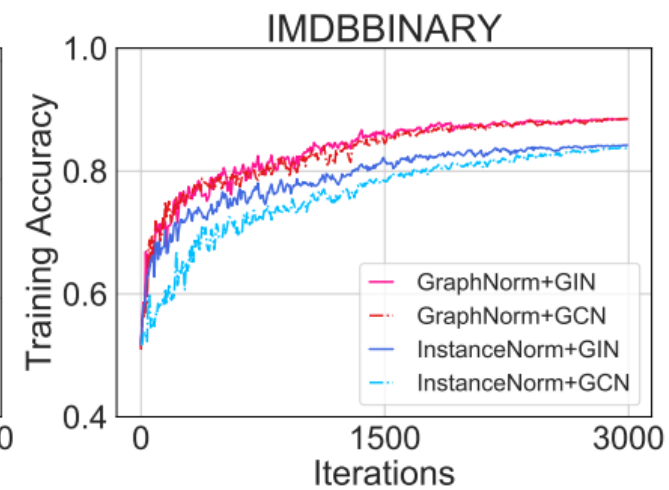
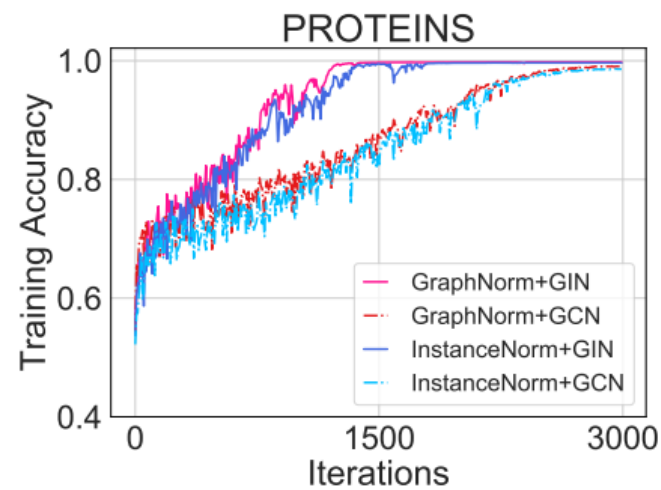
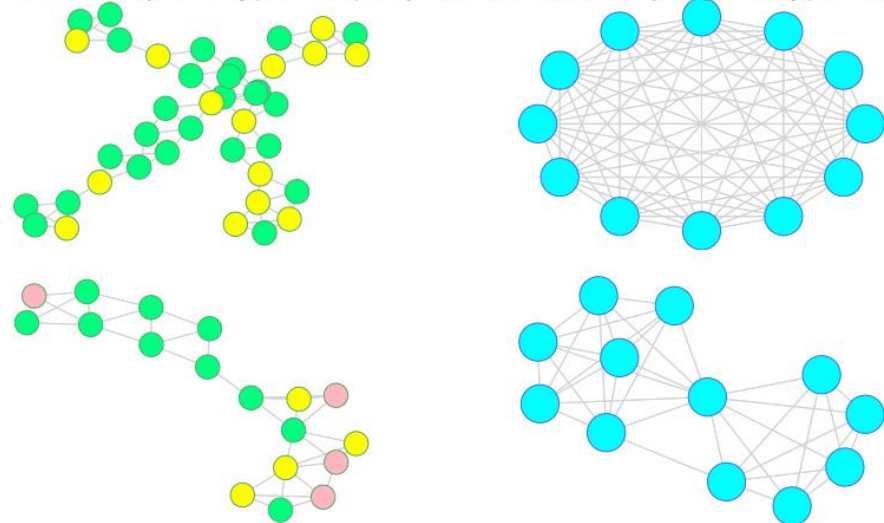
Key: **learnable** parameter to control how much the **information** we need to keep in the **mean**

$$\text{GraphNorm} \left( \hat{h}_{i,j} \right) = \gamma_j \cdot \frac{\hat{h}_{i,j} - \alpha_j \cdot \mu_j}{\hat{\sigma}_j} + \beta_j,$$


- Inheriting the merit of InstanceNorm
- Solving the expressiveness degradation problem

# GraphNorm addresses the issue of InstanceNorm

PROTEINS (Tree-type Graphs) IMDBBINARY (Regular-type Graphs)





# GraphNorm achieves good performance

Datasets	MUTAG	PTC	PROTEINS	NCI1	IMDB-B	RDT-B	COLLAB
# graphs	188	344	1113	4110	1000	2000	5000
# classes	2	2	2	2	2	2	2
Avg # nodes	17.9	25.5	39.1	29.8	19.8	429.6	74.5
WL SUBTREE (SHERVASHIDZE ET AL., 2011)	90.4 ± 5.7	59.9 ± 4.3	75.0 ± 3.1	<b>86.0 ± 1.8</b>	73.8 ± 3.9	81.0 ± 3.1	78.9 ± 1.9
DCNN (ATWOOD & TOWSLEY, 2016)	67.0	56.6	61.3	62.6	49.1	-	52.1
DGCNN (ZHANG ET AL., 2018)	85.8	58.6	75.5	74.4	70.0	-	73.7
AWL (IVANOV & BURNAEV, 2018)	87.9 ± 9.8	-	-	-	74.5 ± 5.9	87.9 ± 2.5	73.9 ± 1.9
GIN+LAYERNORM	82.4 ± 6.4	62.8 ± 9.3	76.2 ± 3.0	78.3 ± 1.7	74.5 ± 4.4	82.8 ± 7.7	80.1 ± 0.8
GIN+BATCHNORM ((XU ET AL., 2019))	89.4 ± 5.6	64.6 ± 7.0	76.2 ± 2.8	82.7 ± 1.7	75.1 ± 5.1	92.4 ± 2.5	<b>80.2 ± 1.9</b>
GIN+INSTANCENORM	90.5 ± 7.8	64.7 ± 5.9	76.5 ± 3.9	81.2 ± 1.8	74.8 ± 5.0	93.2 ± 1.7	80.0 ± 2.1
<b>GIN+GraphNorm</b>	<b>91.6 ± 6.5</b>	<b>64.9 ± 7.5</b>	<b>77.4 ± 4.9</b>	81.4 ± 2.4	<b>76.0 ± 3.7</b>	<b>93.5 ± 2.1</b>	<b>80.2 ± 1.0</b>

Table 2. Test performance on OGB.

Datasets	OGBG-MOLHIV
# graphs	41,127
# classes	2
Avg # nodes	25.5
GCN (Hu et al., 2020)	76.06 ± 0.97
GIN (Hu et al., 2020)	75.58 ± 1.40
GCN+LayerNorm	75.04 ± 0.48
GCN+BatchNorm	76.22 ± 0.95
GCN+InstanceNorm	78.18 ± 0.42
<b>GCN+GraphNorm</b>	<b>78.30 ± 0.69</b>
GIN+LayerNorm	74.79 ± 0.92
GIN+BatchNorm	76.61 ± 0.97
GIN+InstanceNorm	77.54 ± 1.27
<b>GIN+GraphNorm</b>	<b>77.73 ± 1.29</b>



# Thank you :)

