



Learning on Graphs

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BRIEF BIO



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■ Research Interest

- **Data-Centric AI, Model-Centric AI, Multimodal Data Mining, AI for Science**
 - *Mining meaningful knowledge from multimodal data to develop artificial intelligence solutions for various real-world applications across different disciplines*
 - Multi-modal Learning, Graph neural network
 - **Application domains:** Recommendation system, Social network analysis, Fraud detection, Sentiment analysis, Purchase/Click prediction, Anomaly detection, Knowledge-graph construction, Time-series analysis, Bioinformatics, Chemistry etc.

■ Professional Experience

- Assistant Professor, **KAIST** (2020.11 – Present)
- Postdoctoral Research Fellow, **University of Illinois at Urbana-Champaign**, Dept. of Computer Science (2019. 1 – 2020. 10)
- Research Intern, **Microsoft Research Asia** (2017. 9 – 2017. 12)
- Research Intern, **NAVER** (2017. 3 – 2017. 6)

Outline

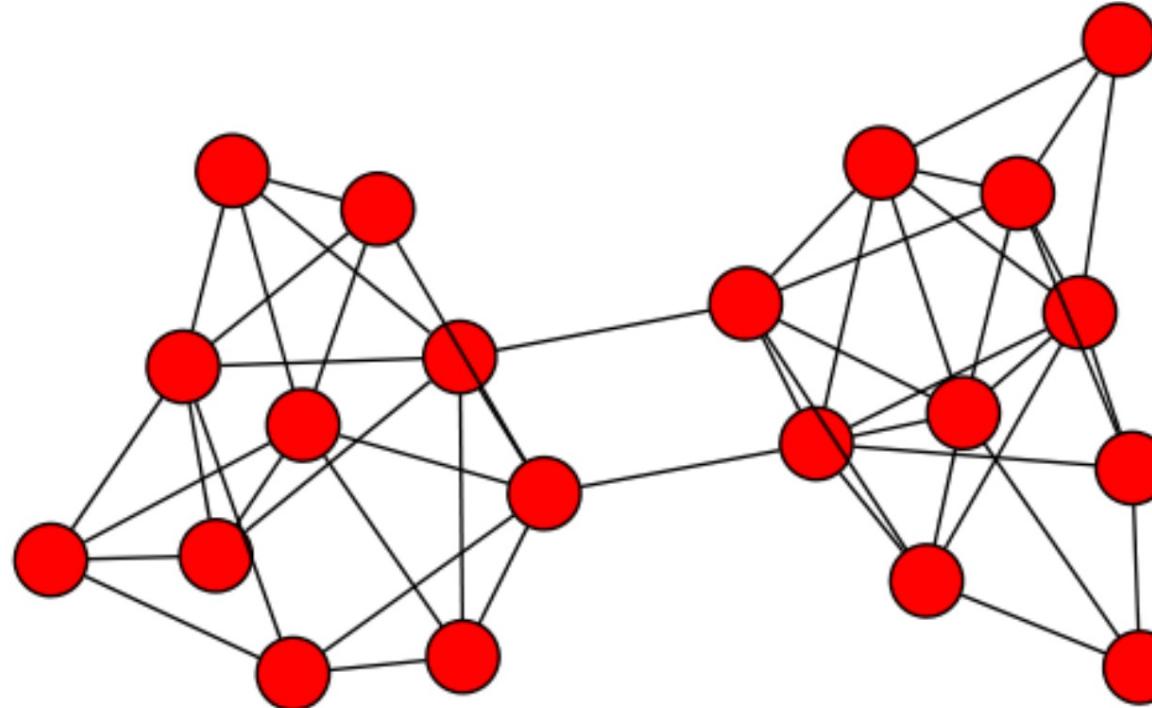
- Overview
- Random walk-based Methods
- Graph Neural Networks (GNNs)
- How to effectively train GNNs?
 - Self-supervised Learning (AAAI'22)
 - Explainable Model (NeurIPS'23)
 - Robustness (KDD'23)
- Applications of GNNs
 - GNNs for Science (화학/소재/생물) (ICML'23)
 - ~~GNNs for Medical Data (멀티모달 의료 데이터 분석) (AAAI'23)~~
 - ~~GNNs for Computer vision (Scene Understanding) (AAAI'23)~~

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Graph (network)

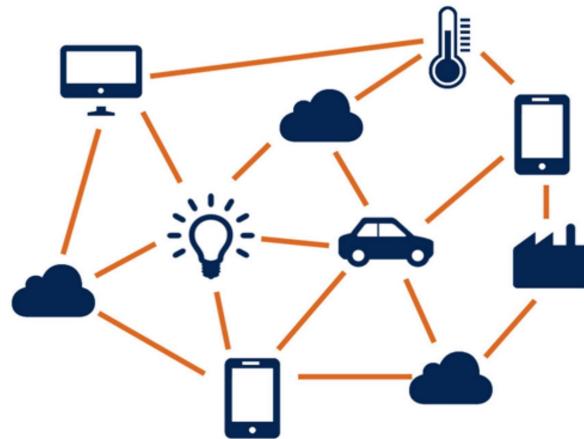
- A general description of data and their relations
- A collection of objects (i.e., nodes), along with a set of interactions (i.e., edges) between pairs of these objects.



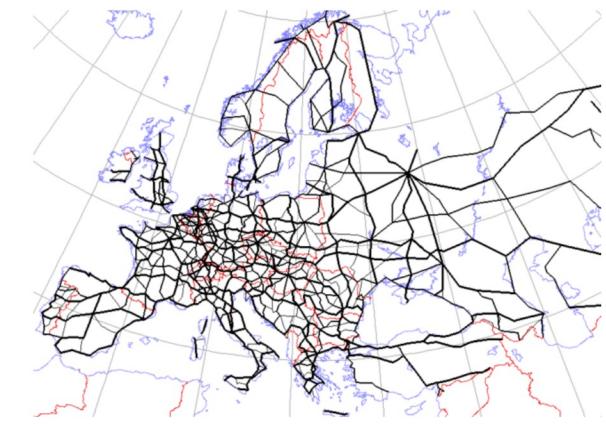
Various real-world graphs (networks)



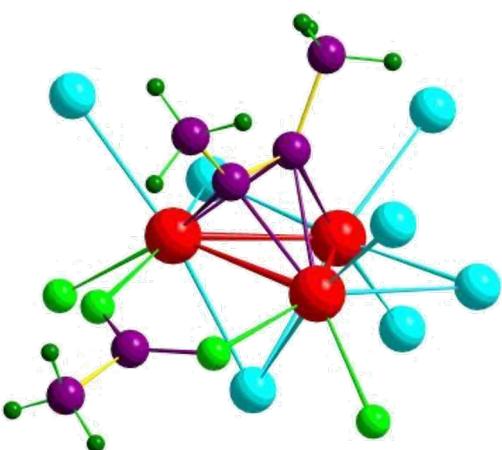
Social graph



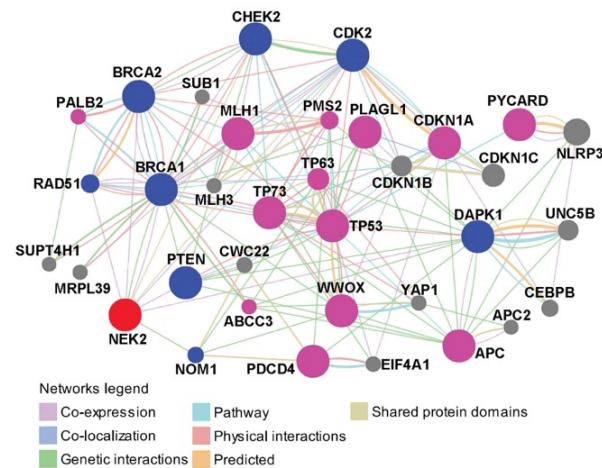
Internet-of-Things



Transportation



Molecular graph



Gene network



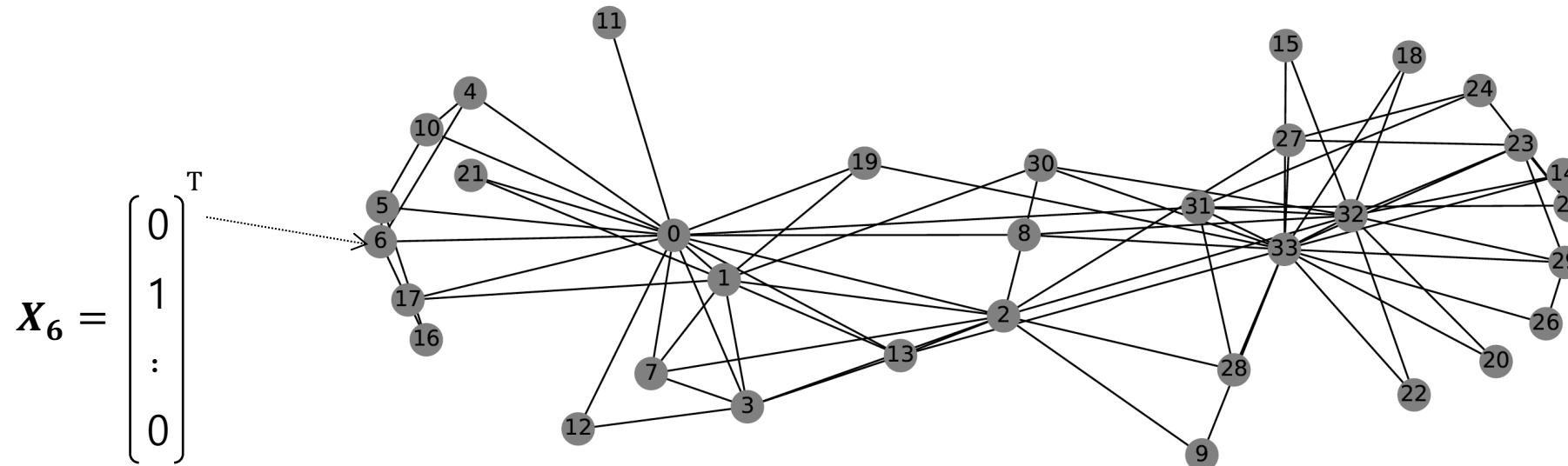
Web graph

Types of graph

- Undirected / Directed
- Homogeneous network
- Heterogeneous network
 - Multiplex network
 - Bipartite graph

Formal definition of graphs

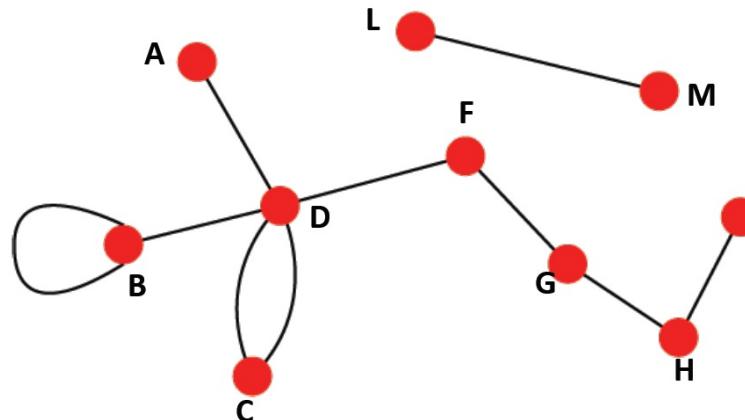
- A graph $G = (V, E)$ is defined by a set of nodes V and a set of edges E between these nodes
- An edge going from node $i \in V$ to node $j \in V$ is denoted as $(i, j) \in E$
- A convenient way to represent graphs is through an adjacency matrix $A \in R^{|V| \times |V|}$
 - $A_{ij} = 1$ if $(i, j) \in E$, and $A_{ij} = 0$ otherwise
 - Some graphs have **weighted** edges, i.e., entries of A are arbitrary real-values rather than $\{0,1\}$
- Feature information $X \in R^{|V| \times F}$



Undirected graph vs. directed graph

▪ Undirected Graph

- Adjacency matrix A is **symmetric**
- $(u, v) \in E \Leftrightarrow (v, u) \in E$



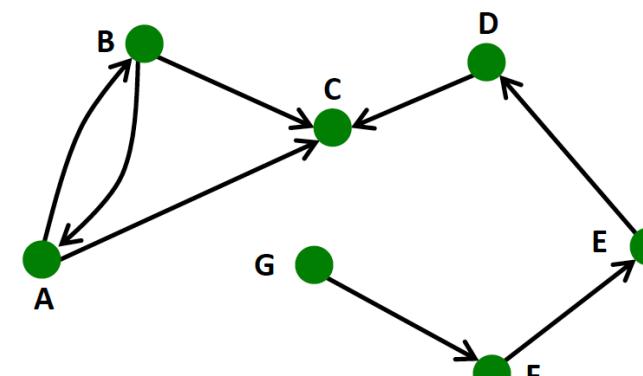
• Examples

- Collaborations
- Friendship on Facebook



▪ Directed Graph

- Adjacency matrix A is **not symmetric**
- $(u, v) \in E \nRightarrow (v, u) \in E$



• Examples

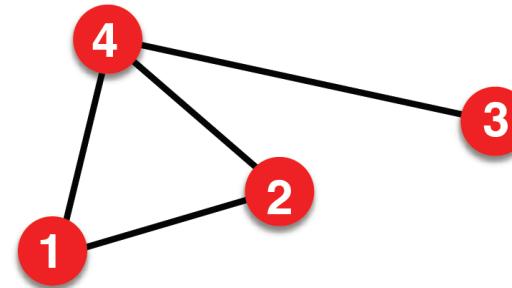
- Paper citation
- Follow on Twitter



Adjacency matrix

$A_{ij} = 1$ if there is a link from node i to node j
 $A_{ij} = 0$ otherwise

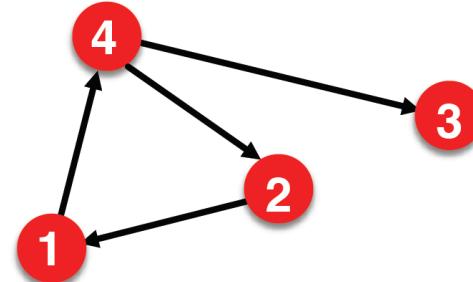
Undirected Graph



$$A_{ij} = A_{ji}$$
$$A_{ii} = 0$$

$$A = \begin{pmatrix} 0 & 1 & 0 & 1 \\ 1 & 0 & 0 & 1 \\ 0 & 0 & 0 & 1 \\ 1 & 1 & 1 & 0 \end{pmatrix}$$

Directed Graph



$$A_{ij} \neq A_{ji}$$
$$A_{ii} = 0$$

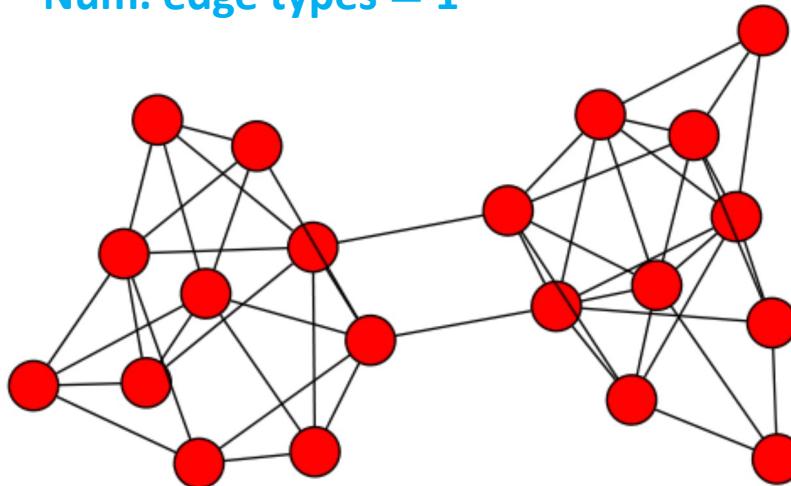
$$A = \begin{pmatrix} 0 & 0 & 0 & 1 \\ 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & 1 & 1 & 0 \end{pmatrix}$$

Homogeneous network

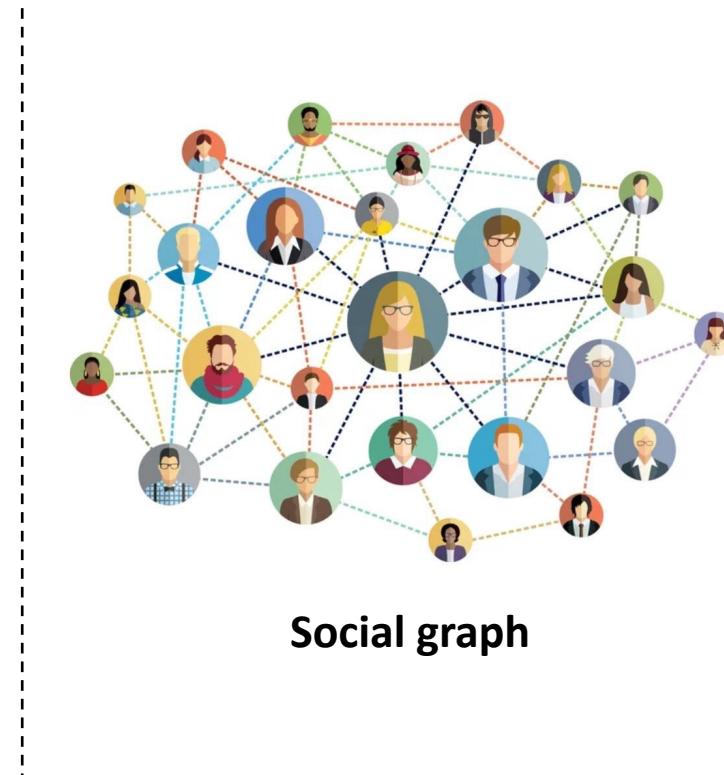
- A graph with a single type of node and a single type of edge

Num. node types = 1

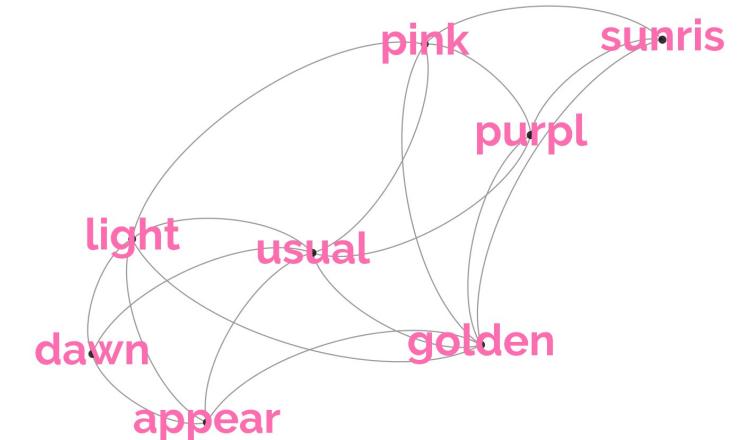
Num. edge types = 1



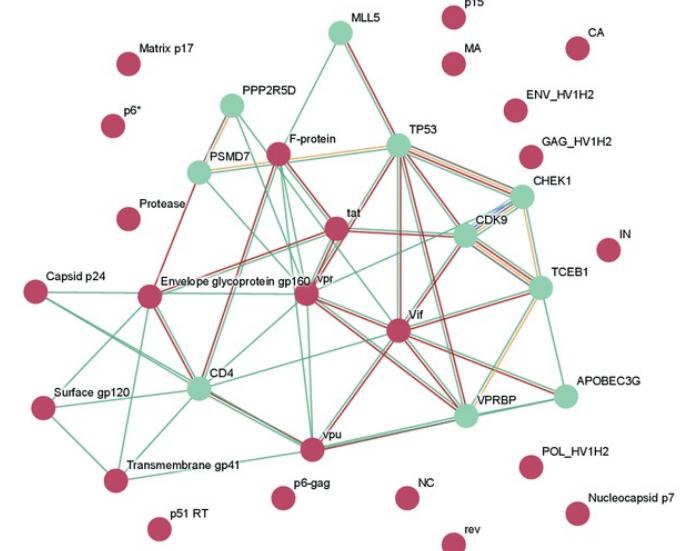
Homogeneous network



Social graph



Word cooccurrence graph

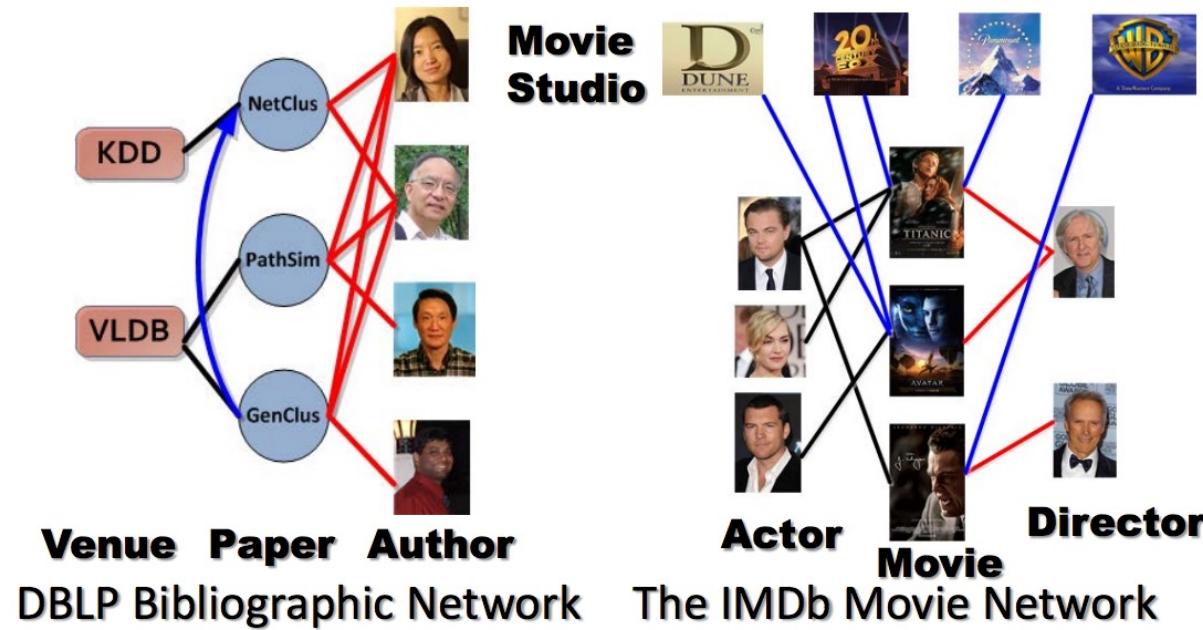


Protein-Protein Interaction Graph

Heterogeneous network (HetNet)

- In reality a lot of graphs have **multiple types of nodes** and **multiple types of edges**
- Such networks are called “**heterogeneous network**”

Num. node types > 1
Num. edge types > 1



Multiplex (Multi-layer) network

- A type of heterogeneous network
 - A single node type, multiple edge types

- **Example 1: Social network**

- Relationship between users

- **Example 2: E-commerce**

- Relationship between items

- **Example 3: Publication network**

- Relationship between papers (Citation, share authors)
 - Relationship between authors (Co-author, co-citation)

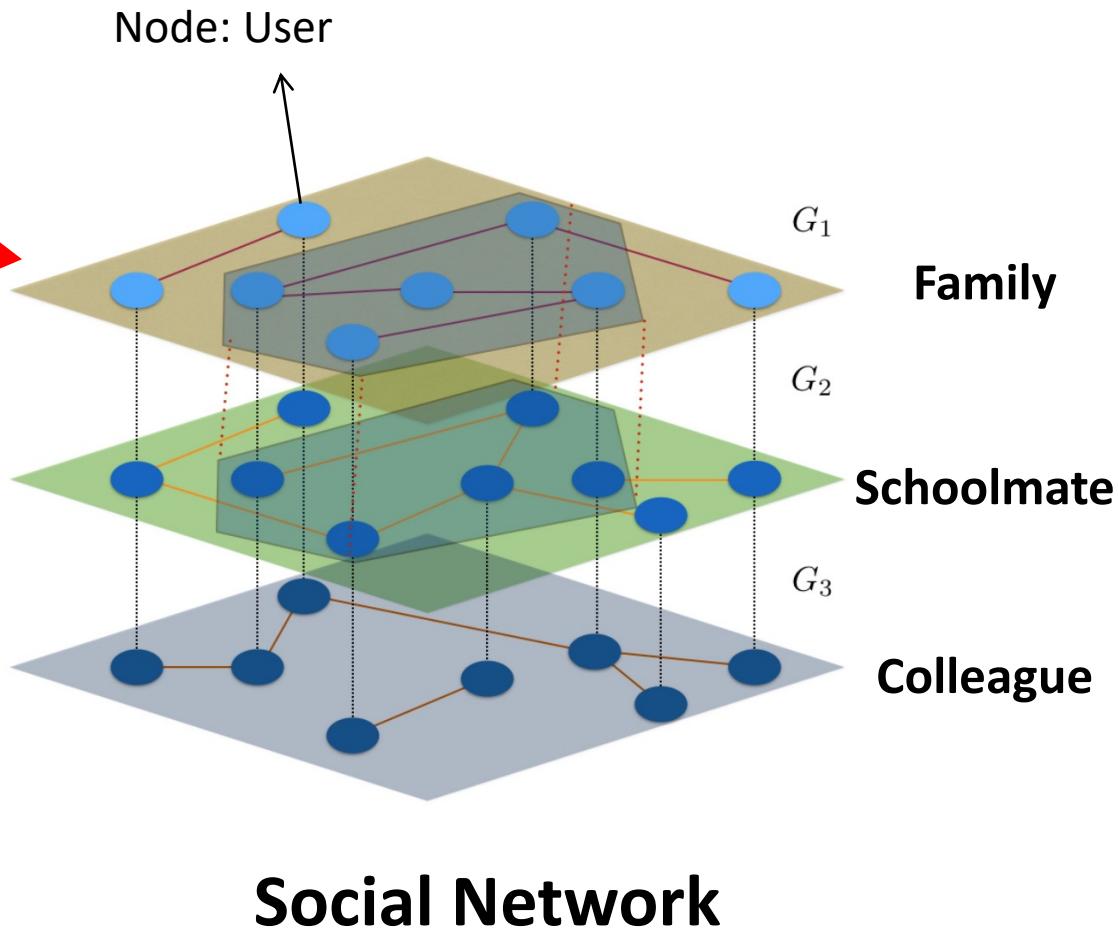
- **Example 4: Movie database**

- Relationship between movies
 - Common director, common actor

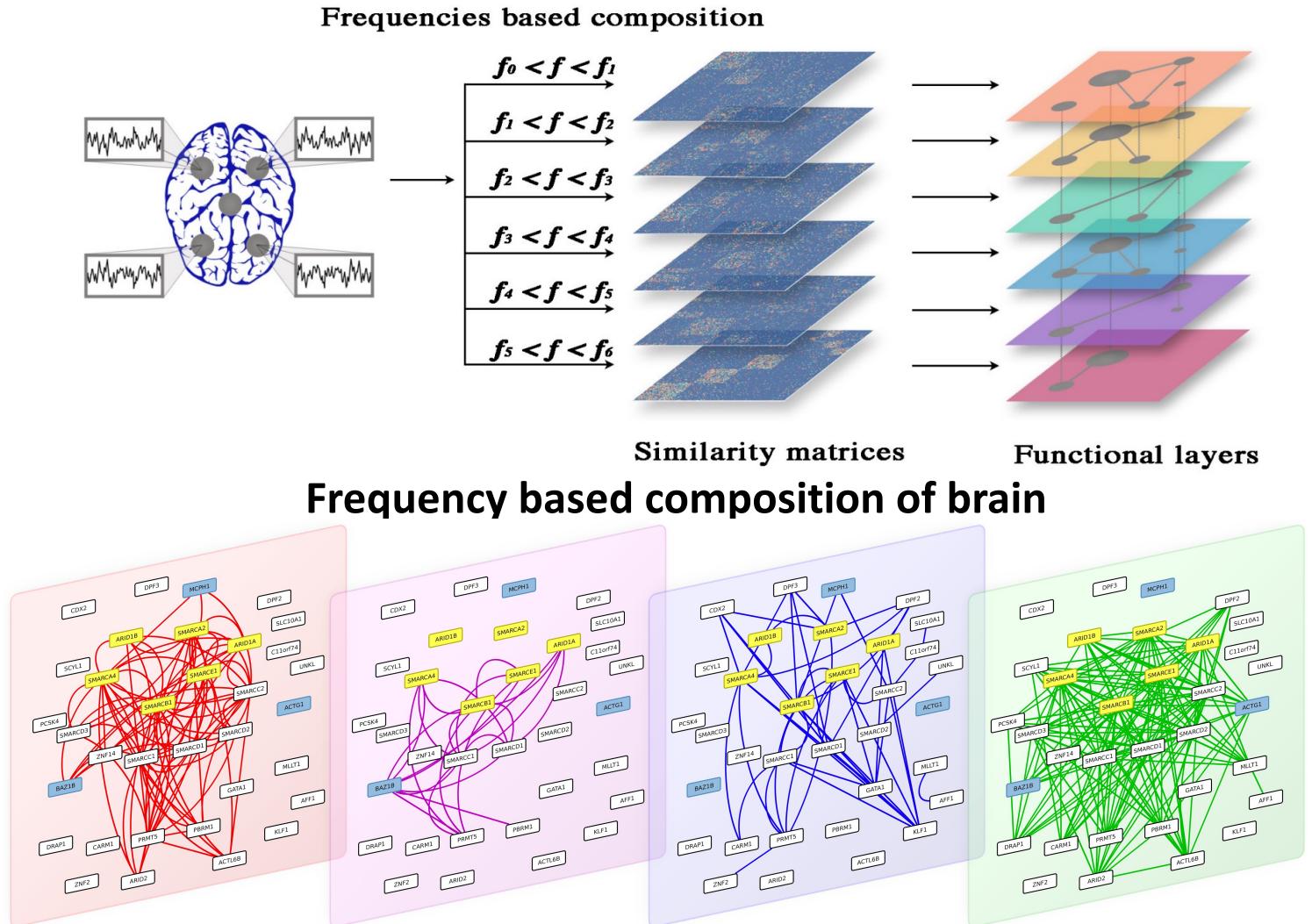
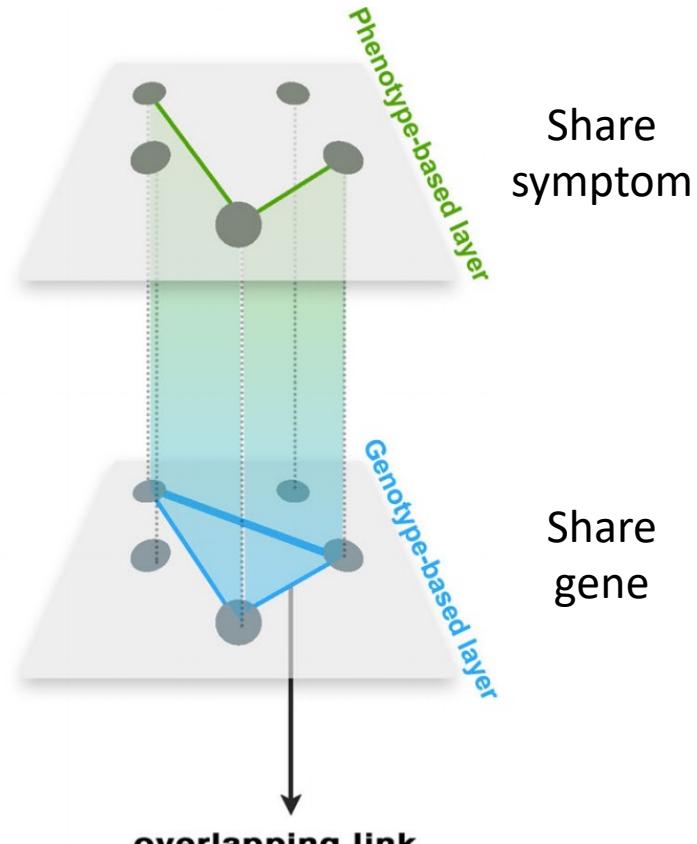
- **Example 5: Transportation network in a city**

- Relation between locations in a city
 - Bus, train, car, taxi

Num. node types = 1
Num. edge types > 1



Other examples of multiplex network



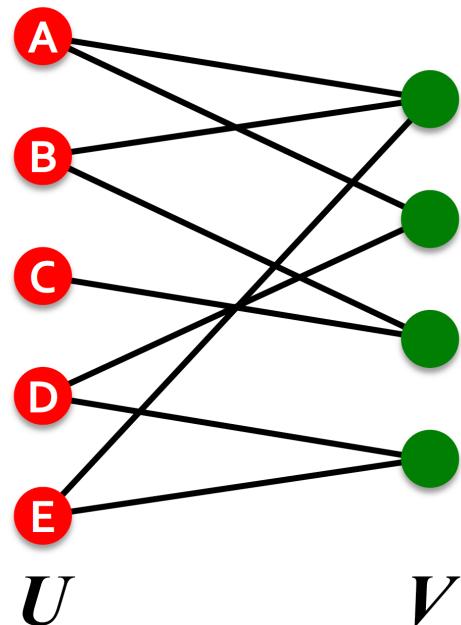
De Domenico, Manlio. "Multilayer modeling and analysis of human brain networks." *Giga Science* 6.5 (2017): gix004.

Halu, Arda, et al. "The multiplex network of human diseases." *NPJ systems biology and applications* 5.1 (2019): 1-12.

Didier, Gilles, Christine Brun, and Anaïs Baudot. "Identifying communities from multiplex biological networks." *PeerJ* 3 (2015): e1525.

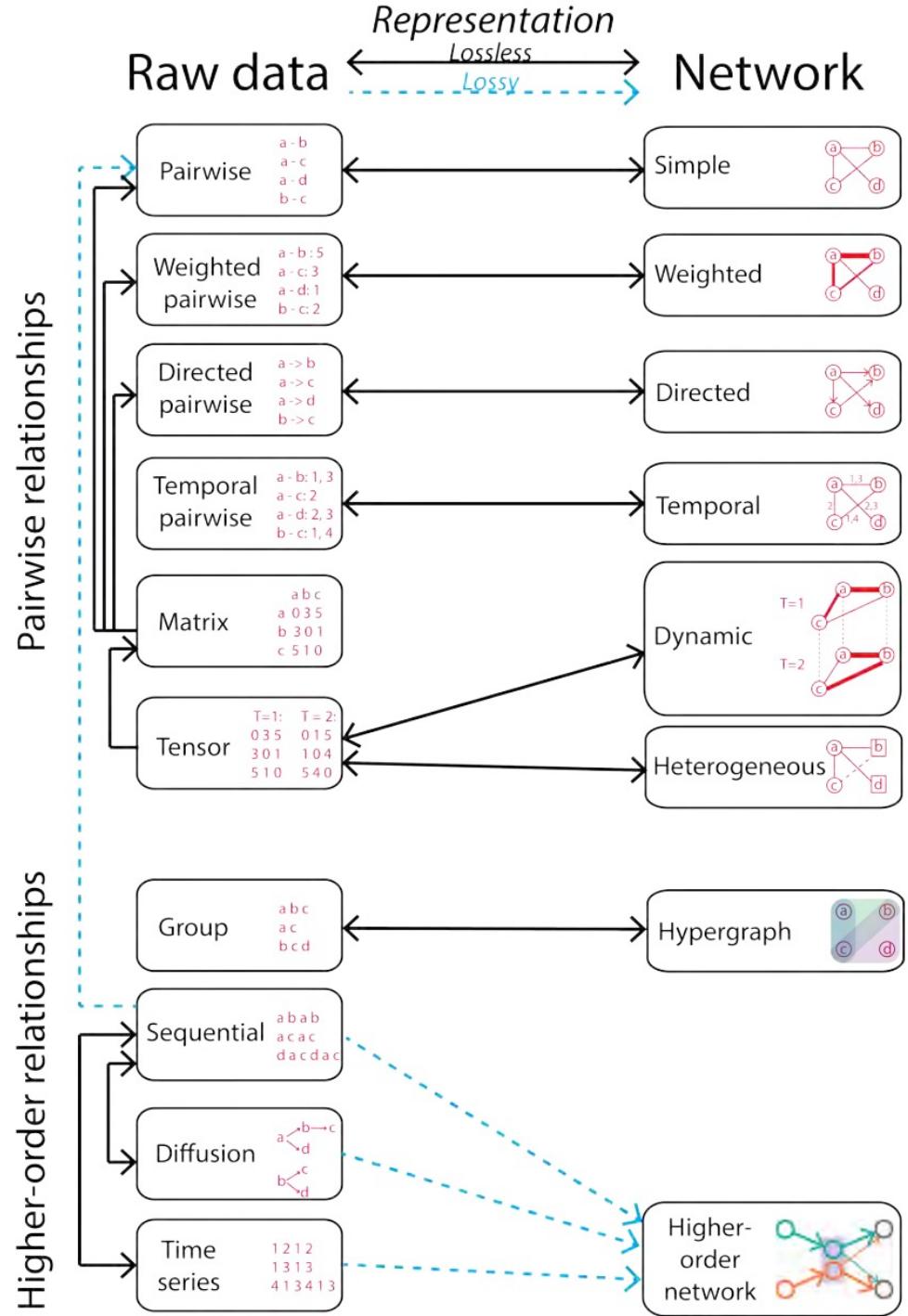
Bipartite graph

- Nodes can be divided into two disjoint sets U and V such that every link connects a node in U to one in V
 - U and V are independent sets



- Examples
 - Authors-to-Papers (they authored)
 - Actors-to-Movies (they appeared in)
 - Users-to-Movies (they rated)
 - Recipes-to-Ingredients (they contain)

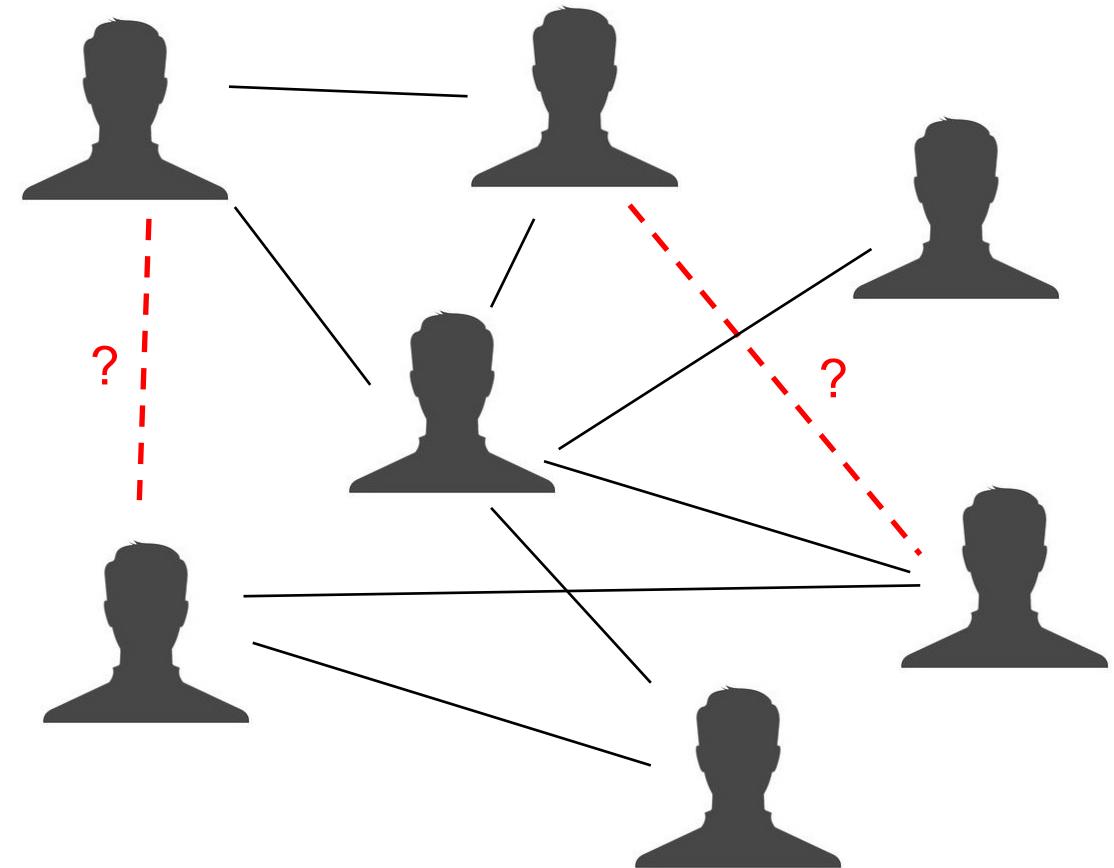
Overview: Data as Graphs



Machine learning on graphs

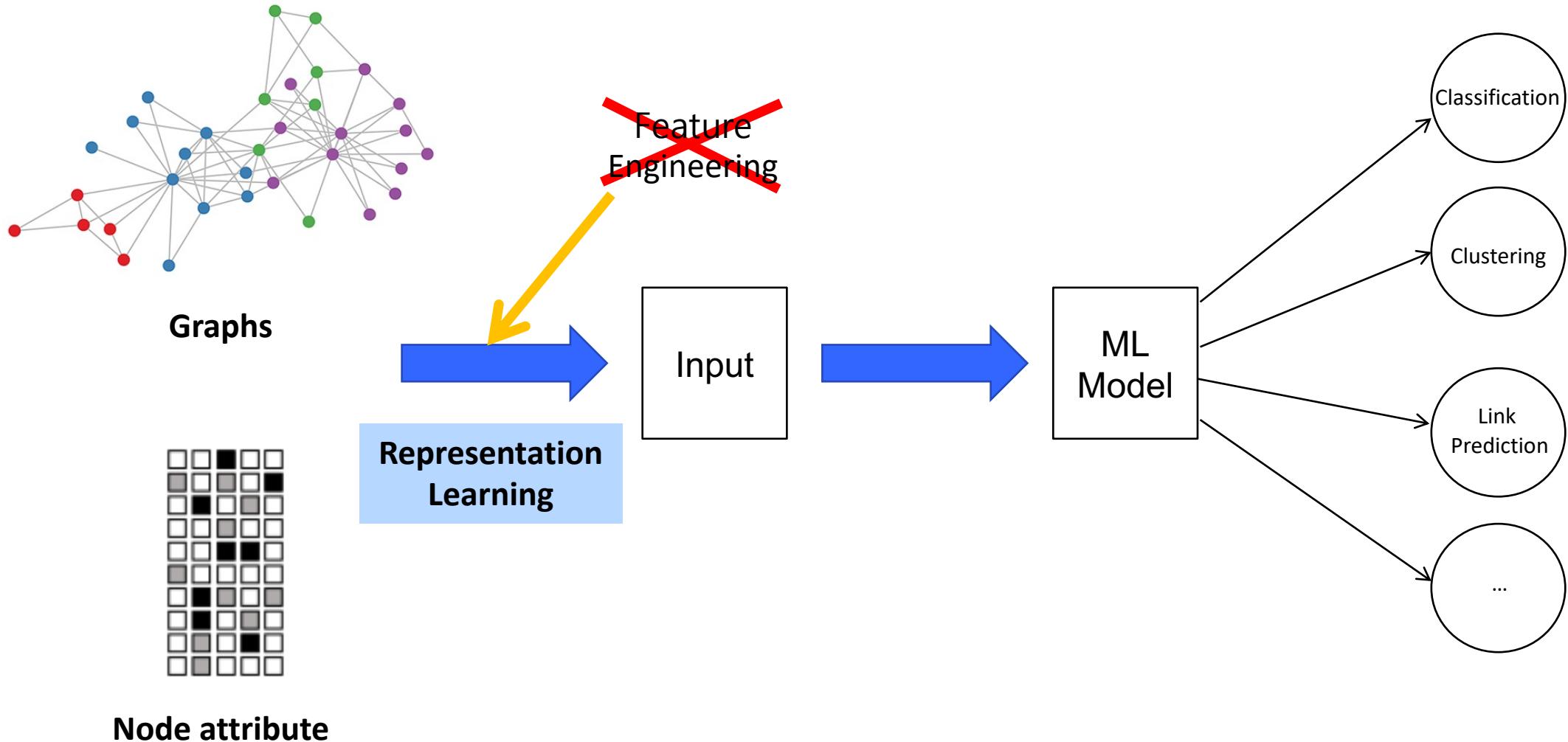
Classical ML tasks in graphs:

- Node classification
 - Predict a type of a given node
- Link prediction
 - Predict whether two nodes are linked
- Community detection
 - Identify densely linked clusters of nodes
- Network similarity
 - How similar are two (sub)networks



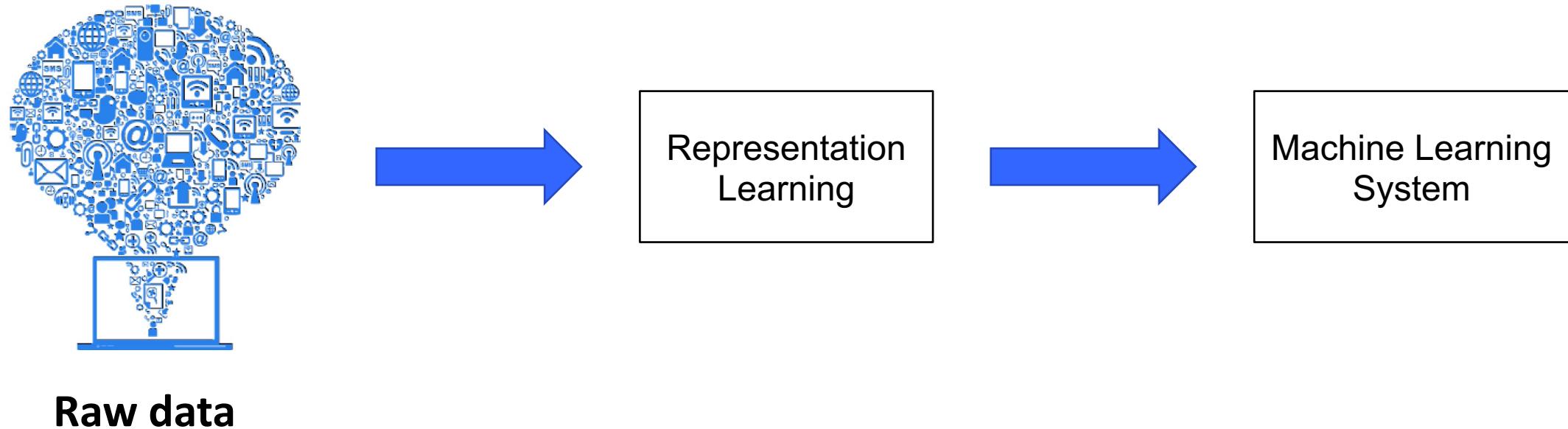
**Link Prediction
(Friend Recommendation)**

Machine learning on graphs



Machine learning in general

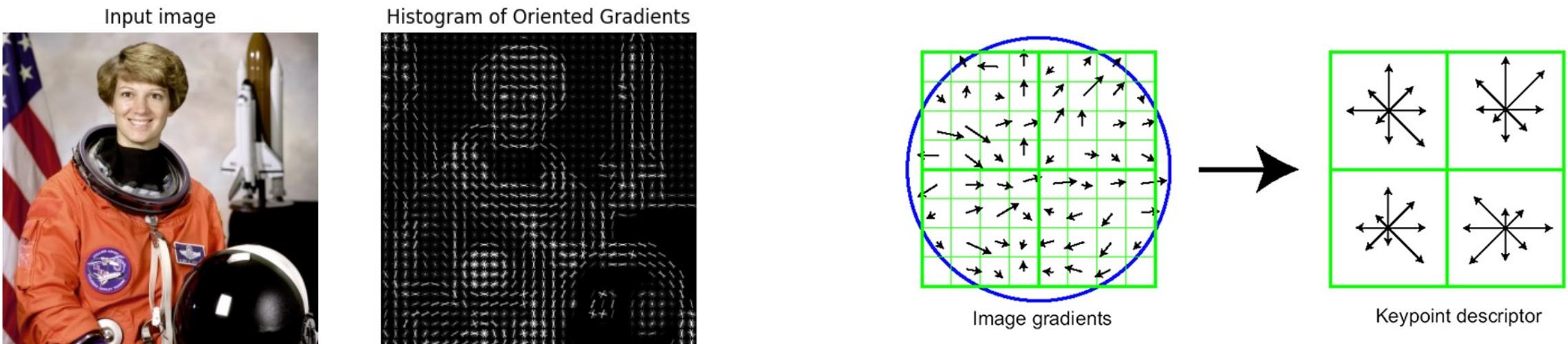
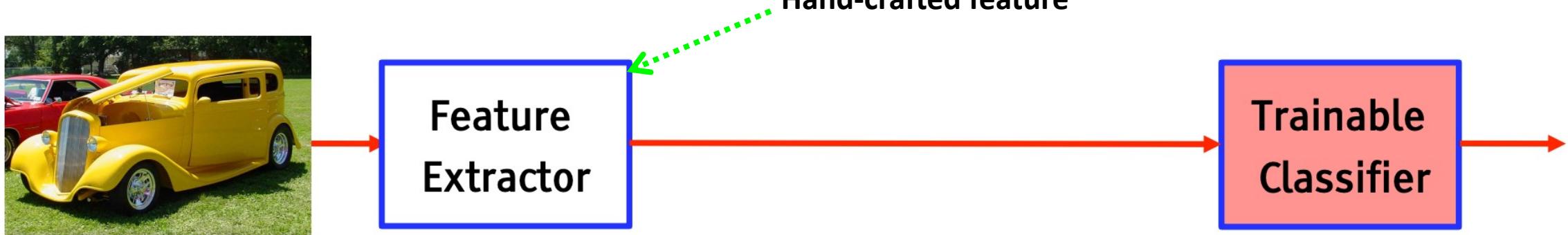
- Machine Learning = **Representation** + Objective + Optimization



Good **Representation** is Essential for
Good **Machine Learning**

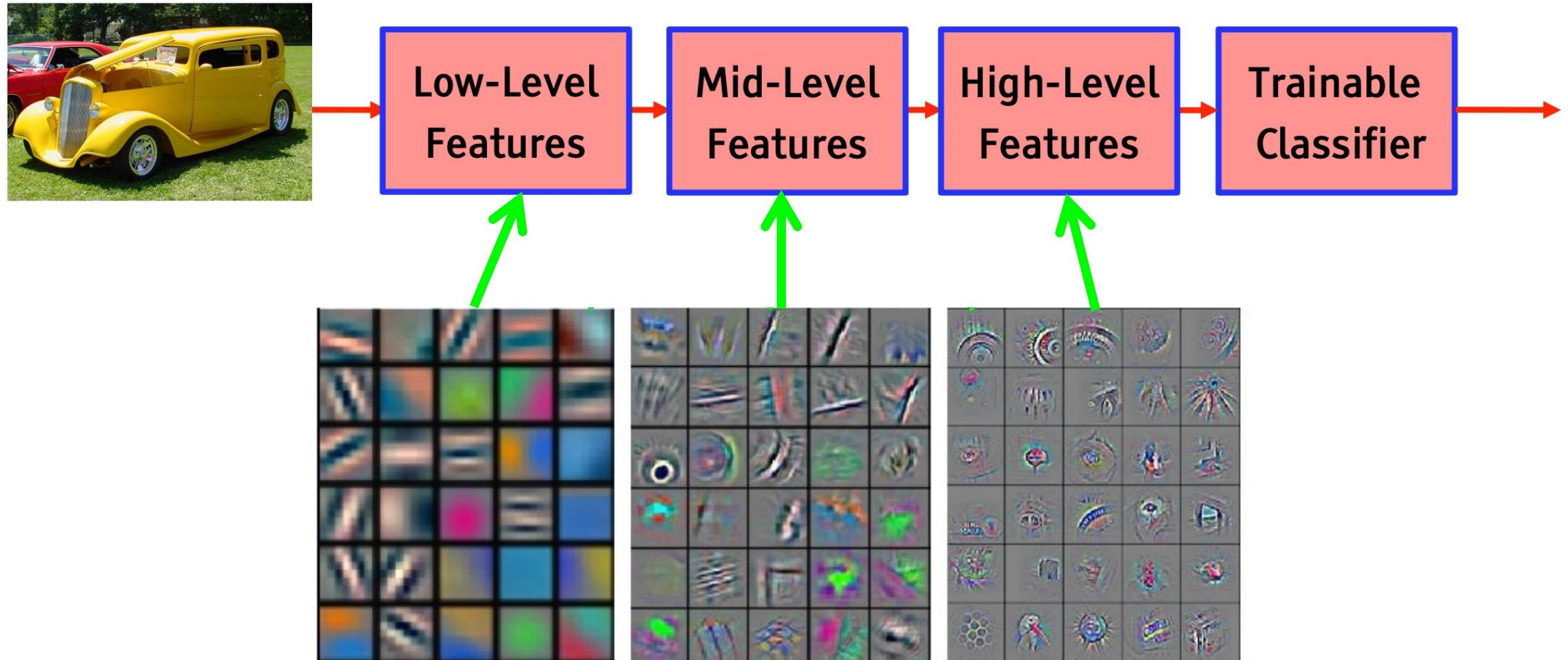
Traditional feature extraction for images

- Fixed/Hand-crafted Feature Extractor

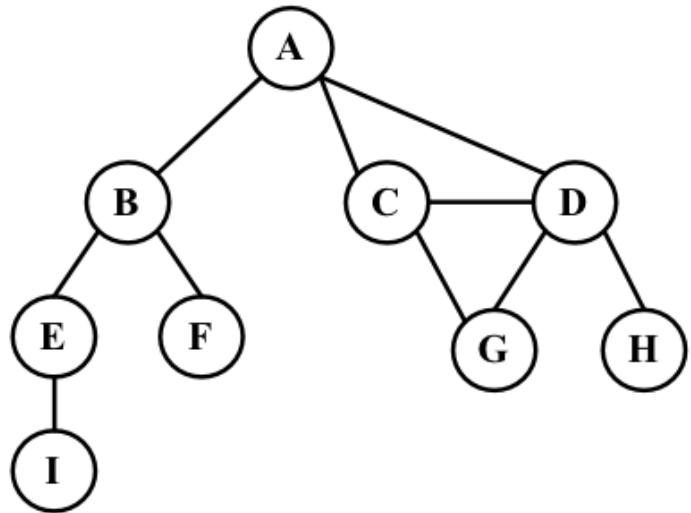


Machine (Deep) learning based Representation learning

- Multiple layers trained end-to-end



Traditional Graph Representation



	A	B	C	D	E	F	G	H	I
A	0	1	1	1	0	0	0	0	0
B	1	0	0	0	1	1	0	0	0
C	1	0	0	1	0	0	1	0	0
D	1	0	1	0	0	0	1	1	0
E	0	1	0	0	0	0	0	0	1
F	0	1	0	0	0	0	0	0	0
G	0	0	1	1	0	0	0	0	0
H	0	0	0	1	0	0	0	0	0
I	0	0	0	0	1	0	0	0	0

Adjacency matrix

How to effectively and efficiently represent graphs is the key!

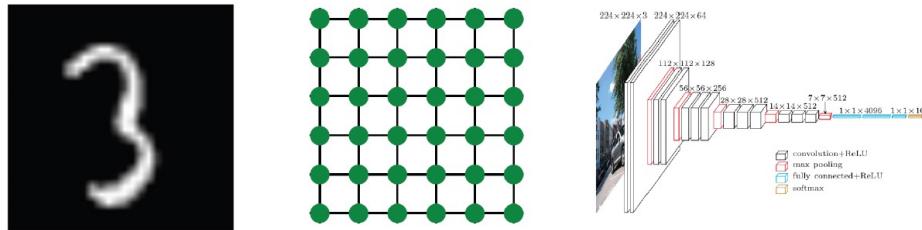
→ Deep learning-based approach?

Problems

- Suffer from data sparsity
- Suffer from high dimensionality
- High complexity for computation
- Does not represent “semantics”
- ...

Challenges of Graph Representation Learning

- Existing deep neural networks are designed for data with regular-structure (grid or sequence)
 - CNNs for fixed-size images/grids ...



- RNNs for text/sequences ...

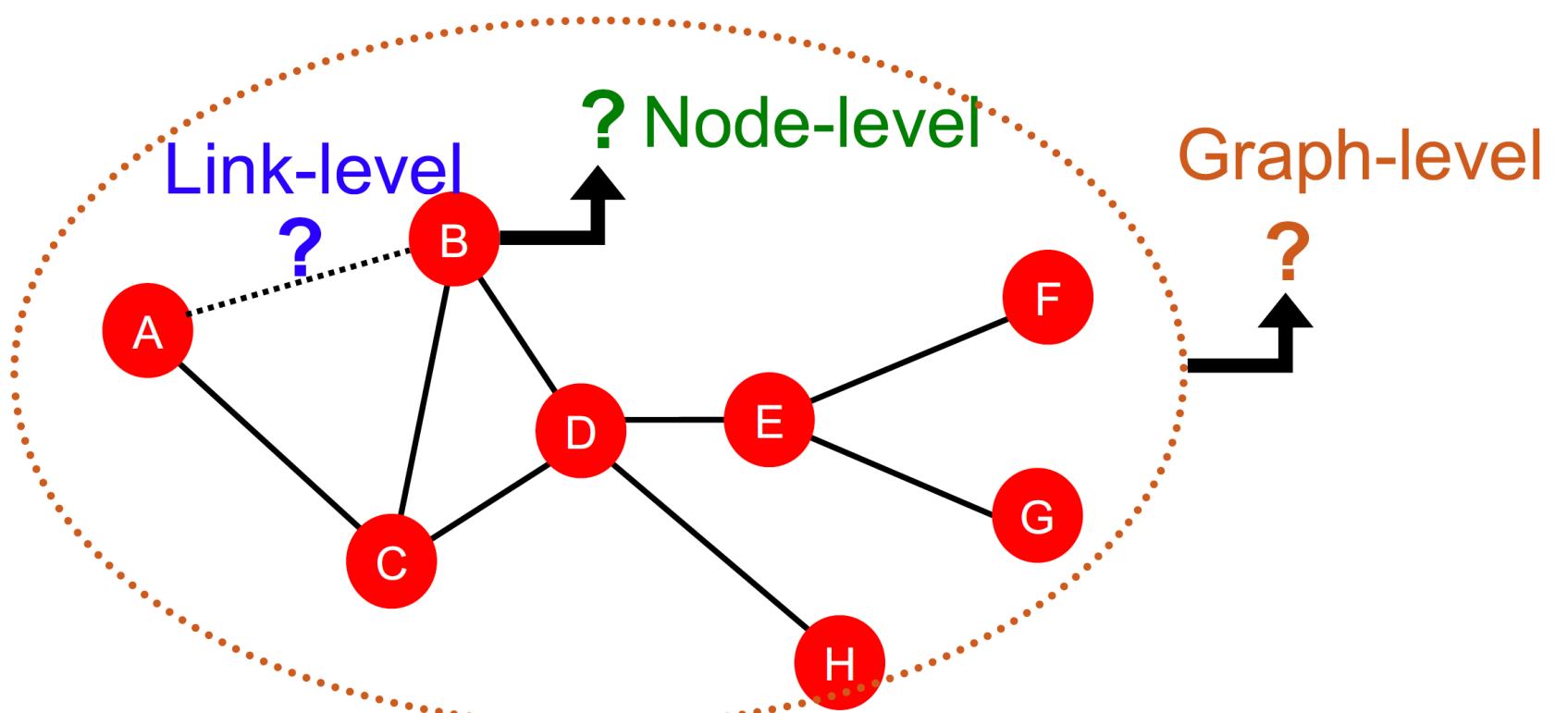


▪ Graphs are very complex

- Arbitrary structures (no spatial locality like grids / no fixed orderings)
- Heterogeneous: Directed/undirected, binary/weighted/typed, multimodal features
- Large-scale: More than millions of nodes and billions of edges

Typical tasks

- **Node-level** prediction
- **Edge-level** prediction
- **Graph-level** prediction



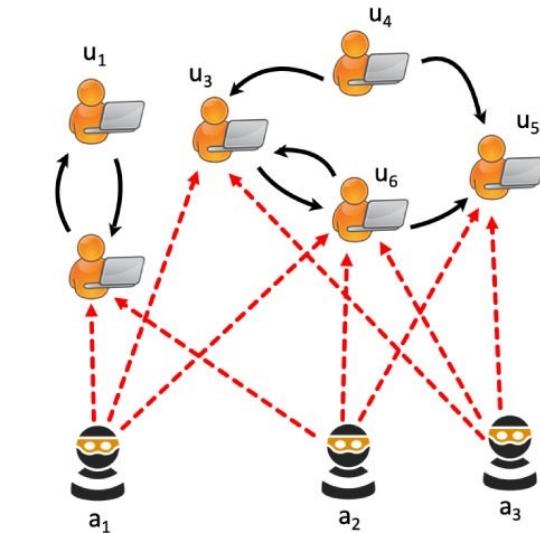
Typical tasks

▪ Node-level tasks (or edge-level tasks)

- Node label classification, including node-level anomaly detection
- Node label regression
- Link label binary classification, i.e., link prediction
- Link label multi-class classification, i.e., relation classification

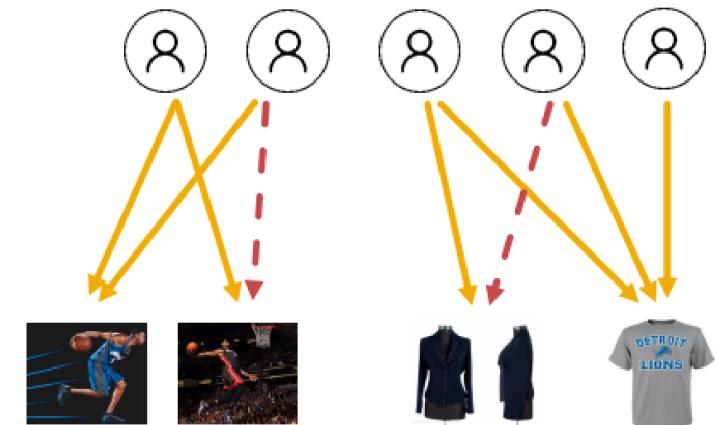


- Social network analysis (e.g., demographic info prediction)
- Spam / fraud detection (e.g., transaction networks)
- Link prediction (e.g., social networks, chemical interaction networks, biological networks, transportation networks)
- Knowledge graph population / completion / relation reasoning
- Recommender system (bipartite graphs, hyper-graphs)



Users

Items



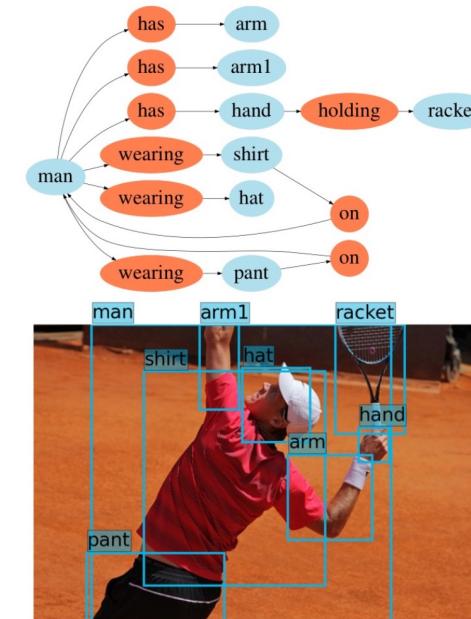
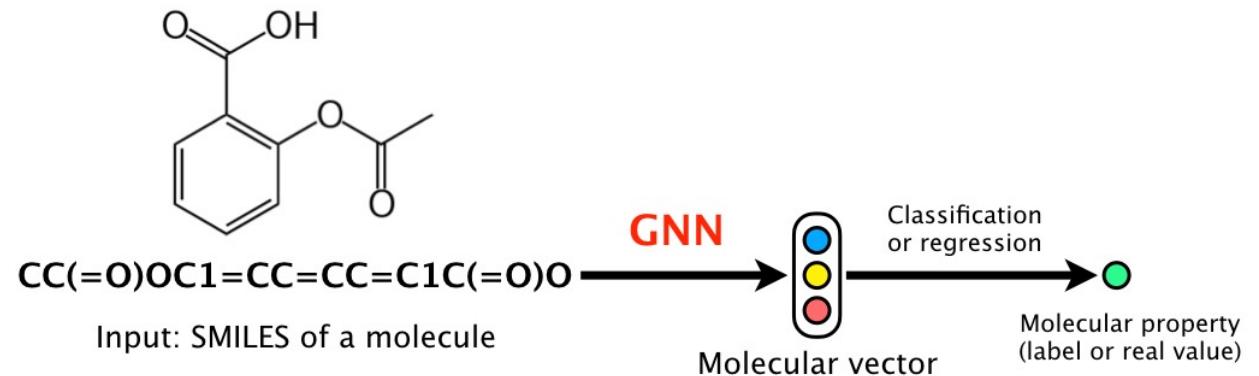
Typical tasks

▪ Graph-level tasks

- Graph label classification
- Graph label regression



- Molecular property prediction
- Drug discovery
- Scene understanding (i.e., objects graph)

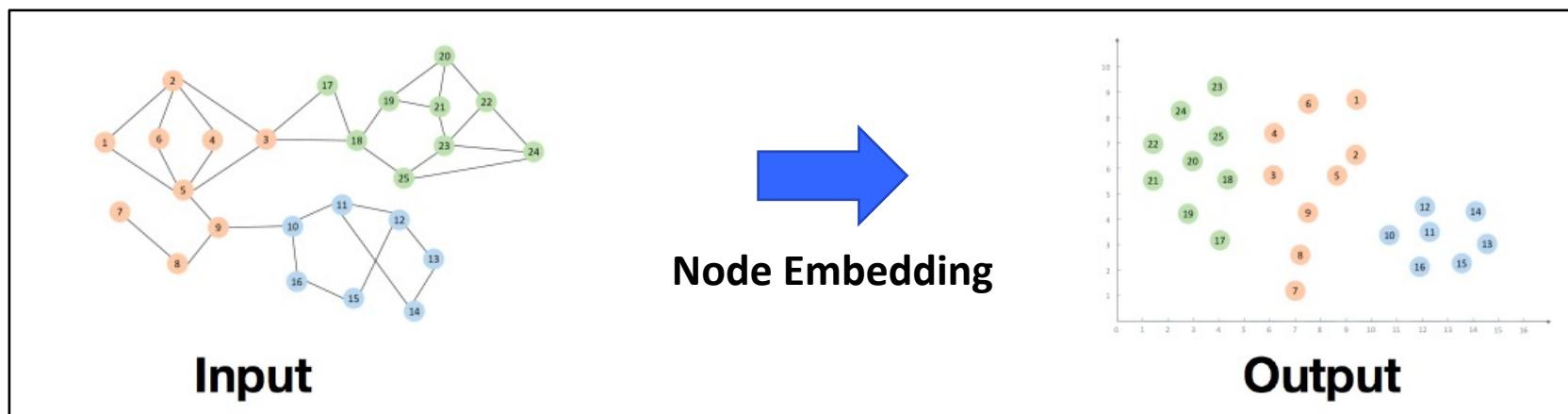
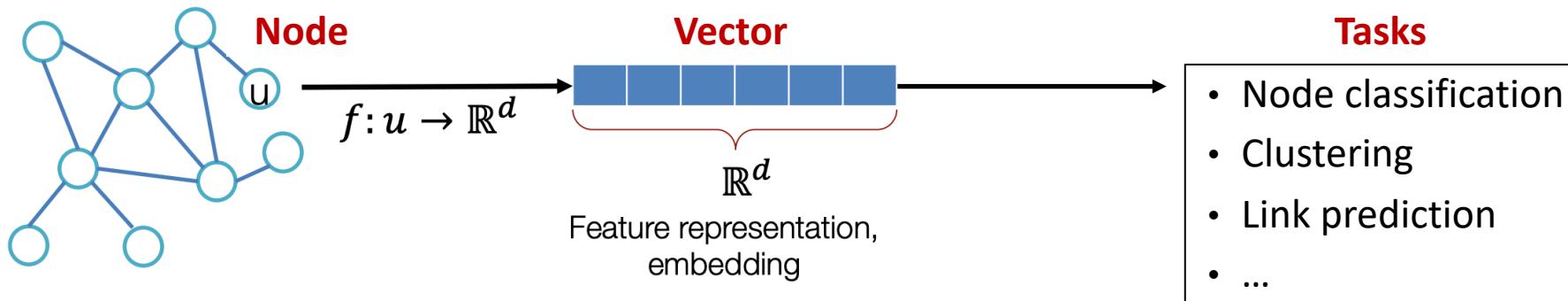


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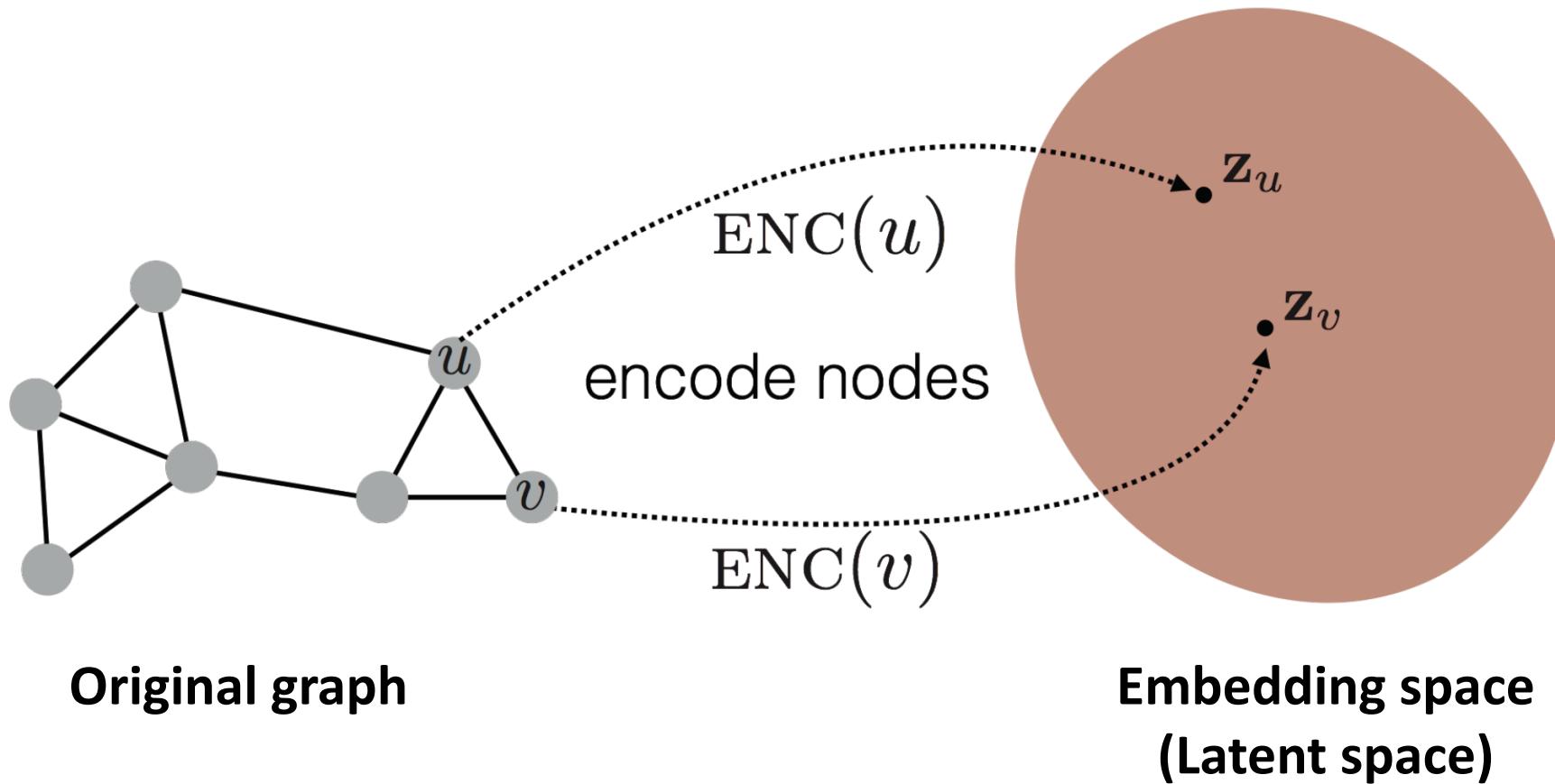
GRAPH REPRESENTATION LEARNING

- Goal: Encode nodes so that **similarity in the embedding space** approximates **similarity in the original network**
- Similar nodes in a network have similar vector representations



NODE EMBEDDING

- Main idea: Encode nodes so that **similarity in the embedding space** approximates **similarity in the graph**



- Two things to consider
 - 1. How to encode nodes?
 - Encoder
 - 2. How to define similarity in the embedding space?
 - Decoder (Similarity function)

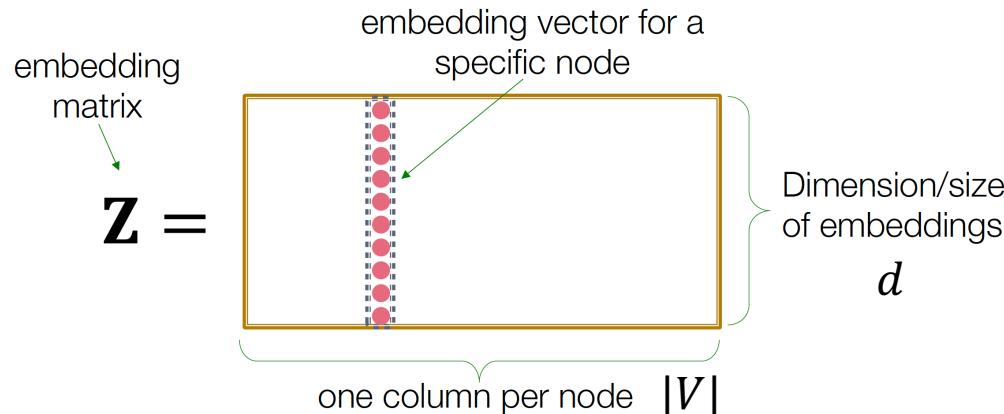
ENCODER

- Maps each node to a low-dimensional vector

$$ENC(v) = z_v \quad \begin{matrix} \text{d-dimensional embedding vector} \\ \uparrow \\ \text{Node in the input graph} \end{matrix}$$

- Simplest encoding approach:** Encoder is just an embedding-lookup (Shallow model)

$$ENC(v) = z_v = Z \cdot v \quad \begin{matrix} Z \in R^{d \times |V|} \\ v \in I^{|V|} \end{matrix}$$



$$v = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 1 \\ 0 \\ 0 \\ 0 \\ 0 \end{bmatrix} \quad \begin{matrix} \text{ex) node 4} \\ \uparrow \end{matrix}$$

Each node is assigned a unique embedding vector (i.e., we directly optimize the embedding of each node)

DECODER (SIMILARITY FUNCTION)

- Specifies how the **relationships in the original graph** map to **relationships in embedding space**

Relationships in the original graph

$$\text{similarity } (u, v)$$

Similarity between
node u and node v in
the original network



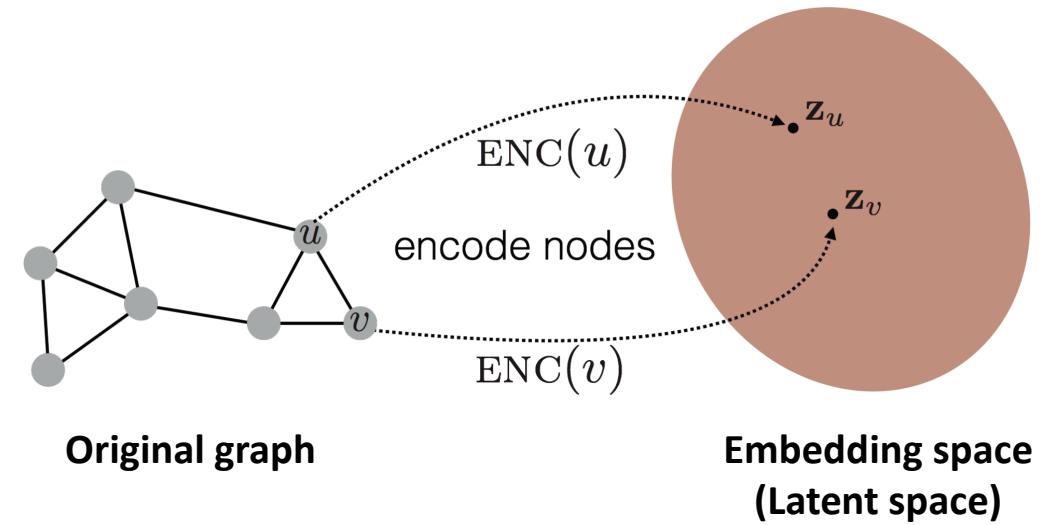
Relationships in embedding space

$$\mathbf{z}_v^T \mathbf{z}_u$$

Dot product between
embeddings of node u
and node v

ENCODER + DECODER FRAMEWORK

- **Encoder:** Embedding look-up (Shallow model)
 - Deep encoders (GNNs) later in the lecture
- **Decoder:** Based on dot product



Objective

Maximize $\mathbf{z}_v^T \mathbf{z}_u$ for node pairs (u, v) that are **similar**

- How can we define **node similarity**?

- Possible choice

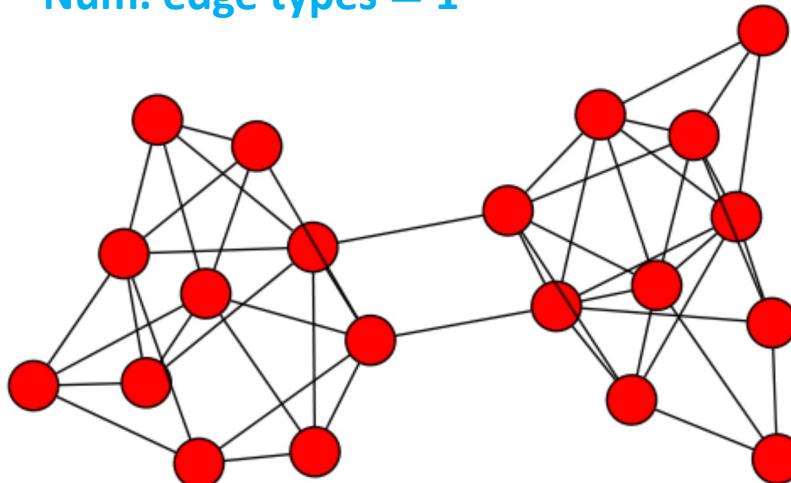
- Are two nodes linked?
- Do they share neighbors?
- Do they have similar structural roles?
- ...

RECALL: HOMOGENEOUS NETWORK

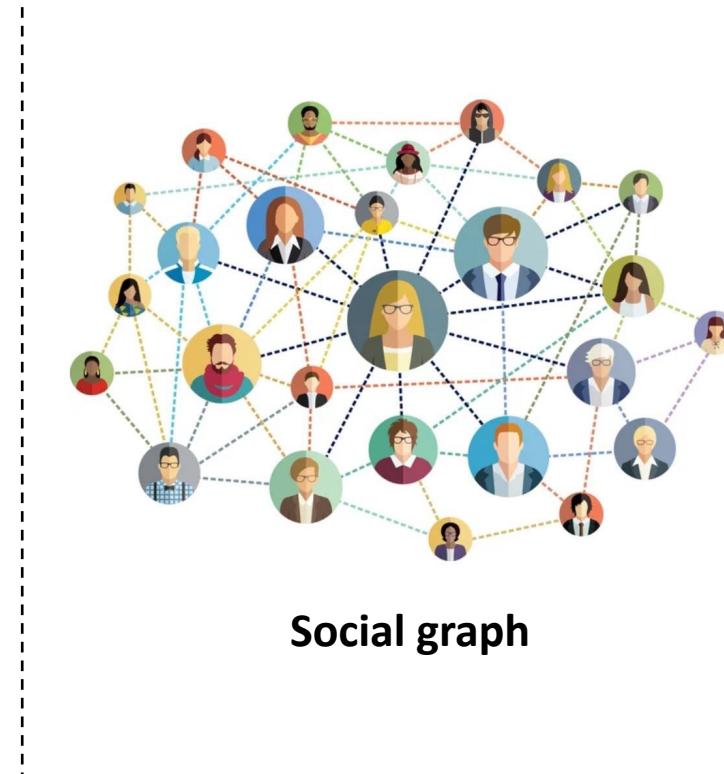
- A graph with a single type of node and a single type of edge

Num. node types = 1

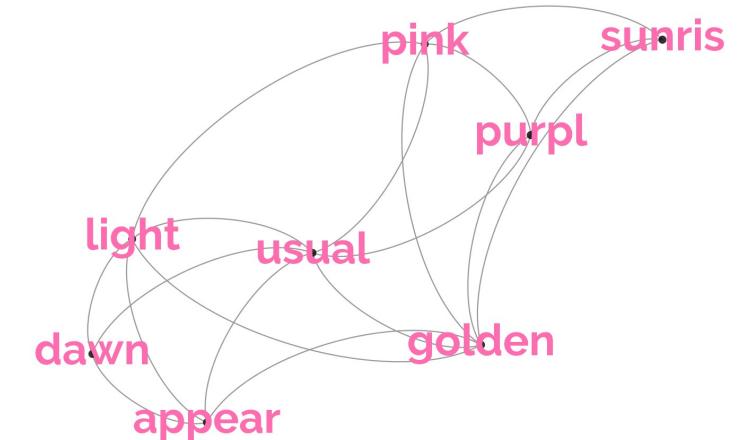
Num. edge types = 1



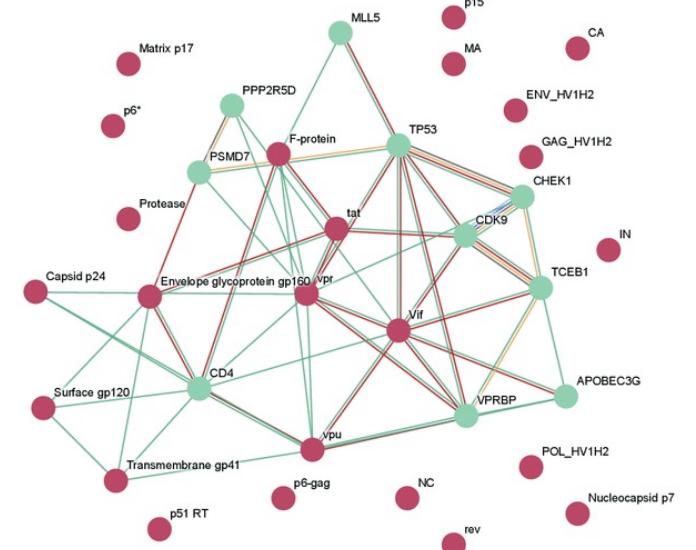
Homogeneous network



Social graph



Word cooccurrence graph



Protein-Protein Interaction Graph

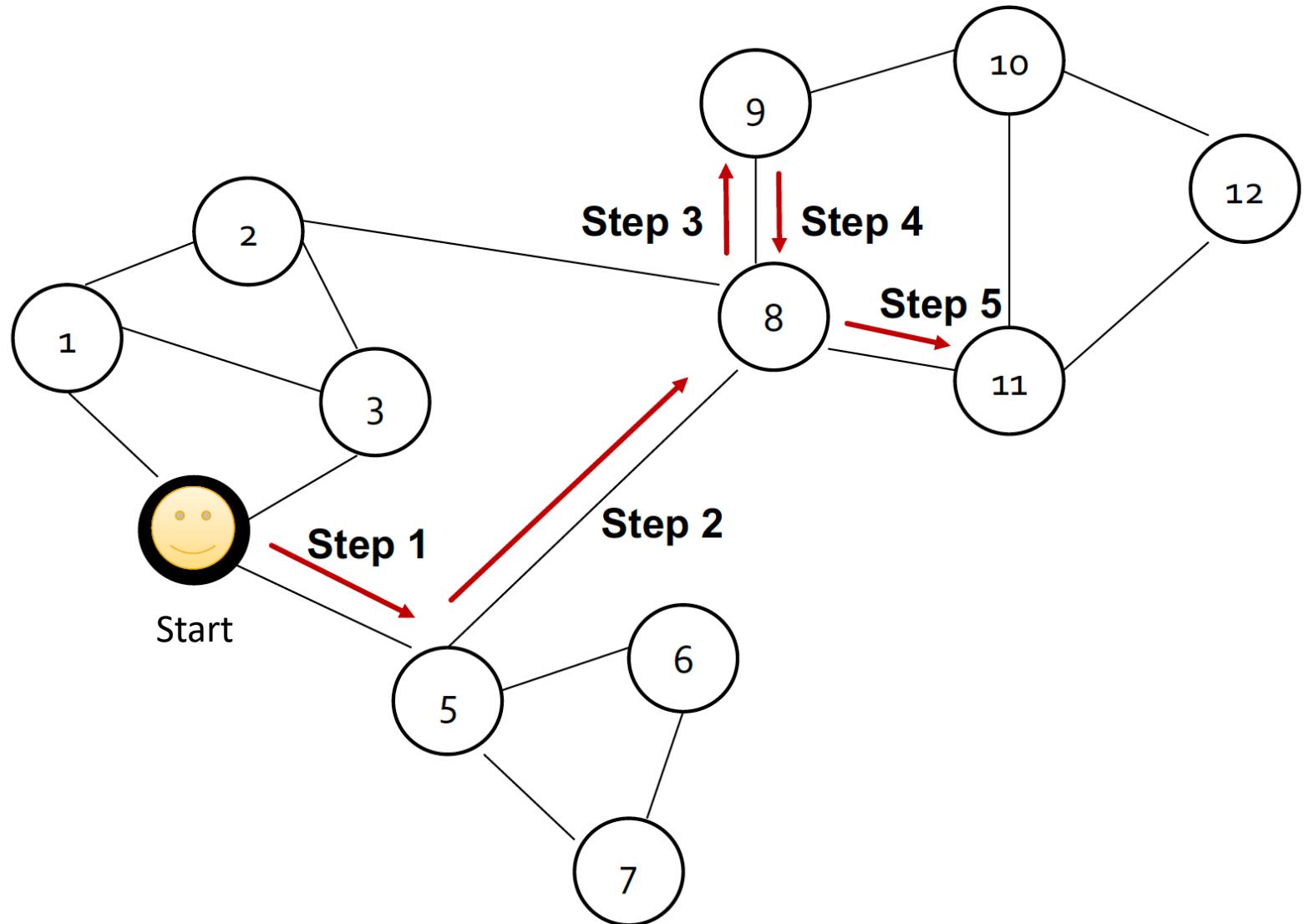
(Figure credit) <https://medium.com/analytics-vidhya/social-network-analytics-f082f4e21b16>

<https://www.researchgate.net/publication/327854066/figure/fig2/AS:674567748075520@1537840892354/HIV-1-and-Homo-sapiens-interaction-network-in-virusesSTRING-HIV-1-and-Homo-sapiens.png>

[https://commons.wikimedia.org/wiki/File:Word_co-occurrence_network_\(range_3_words\)_-_ENG.jpg](https://commons.wikimedia.org/wiki/File:Word_co-occurrence_network_(range_3_words)_-_ENG.jpg)

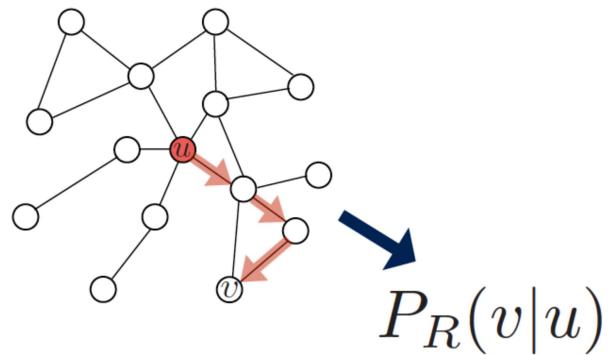
WHAT IS RANDOM WALK?

- Given a graph and a starting node,
 - 1. Select a neighbor of it at random,
 - 2. Move to this neighbor
 - Repeat 1,2
- Example of random walk
 - Start \rightarrow 5 \rightarrow 8 \rightarrow 9 \rightarrow 8 \rightarrow 11
 - (Random) Sequence of nodes



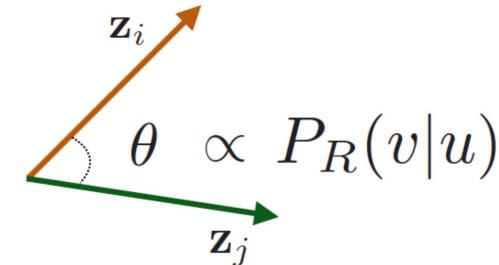
RANDOM WALK-BASED NODE EMBEDDINGS: OVERVIEW

- Idea: Learn node embedding such that nearby nodes in the graph are close together in the embedding space
- Q. Given a node u , how do we define nearby nodes?
- A. Through **random walk**!
- Step 1. Estimate the probability of visiting node v on a random walk starting from node u using some random walk strategy R
- Step 2. Optimize embeddings to encode these random walk statistics
 - e.g., If two nodes co-occur, maximize their similarity



Why random walk?

Random walk can reflect both **local** and **high-order** neighborhood information



cf) Dot product = cosine similarity, if node embeddings are unit vectors

RANDOM WALK-BASED NODE EMBEDDINGS: DETAILED ALGORITHM

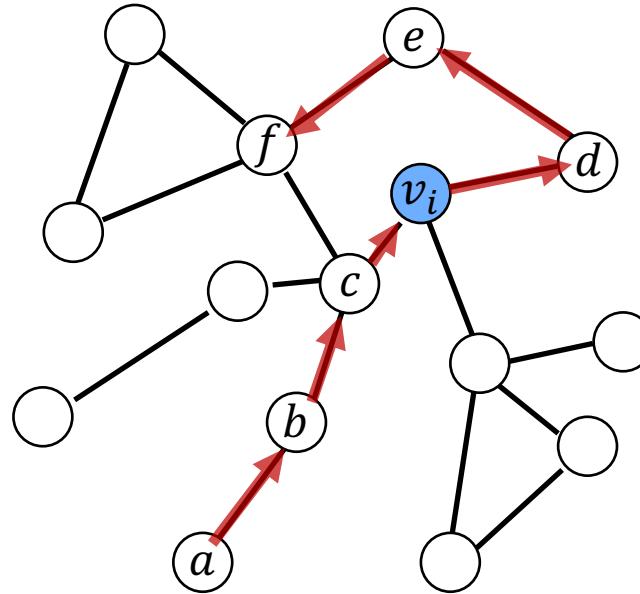
- **Given:** $G = (V, E)$
- **Goal:** To learn a mapping function $f: u \rightarrow R^d$ for $u \in V$
 - $f(u) = \mathbf{z}_u \in R^d$
- **Step 1:** Run fixed-length random walks starting from each node u in the graph using some random walk strategy R
- **Step 2:** For each node u collect $N_R(u)$, the multiset of nodes visited on random walks starting from u
 - $N_R(u)$: Neighboring nodes of node u under random walk strategy R
- **Step 3:** Optimize embeddings according to the following objective
 - Objective: Given node u , predict its neighbors $N_R(u)$

$$\max_f \sum_{u \in V} \log P(N_R(u) | \mathbf{z}_u) \quad (\text{Maximum likelihood objective})$$

Given node u , we aim to maximize the probability of its neighboring nodes

i.e., we want to learn embedding of node u that is predictive of its neighboring nodes

HOW TO DEFINE NEIGHBORING NODES?



Random walk strategy

Example sequence

Window size=2

R

$a \rightarrow b \rightarrow c \rightarrow v_i \rightarrow d \rightarrow e \rightarrow f$

$a \rightarrow \boxed{b} \rightarrow c \rightarrow v_i \rightarrow d \rightarrow e \rightarrow f$

Center node

v_i

Neighborhood

$N_R(v_i) = b, c, d, e$

RANDOM WALK-BASED NODE EMBEDDINGS: OPTIMIZATION

$$\max_f \sum_{u \in V} \log P(N_R(u) | \mathbf{z}_u) \quad \xrightarrow{\text{Equivalent}} \quad L = \sum_{u \in V} \sum_{v \in N_R(u)} -\log(P(v | \mathbf{z}_u))$$

- **Intuition:** Optimize embeddings \mathbf{z}_u to maximize the likelihood of random walk co-occurrences
- **Approach:** Parameterize $P(v | \mathbf{z}_u)$ using **softmax**

$$P(v | \mathbf{z}_u) = \frac{\exp(\mathbf{z}_u^T \mathbf{z}_v)}{\sum_{j \in V} \exp(\mathbf{z}_u^T \mathbf{z}_j)}$$

$$L = \sum_{u \in V} \sum_{v \in N_R(u)} -\log\left(\frac{\exp(\mathbf{z}_u^T \mathbf{z}_v)}{\sum_{j \in V} \exp(\mathbf{z}_u^T \mathbf{z}_j)}\right)$$

Sum over
all nodes u

Sum over nodes v seen
on random walks
starting from u

Predicted probability of
 u and v cooccurring on
random walk

Optimizing random walk embeddings
= Finding embeddings \mathbf{z}_u that
minimizes the loss L

NEGATIVE SAMPLING

- But, optimizing the loss L is expensive!

- $O(|V|^2)$ complexity
- We can approximate this normalization term

$$L = \sum_{u \in V} \sum_{v \in N_R(u)} -\log\left(\frac{\exp(\mathbf{z}_u^T \mathbf{z}_v)}{\sum_{j \in V} \exp(\mathbf{z}_u^T \mathbf{z}_j)}\right)$$

$$\log\left(\frac{\exp(\mathbf{z}_u^T \mathbf{z}_v)}{\sum_{j \in V} \exp(\mathbf{z}_u^T \mathbf{z}_j)}\right) \approx \log(\sigma(\mathbf{z}_u^T \mathbf{z}_v)) - \sum_{j=1}^k \log(\sigma(\mathbf{z}_u^T \mathbf{z}_j)), \quad j \sim P_V$$

- $\sigma(x) = \frac{1}{1+e^{-x}}$ (Sigmoid function)
 - Makes each term a “probability” between 0 and 1
- P_V : Random distribution over nodes

<https://arxiv.org/pdf/1402.3722.pdf>

- Instead of normalizing w.r.t. all nodes, just normalize against k random “negative samples” j
- How do we sample from P_V to help the training process?
 - Sample k negative nodes considering the degree of each node

RANDOM WALK-BASED NODE EMBEDDINGS: OPTIMIZATION

- After we obtained the objective function, **how do we optimize (minimize) it?**

$$L = \sum_{u \in V} \sum_{v \in N_R(u)} -\log(P(v|\mathbf{z}_u))$$

- Gradient descent:** A simple and the most common way to minimize L

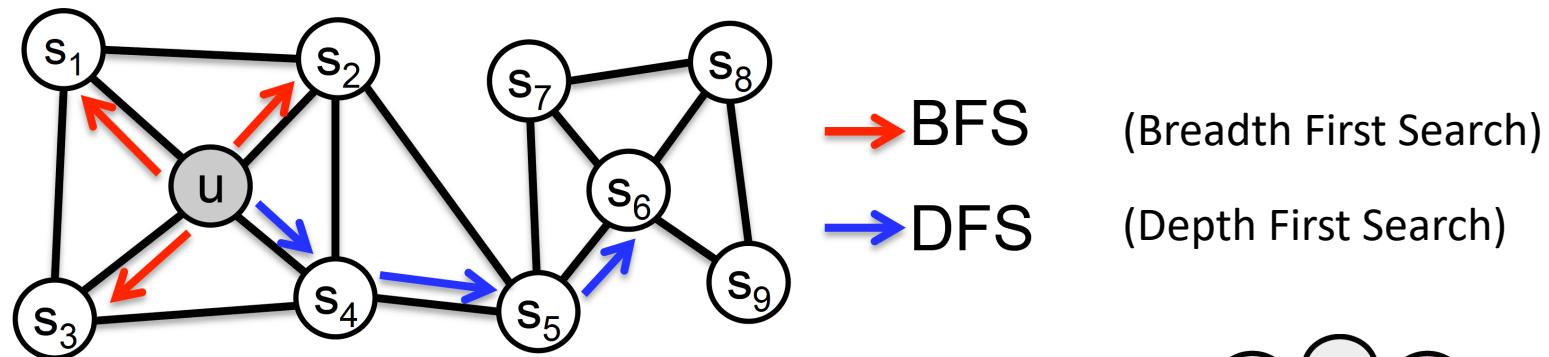
- Step 1: Randomly initialize \mathbf{z}_i for all $i \in V$
- Step 2: Iterate until convergence
 - For all $i \in V$, compute the derivative w.r.t. the loss L , i.e., $\frac{\partial L}{\partial \mathbf{z}_i}$
 - For all $i \in V$, update $\mathbf{z}_i \leftarrow \mathbf{z}_i - \eta \frac{\partial L}{\partial \mathbf{z}_i}$

- Stochastic Gradient descent:** Instead of evaluating gradients over all examples, evaluate for a **single** node

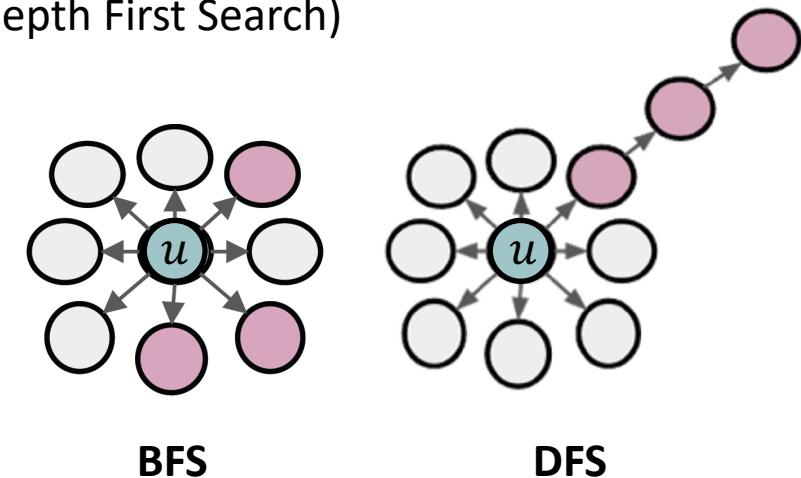
- Step 1: Randomly initialize \mathbf{z}_i for all $i \in V$
- Step 2: Iterate until convergence: $L^{(u)} = \sum_{v \in N_R(u)} -\log(P(v|\mathbf{z}_u))$
 - Sample a node i , for all $j \in N_R(i)$ compute the derivative w.r.t. the loss L , i.e., $\frac{\partial L^{(i)}}{\partial \mathbf{z}_j}$
 - For all $j \in N_R(i)$, update $\mathbf{z}_j \leftarrow \mathbf{z}_j - \eta \frac{\partial L^{(i)}}{\partial \mathbf{z}_j}$

NODE2VEC: BIASED WALKS

- Idea: Use flexible, biased random walks that can trade off between **local** and **global** views of the network
 - Deepwalk's simple random walk mainly focuses on the global view



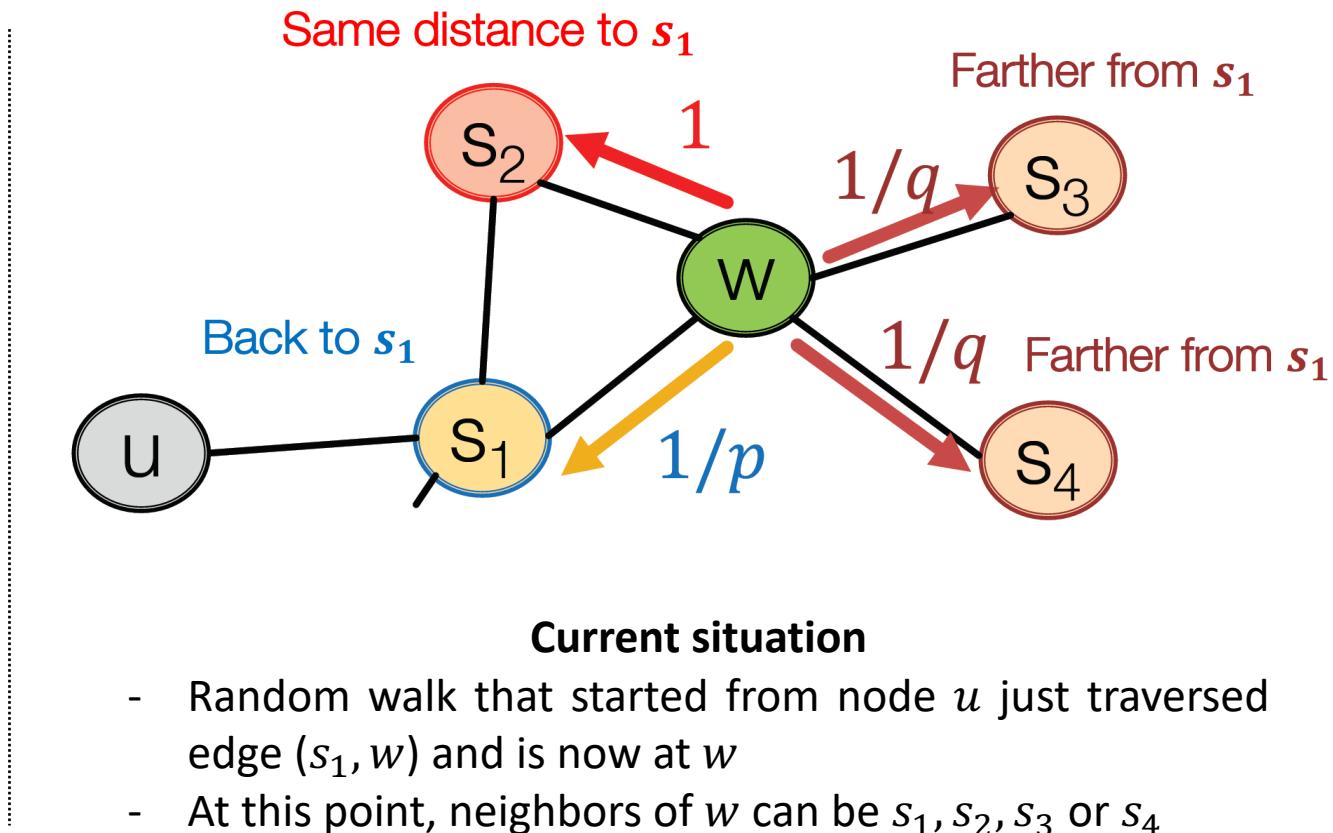
- Two strategies to define a neighborhood $N_R(u)$ of node u : BFS and DFS
- Example: Walk of length 3 from node u
 - $N_{BFS}(u) = \{s_1, s_2, s_3\}$ **Local** microscopic view
 - $N_{DFS}(u) = \{s_4, s_5, s_6\}$ **Global** macroscopic view



How can we interpolate between BFS and DFS?

NODE2VEC: INTERPOLATING BFS AND DFS

- Biased fixed-length random walk R starting from node u generates neighborhood $N_R(u)$
- **Two parameters to control the interpolation**
 - **Return parameter p**
 - Return back to the previous node
 - **In-out parameter q**
 - Moving outwards (DFS) vs. inwards (BFS)
 - Intuitively, q is the “ratio” of BFS vs. DFS

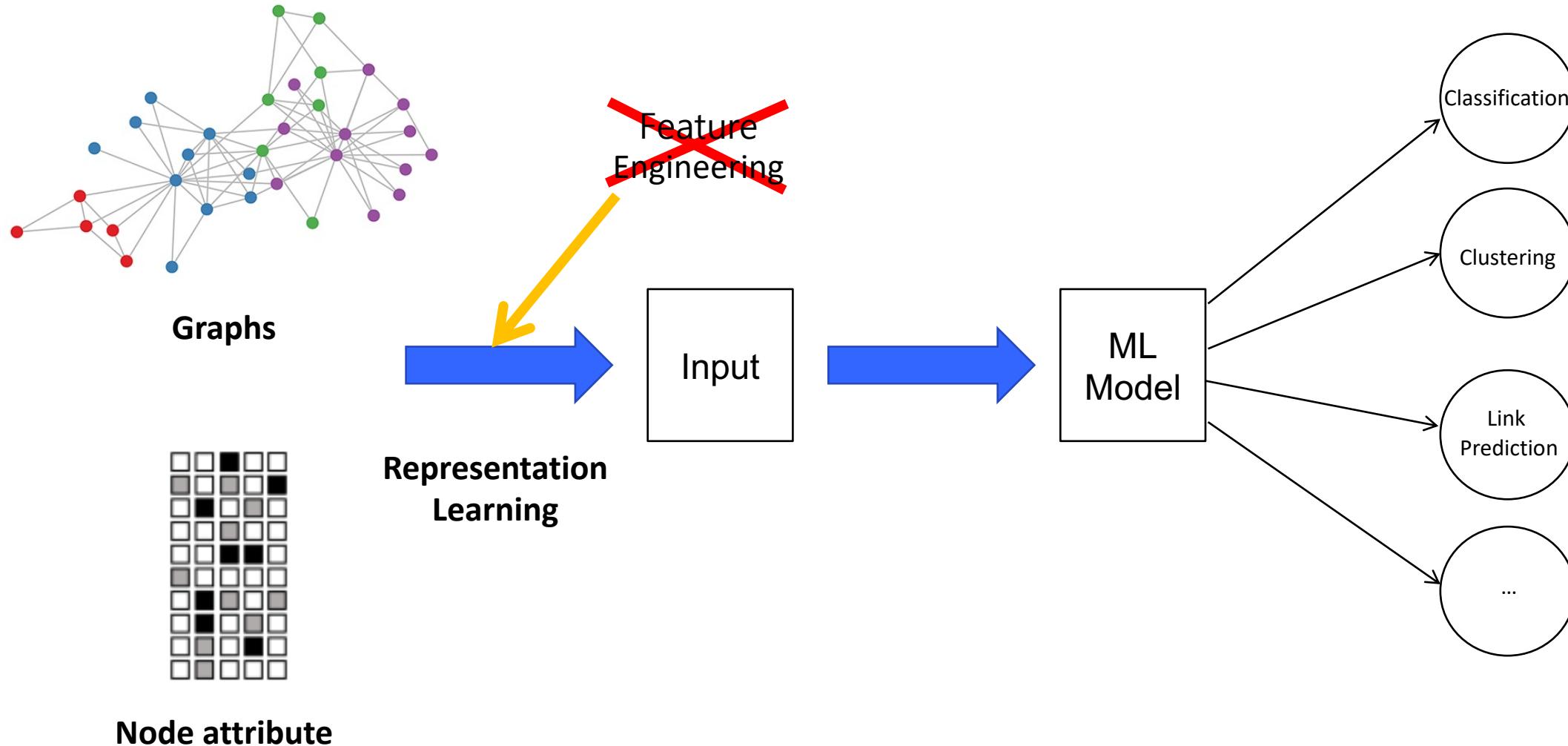


Idea of biased random walk
Remember where the walk came from!

OUTLINE

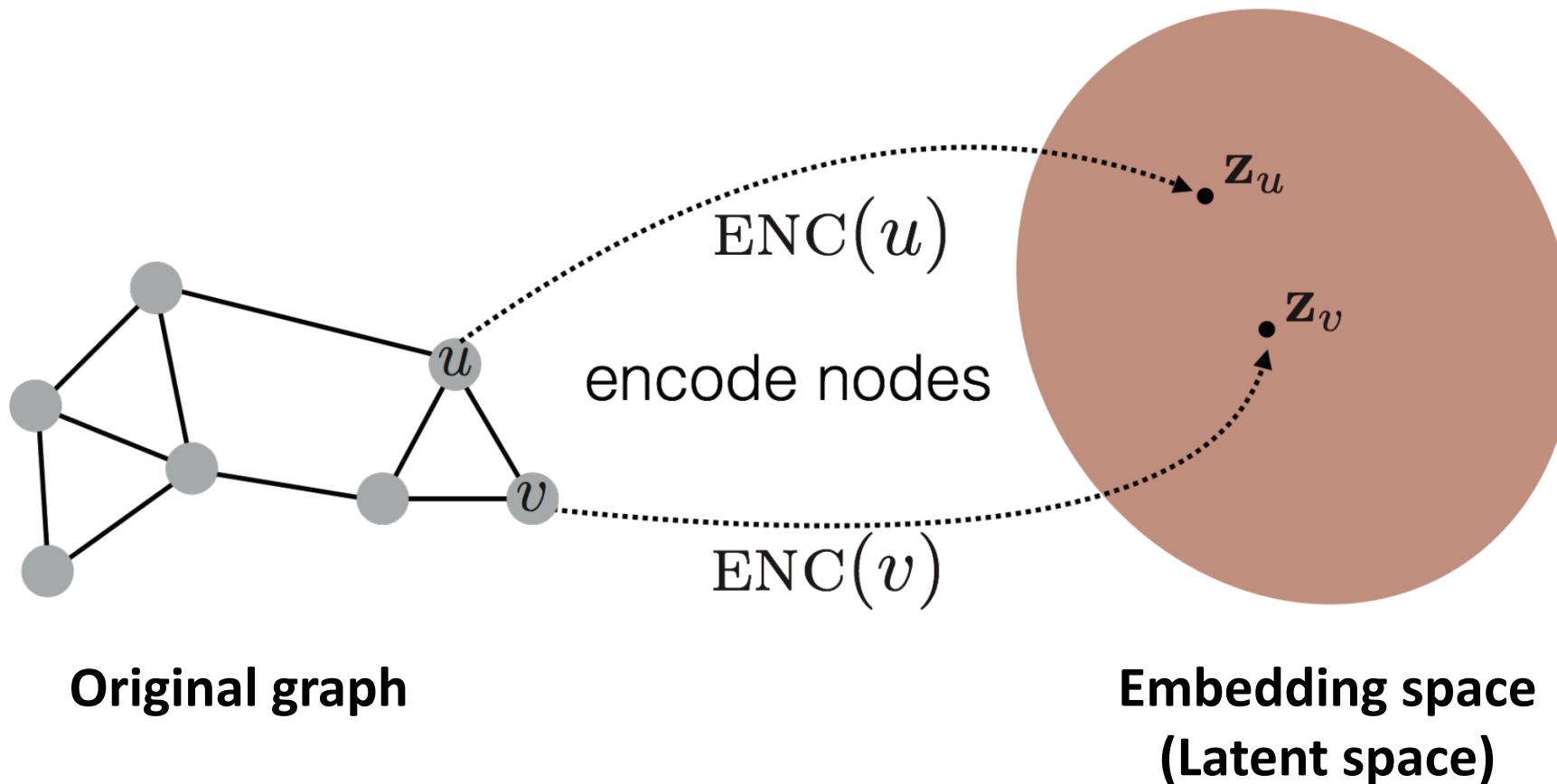
- Overview
- Random walk-based Methods
- Graph Neural Networks (GNNs)
- How to effectively train GNNs?
 - Self-supervised Learning (AAAI'22)
 - Explainable Model (NeurIPS'23)
 - Robustness (KDD'23)
- Applications of GNNs
 - GNNs for Science (화학/소재/생물) (ICML'23)
 - GNNs for Medical Data (멀티모달 의료 데이터 분석) (AAAI'23)
 - GNNs for Computer vision (Scene Understanding) (AAAI'23)

RECALL: MACHINE LEARNING ON GRAPHS



RECAP: NODE EMBEDDING

- Main idea: Encode nodes so that **similarity in the embedding space** approximates **similarity in the graph**



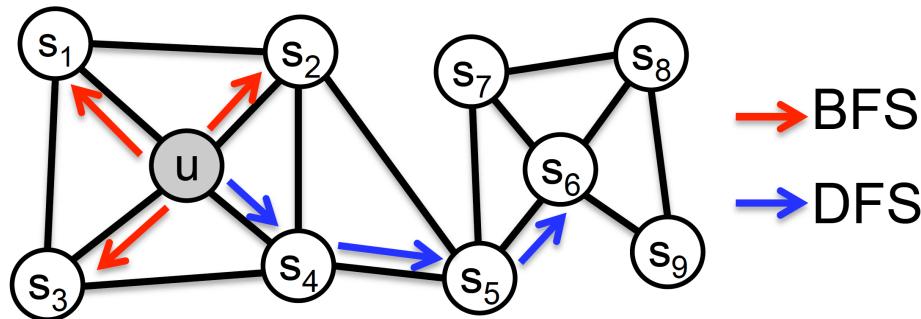
- **Two things to consider**
 - 1. How to encode nodes?
 - Encoder
 - 2. How to define similarity in the embedding space?
 - Decoder (Similarity function)

RECAP: DEEPWALK/NODE2VEC

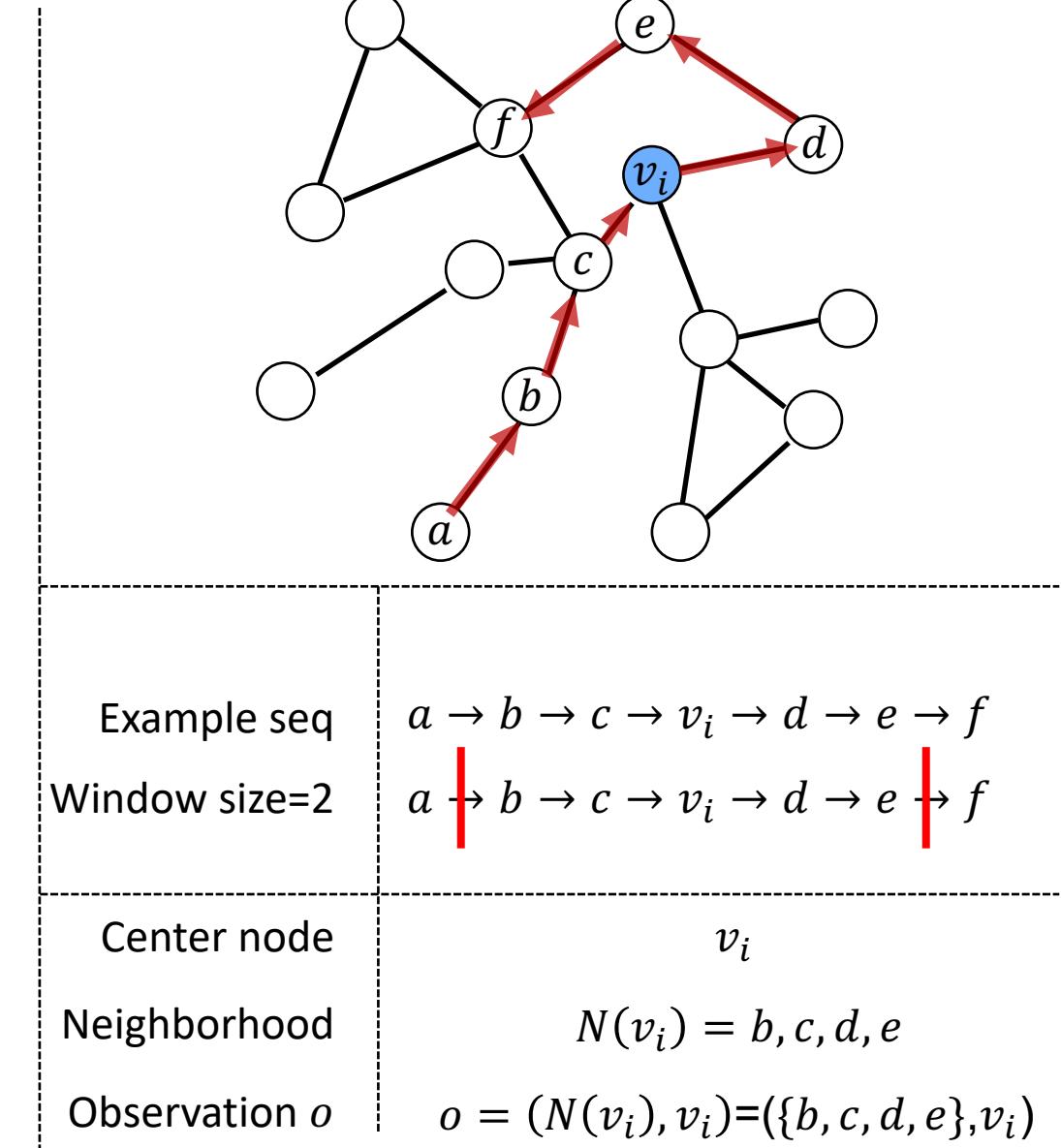
$$\begin{aligned}\mathcal{L}_{DW}(\theta) &= \sum_{o \in O} \log p(o|\theta) = \sum_{o \in O} \log p((N(v_i), v_i)|\theta) \\ &= \sum_{o \in O} \sum_{v_j \in N(v_i)} \log p(v_j|v_i),\end{aligned}$$

- \mathcal{O} : The set of all observations obtained from random walks
- $o = (N(v_i), v_i) \in \mathcal{O}$
 - Center node v_i
 - Neighboring nodes $N(v_i)$

▪ Biased random walk



BFS
DFS



RECALL: ENCODER

- Maps each node to a low-dimensional vector

$$ENC(v) = z_v \quad \begin{matrix} \text{d-dimensional embedding vector} \\ \uparrow \\ \text{Node in the input graph} \end{matrix}$$

- Simplest encoding approach:** Encoder is just an embedding-lookup (Shallow model)

$$ENC(v) = z_v = Z \cdot v$$

$Z \in R^{d \times |V|}$
 $v \in I^{|V|}$

embedding matrix
embedding vector for a specific node
Dimension/size of embeddings d
one column per node $|V|$

$$v = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 1 \\ 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

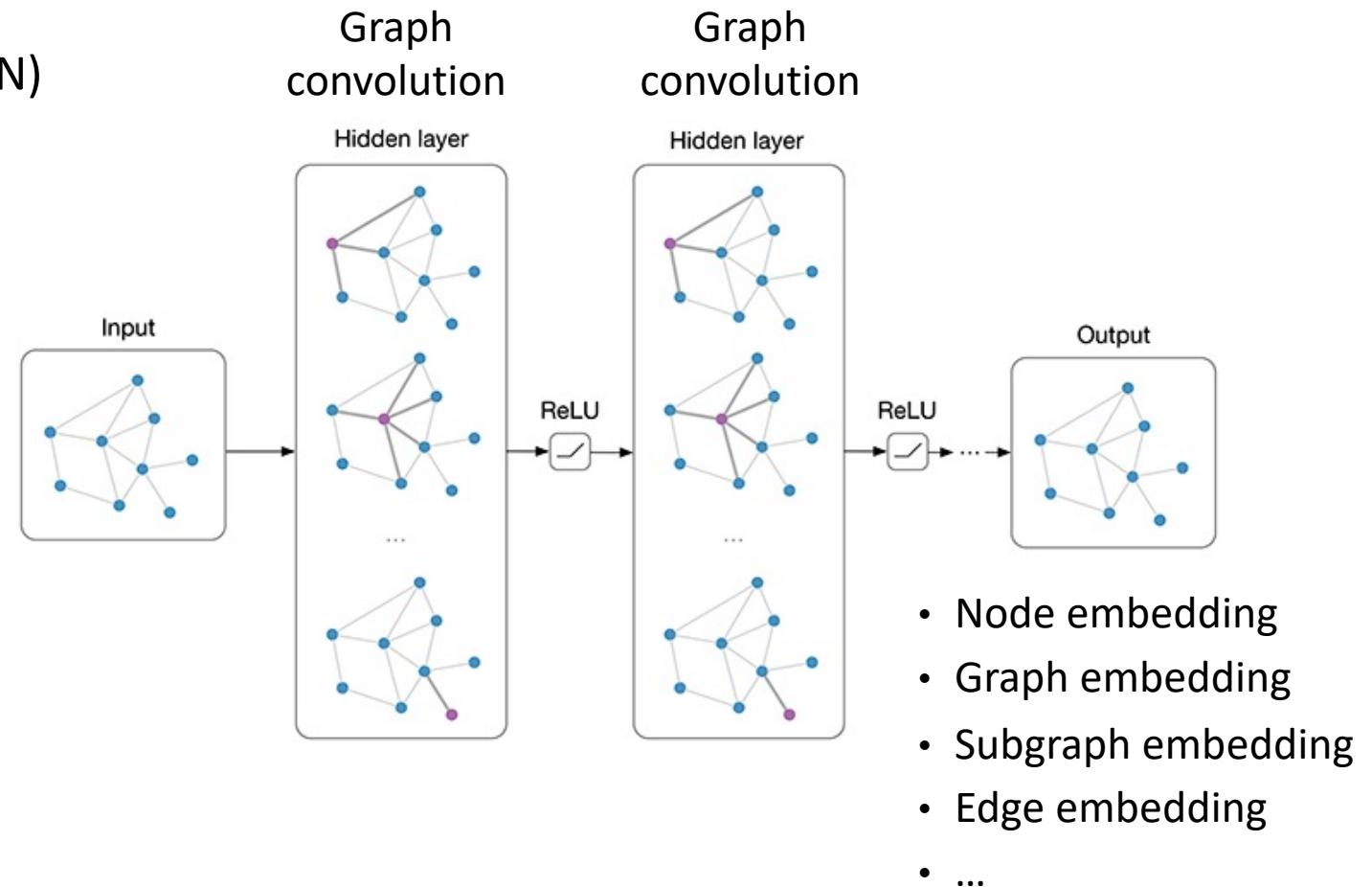
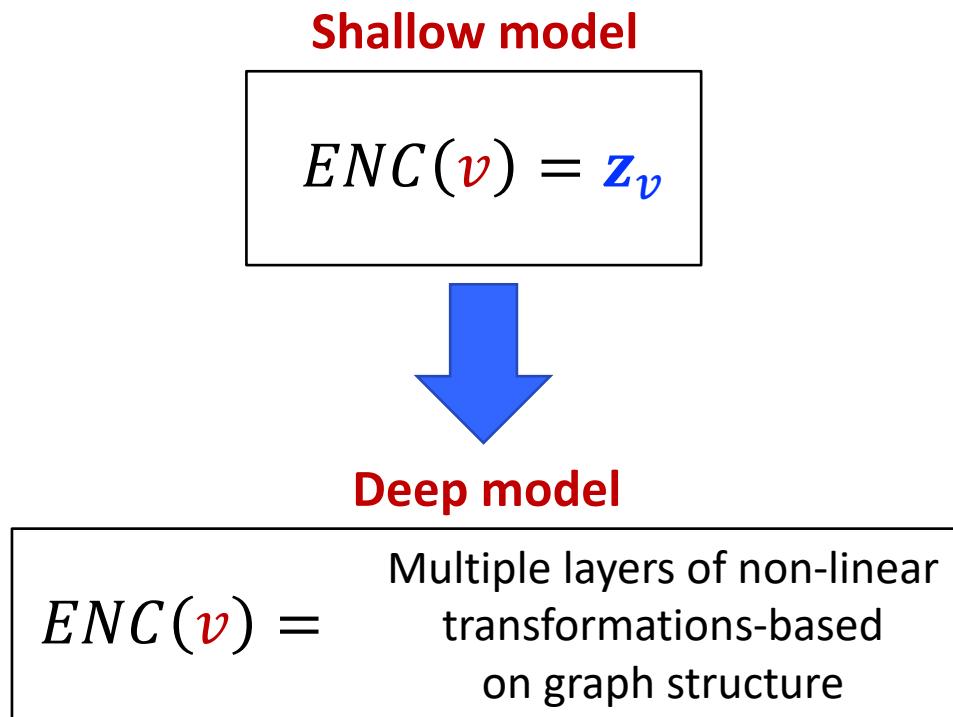
ex) node 4

Each node is assigned a unique embedding vector (i.e., we directly optimize the embedding of each node)

From now on: Deep encoder

DEEP GRAPH ENCODER

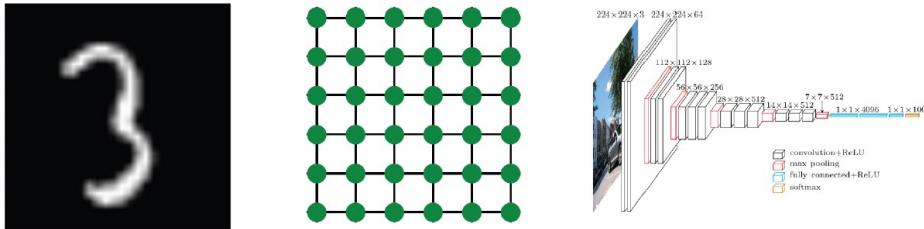
- Deep Encoder = Graph neural network (GNN)



Can we use existing deep learning models? e.g., CNN, RNN, etc

RECAP: CHALLENGES OF GRAPH REPRESENTATION LEARNING

- Existing deep neural networks are designed for data with regular-structure (grid or sequence)
 - CNNs for fixed-size images/grids ...



- RNNs for text/sequences ...

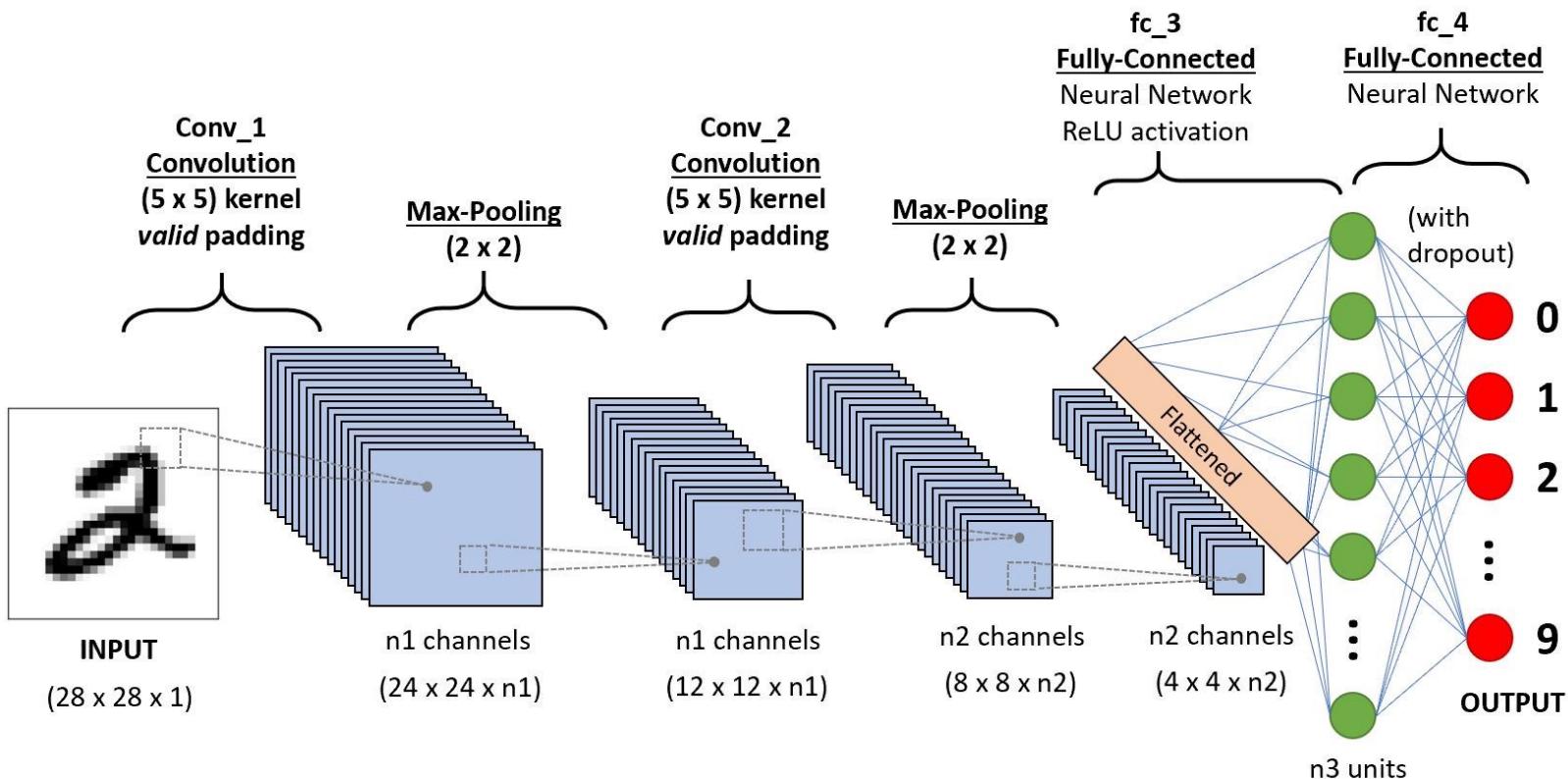


▪ Graphs are very complex

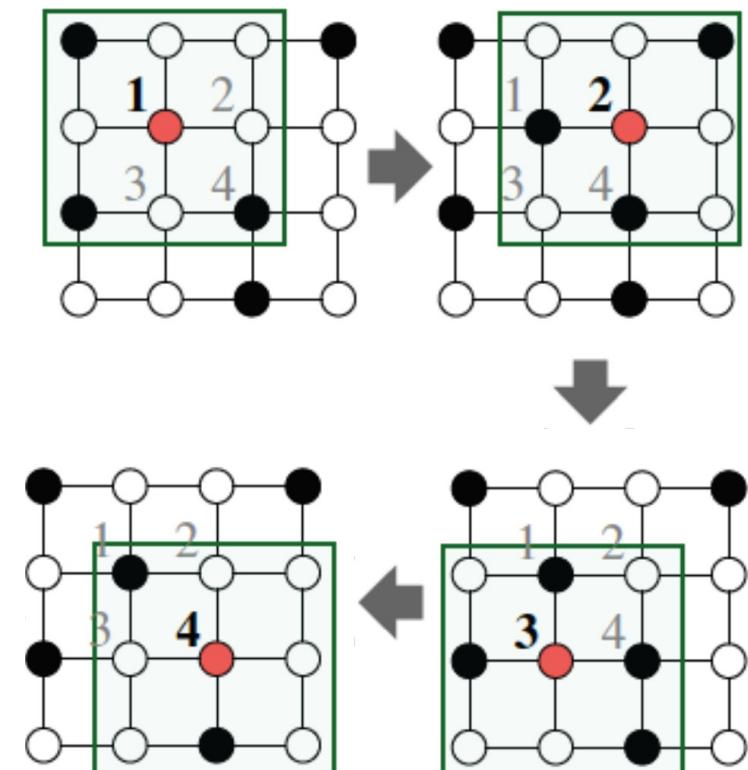
- Arbitrary structures (no spatial locality like grids / no fixed orderings)
- Heterogeneous: Directed/undirected, binary/weighted/typed, multimodal features
- Large-scale: More than millions of nodes and billions of edges

BACKGROUND: CONVOLUTIONAL NEURAL NETWORKS FOR IMAGES

- Convolutional filters
 - Local feature detectors
 - A feature is learned in each **local receptive field** by a convolutional filter

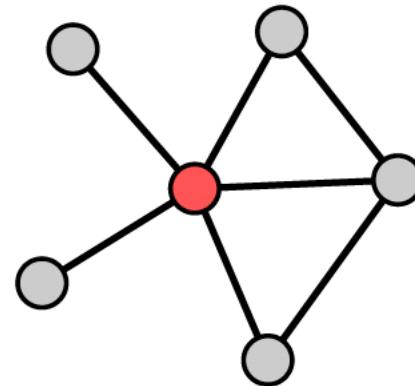
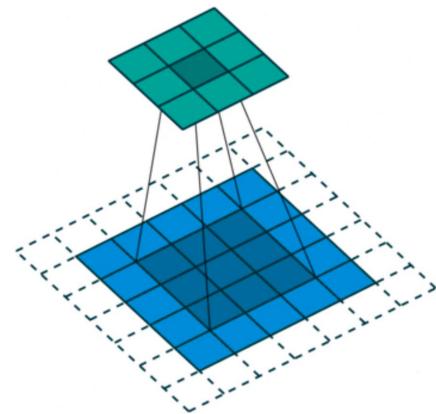
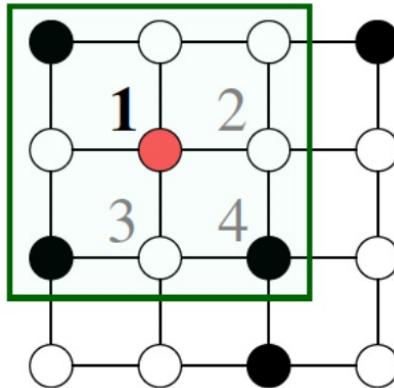


CNN on an image



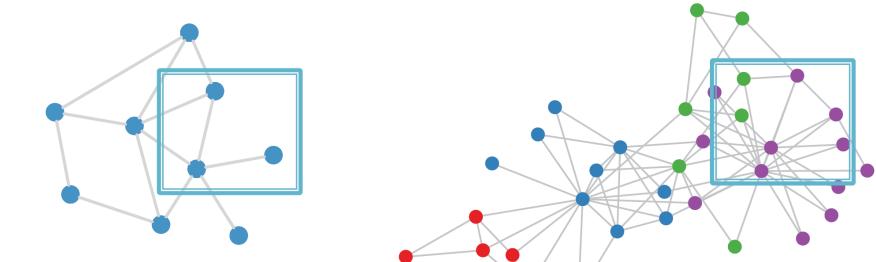
FROM IMAGES TO GRAPHS: LOCAL RECEPTIVE FIELD ON GRAPHS

- How should we define local receptive fields on graphs?
 - Local subgraphs



Graph

Graphs look like this

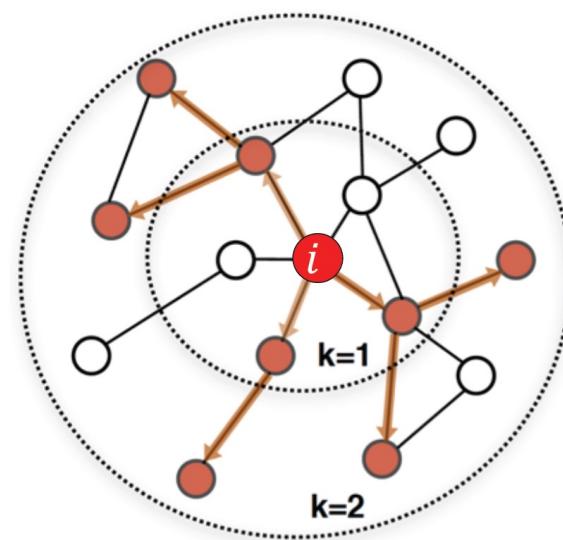


- There is no fixed notion of locality or sliding window on the graph
- No order among neighboring nodes
 - Permutation invariant

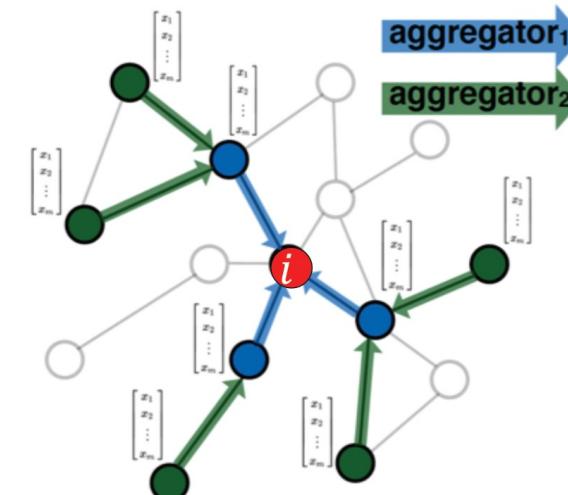
- Idea: Transform information from the neighboring nodes and combine it
 - Step 1: For each node v_i , transform “messages” from neighbors $N(i)$
 - $W_j h_j$ for $v_j \in N(i)$, h_j : “Message” from v_j
 - Step 2: Add them up: $\sum_{v_j \in N(i)} W_j h_j$

GRAPH CONVOLUTIONAL NETWORK (GCN)

- Idea: Node's neighborhood defines a computation graph
 - Messages contain **relational information** + **attribute information**



Determine node
computation graph

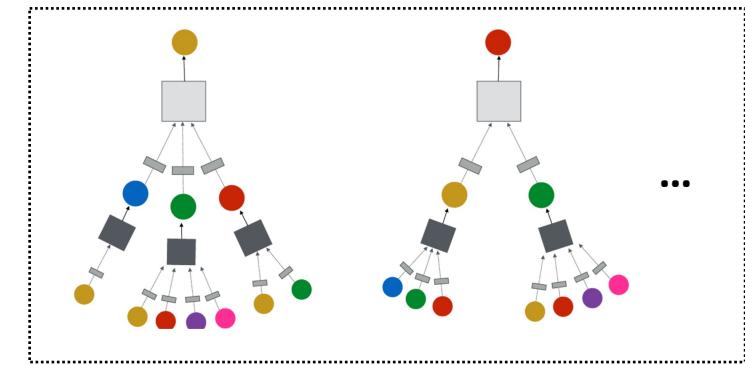


Propagate messages and
transform information

Learn how to propagate information across the graph to compute node features

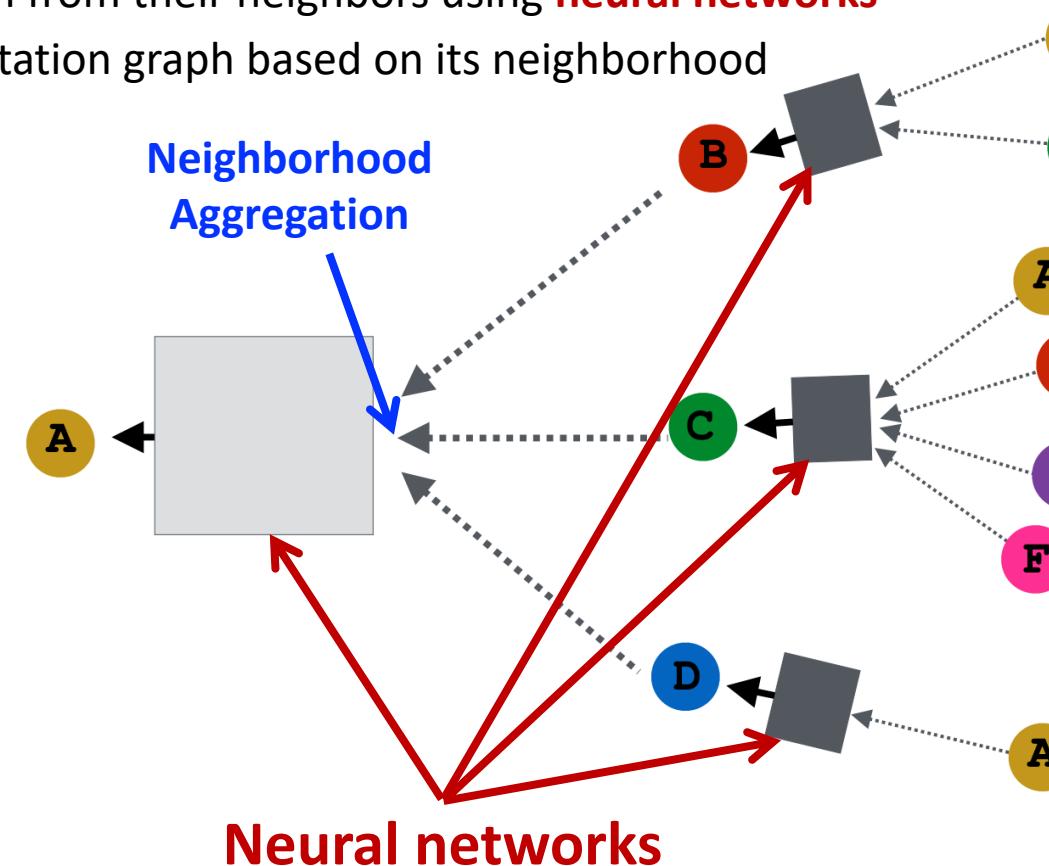
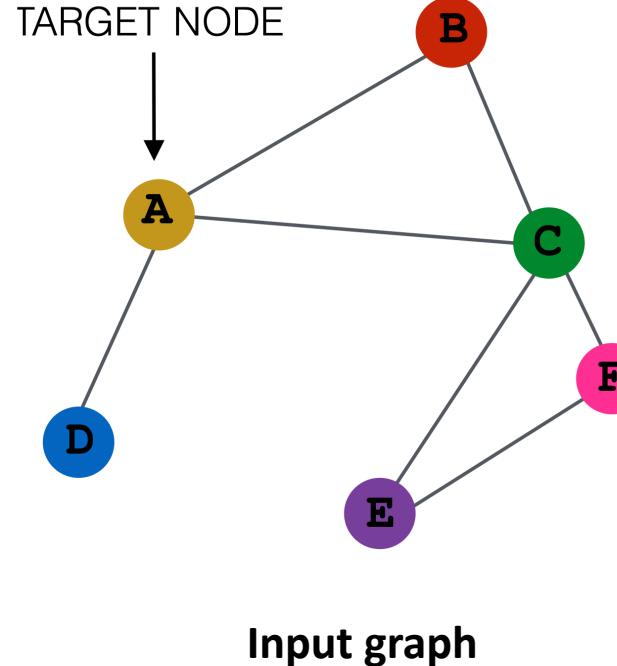
GCN: NEIGHBORHOOD AGGREGATION

- Generate node embeddings based on local network neighborhoods



- **Neighborhood aggregation**

- Nodes aggregate information from their neighbors using **neural networks**
- Every node defines a computation graph based on its neighborhood



- **Things to consider**
- 1. What kind of neural network?
 - 2. How do we aggregate neighboring nodes?

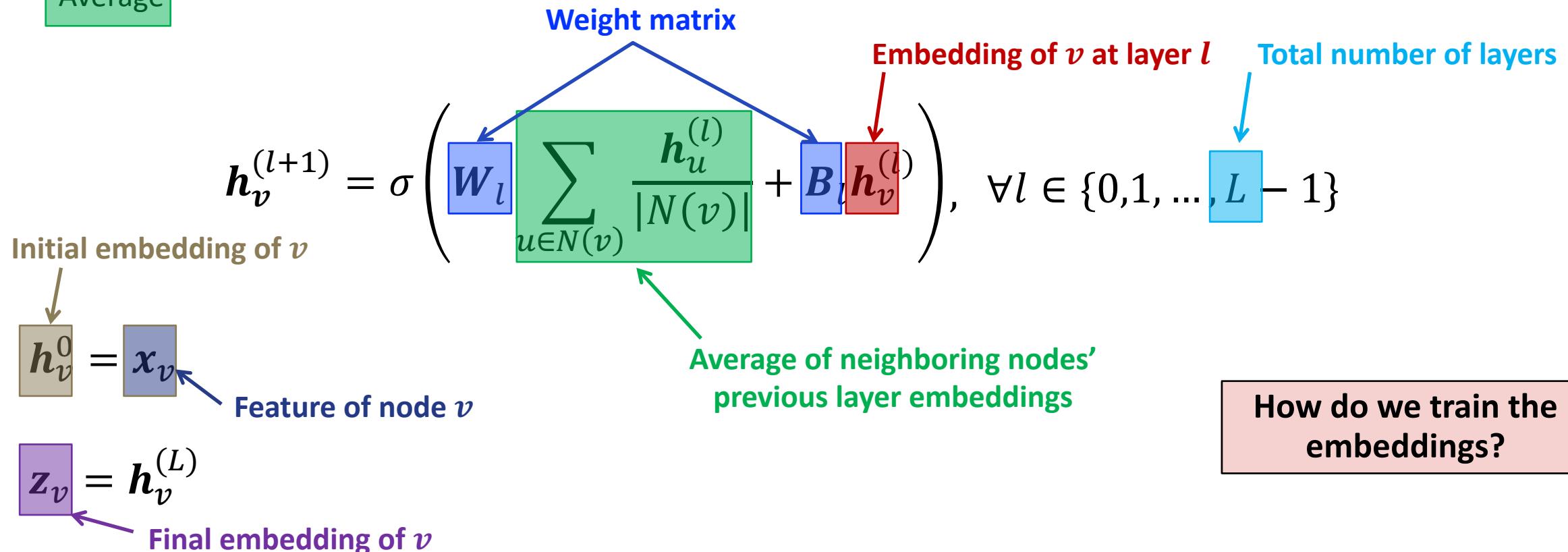
GCN: BASIC APPROACH

- 1. What kind of neural network?

- Simple multiplication of weight matrices (B and W)

- 2. What kind of aggregation?

- Average



What kind of neural network?

Simple multiplication of weight matrices (B and W)

...

How do we train the embeddings?

GCN: MATRIX FORMULATION

- GCN can be efficiently computed in a matrix form

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v)} \frac{\mathbf{h}_u^{(l)}}{|N(v)|} + \mathbf{B}_l \mathbf{h}_v^{(l)} \right), \quad h_v^0 = x_v, \quad z_v = h_v^{(L)}$$



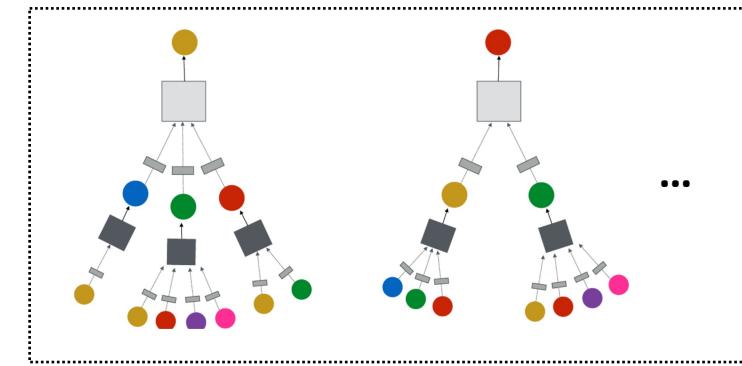
$D^{-1} \mathbf{A} \mathbf{H}^{(l)}$ (Matrix form)

$$\mathbf{H}^{(l+1)} = \sigma(\tilde{\mathbf{A}} \mathbf{H}^{(l)} \mathbf{W}_l^T + \mathbf{H}^{(l)} \mathbf{B}_l^T) \text{ where } \tilde{\mathbf{A}} = \mathbf{D}^{-1} \mathbf{A}$$

Neighborhood aggregation

Self transformation

Since $\tilde{\mathbf{A}}$ is sparse, sparse matrix multiplication can be used (efficient)



$$\mathbf{H}^{(k-1)}$$

$\mathbf{h}_i^{(k-1)}$

$$\sum_{u \in N(v)} \mathbf{h}_u^{(l)} = \mathbf{A}_v \mathbf{H}^{(l)}$$

$$\mathbf{D}^{-1}$$

$\mathbf{D}_{v,v}^{-1} = \frac{1}{|N(v)|}$

$$\mathbf{D}_{v,v} = \text{Deg}(v) = |N(v)|$$

$$\mathbf{D}_{v,v}^{-1} = \frac{1}{|N(v)|}$$

GCN: TRAINING

- We need to define the loss function on the embeddings
- We can feed the **final embeddings \mathbf{z}_v** into any loss function and run SGD to train the weight parameters
- **Types of loss function:** 1) Supervised loss, 2) Unsupervised loss
- **1) Supervised loss**

$$\min_{\theta} \sum_{v \in V} \mathcal{L}(y_v, f_{\theta}(\mathbf{z}_v))$$

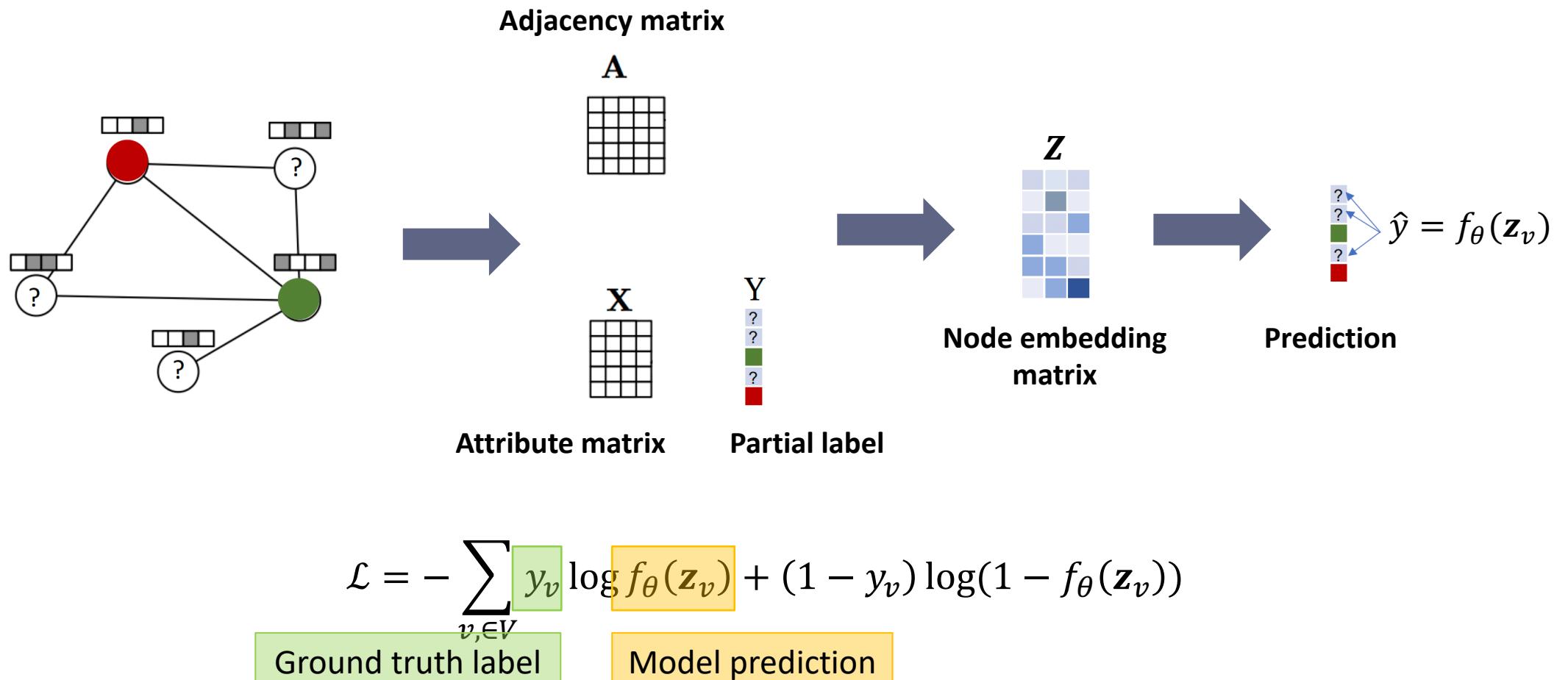
- y_v : Label of node v
 - f_{θ} : Classifier with parameter θ
 - \mathcal{L} could be squared error if y is real number (regression), or cross entropy if y is categorical (classification)
- **2) Unsupervised loss**

- No node label available
- We can use the graph structure as the supervision
 - e.g., adjacency information
 - In this case, \mathcal{L} is cross entropy ($A_{v,u} = 1$ if an edge exists between node v and node u , otherwise 0)

$$\min_{\theta} \sum_{v,u \in V} \mathcal{L}(A_{v,u}, f_{\theta}(\mathbf{z}_v, \mathbf{z}_u)) \quad f_{\theta}: \text{Encoder}$$

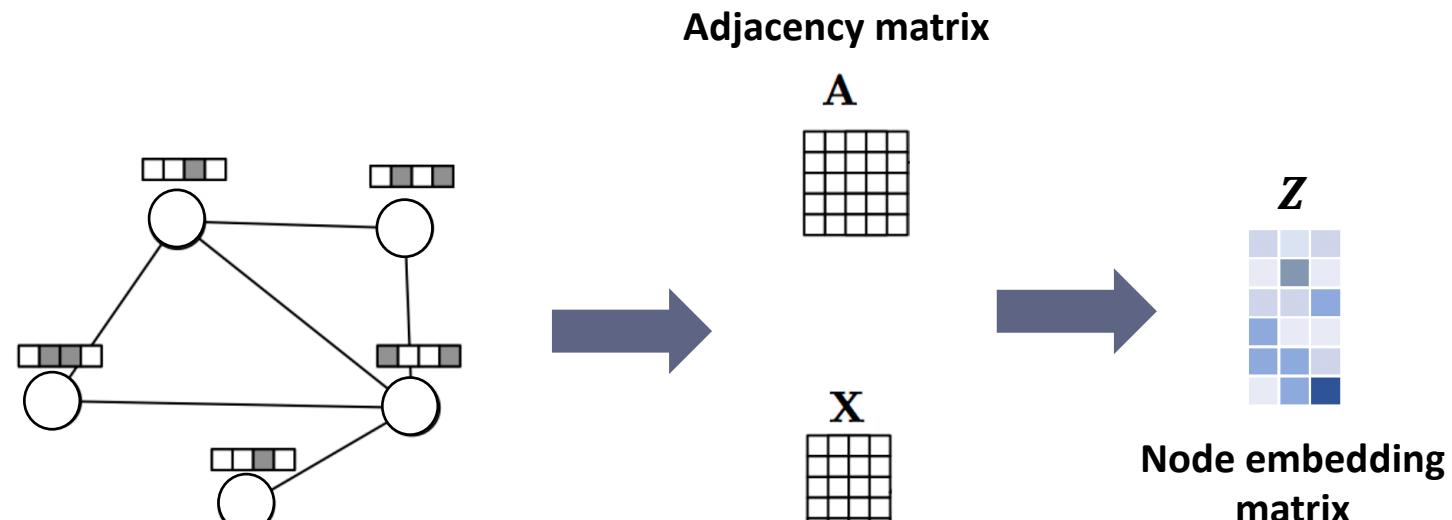
GCN: SUPERVISED TRAINING

- Directly train the model for a supervised task (e.g., node classification)



GCN: UNSUPERVISED TRAINING

- As we are not given node labels, we define our task to reconstruct the graph, i.e., Adjacency matrix



$$\mathcal{L} = - \sum_{v,u \in V} A_{v,u} \log f_\theta(\mathbf{z}_v, \mathbf{z}_u) + (1 - A_{v,u}) \log(1 - f_\theta(\mathbf{z}_v, \mathbf{z}_u))$$

Ground truth label

Model prediction

GRAPH ATTENTION NETWORKS (GAT)

- Idea: Treat different neighboring nodes differently

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu} \mathbf{h}_u^{(l)} \right)$$

Attention weight

$$\boxed{\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v)} \frac{\mathbf{h}_u^{(l)}}{|N(v)|} + \mathbf{B}_l \mathbf{h}_v^{(l)} \right)}$$

(GCN)

- α_{vu} : Importance of node u to node v as its neighboring node
- In GCN, the importance was heuristically defined based on the structural property of the graph (node degree)
 - $\alpha_{vu} = \frac{1}{|N(v)|}$: Does not depend on the neighbors (it is fixed)
 - All neighboring nodes $u \in N(v)$ are **equally important** to node v

Not all neighbors are equally important!

GRAPH ATTENTION NETWORKS (GAT)

- Idea: Treat different neighboring nodes differently

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu} \mathbf{h}_u^{(l)} \right)$$

Attention weight

- α_{vu} : Importance of node u to node v as its neighboring node

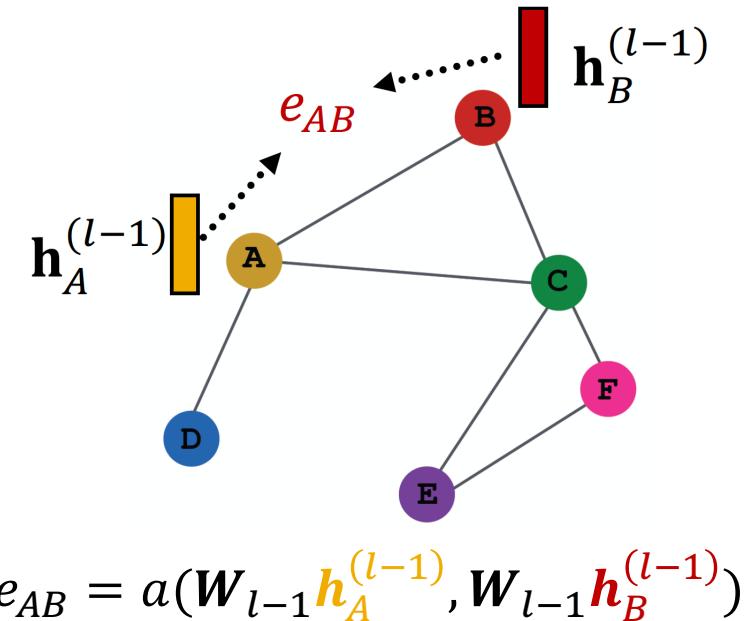
- Computing the attention weight

$$e_{vu} = a(\mathbf{W}_l \mathbf{h}_v^{(l)}, \mathbf{W}_l \mathbf{h}_u^{(l)})$$

(Importance of node u 's message to node v)

$$\alpha_{vu} = \frac{\exp(e_{uv})}{\sum_{k \in N(v)} \exp(e_{vk})}$$

(Normalization)



How do we define $a(\cdot)$?

GRAPH ATTENTION NETWORKS (GAT)

$$\alpha_{vu} = \frac{\exp(e_{uv})}{\sum_{k \in N(v)} \exp(e_{vk})}$$

- Defining the function $a(\cdot)$

$$\begin{aligned} e_{vu} &= a(\mathbf{W}_l \mathbf{h}_v^{(l)}, \mathbf{W}_l \mathbf{h}_u^{(l)}) \quad (\text{Importance of node } u's \text{ message to node } v) \\ &= \text{Linear}(\text{Concat}(\mathbf{W}_l \mathbf{h}_v^{(l)}, \mathbf{W}_l \mathbf{h}_u^{(l)})) \end{aligned}$$

Parameters of Linear layer is jointly trained
end-to-end with other parameters of GAT

- Multi-head attention (Refer to Lecture 5 - Transformer)

- Create multiple attention scores using multiple copies of parameters

$$\mathbf{h}_v^{(l)}[1] = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu}^{[1]} \mathbf{h}_u^{(l)} \right)$$

$$\mathbf{h}_v^{(l)}[2] = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu}^{[2]} \mathbf{h}_u^{(l)} \right)$$

$$\mathbf{h}_v^{(l)}[3] = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu}^{[3]} \mathbf{h}_u^{(l)} \right)$$



$$\mathbf{h}_v^{(l)} = \text{AGG}(\mathbf{h}_v^{(l)}[1], \mathbf{h}_v^{(l)}[2], \mathbf{h}_v^{(l)}[3])$$

The final embedding aggregates the outputs
of multi-head attention

GRAPHSAGE

▪ Generalization of GCN/GAT

- So far we have aggregated the neighbor messages by taking their (weighted) average Can we do better?

Average of neighboring nodes

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \left[\sum_{u \in N(v)} \frac{\mathbf{h}_u^{(l)}}{|N(v)|} \right] + \mathbf{B}_l \mathbf{h}_v^{(l)} \right)$$

GCN

Add self representation

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu} \mathbf{h}_u^{(l)} \right)$$

GAT

Attention weight

GraphSAGE

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\left[\mathbf{W}_l \cdot \text{AGG} \left(\left\{ \mathbf{h}_u^{(l)} | u \in N(v) \right\} \right), \mathbf{B}_l \mathbf{h}_v^{(l)} \right] \right)$$

Generalized representation

Generalized representation

GRAPHSAGE

▪ Generalization of GCN/GAT

- So far we have aggregated the neighbor messages by taking their (weighted) average Can we do better?

$$\mathbf{h}_v^{(l+1)} = \sigma \left(\left[\mathbf{W}_l \cdot \text{AGG} \left(\left\{ \mathbf{h}_u^{(l)} \mid u \in N(v) \right\} \right), \mathbf{B}_l \mathbf{h}_v^{(l)} \right] \right)$$

Generalized representation (Concatenation here)

Generalized representation

▪ Variants of AGG

- **Mean:** Same as GCN

$$\text{AGG} = \sum_{u \in N(v)} \frac{\mathbf{h}_u^{(l)}}{|N(v)|}$$

- **Pool:** Transform neighbor vectors and apply symmetric vector function

$$\text{AGG} = \gamma \left(\left\{ \text{MLP}(\mathbf{h}_u^{(l)}) \mid u \in N(v) \right\} \right), \text{ where } \gamma \text{ is element-wise mean/max/min}$$

- **LSTM:** Apply LSTM to shuffled neighbors

$$\text{AGG} = \text{LSTM} \left(\left[\mathbf{h}_u^{(l)} \mid u \in \pi(N(v)) \right] \right)$$

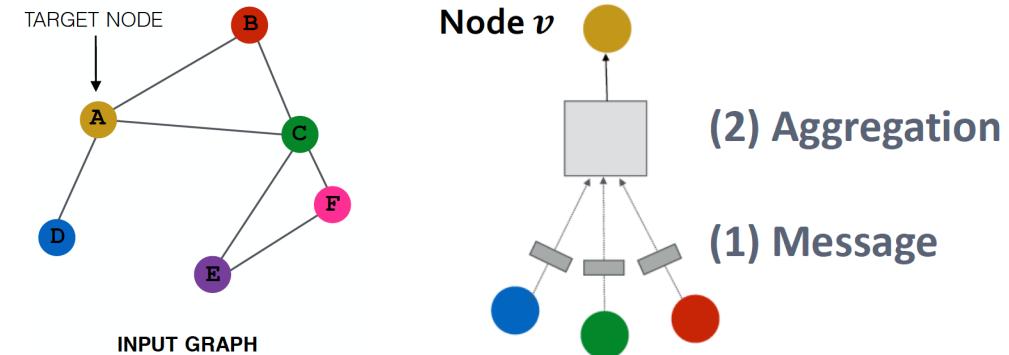
A GNN LAYER: OVERVIEW

- GNN layer = 1) Message + 2) Aggregation
- Compress a set of vectors into a single vector
- Different types of GNN layers
 - GCN, GraphSAGE, GAT, ...

$$(GCN) \quad \mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v)} \frac{\mathbf{h}_u^{(l)}}{|N(v)|} + \mathbf{B}_l \mathbf{h}_v^{(l)} \right)$$

$$(GAT) \quad \mathbf{h}_v^{(l+1)} = \sigma \left(\mathbf{W}_l \sum_{u \in N(v) \cup v} \alpha_{vu} \mathbf{h}_u^{(l)} \right)$$

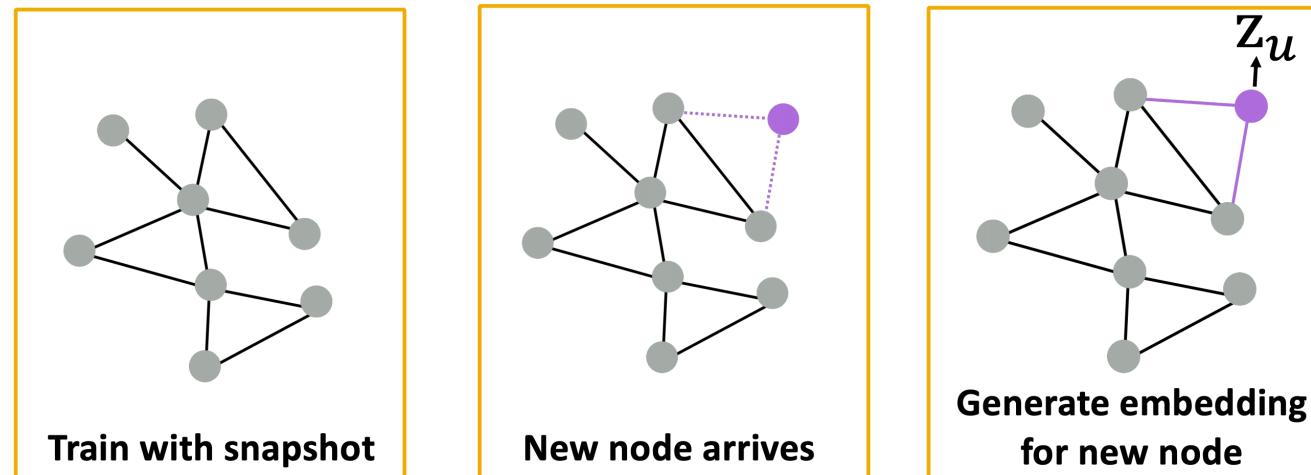
$$(\text{GraphSAGE}) \quad \mathbf{h}_v^{(l+1)} = \sigma \left(\left[\mathbf{W}_l \cdot \text{AGG} \left(\left\{ \mathbf{h}_u^{(l)} | u \in N(v) \right\} \right), \mathbf{B}_l \mathbf{h}_v^{(l)} \right] \right)$$



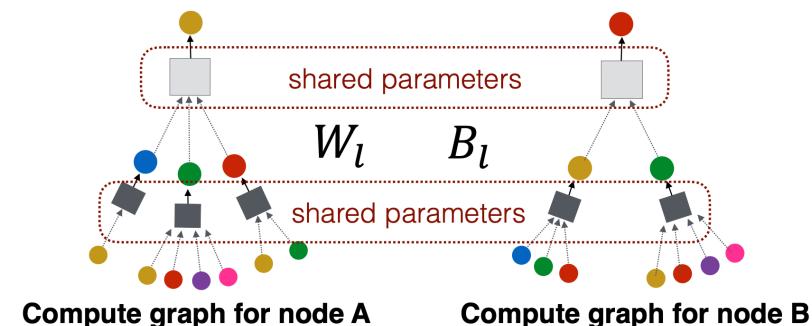
- (1) Message computation
 - Message function: $\mathbf{m}_u^{(l)} = \text{MSG}^{(l)}(\mathbf{h}_u^{(l)})$
 - Example: A linear layer $\mathbf{m}_u^{(l)} = \mathbf{W}_l \mathbf{h}_u^{(l)}$
- (2) Aggregation
 - $\mathbf{h}_v^{(l)} = \text{AGG} \left(\left\{ \mathbf{m}_u^{(l)} | u \in N(v) \right\} \right)$
 - Example: SUM(), MEAN(), or MAX() aggregator
 - $\mathbf{h}_v^{(l)} = \text{SUM} \left(\left\{ \mathbf{m}_u^{(l)} | u \in N(v) \right\} \right) = \sum_{u \in N(v)} \mathbf{W}_l \mathbf{h}_u^{(l)}$
 - $= \sum_{u \in N(v)} \mathbf{m}_u^{(l)}$

INDUCTIVE CAPABILITY OF GNN

- **Inductive learning:** We can obtain embeddings for nodes that have not appeared in the training time
 - e.g., In Amazon, new users are consistently added to the system, and it is impractical to re-train the system to get the embeddings for the new users



- This is possible because we do not train an embedding matrix as done in Deepwalk/node2vec
- Instead, we train **aggregator** and **transformer**



OUTLINE

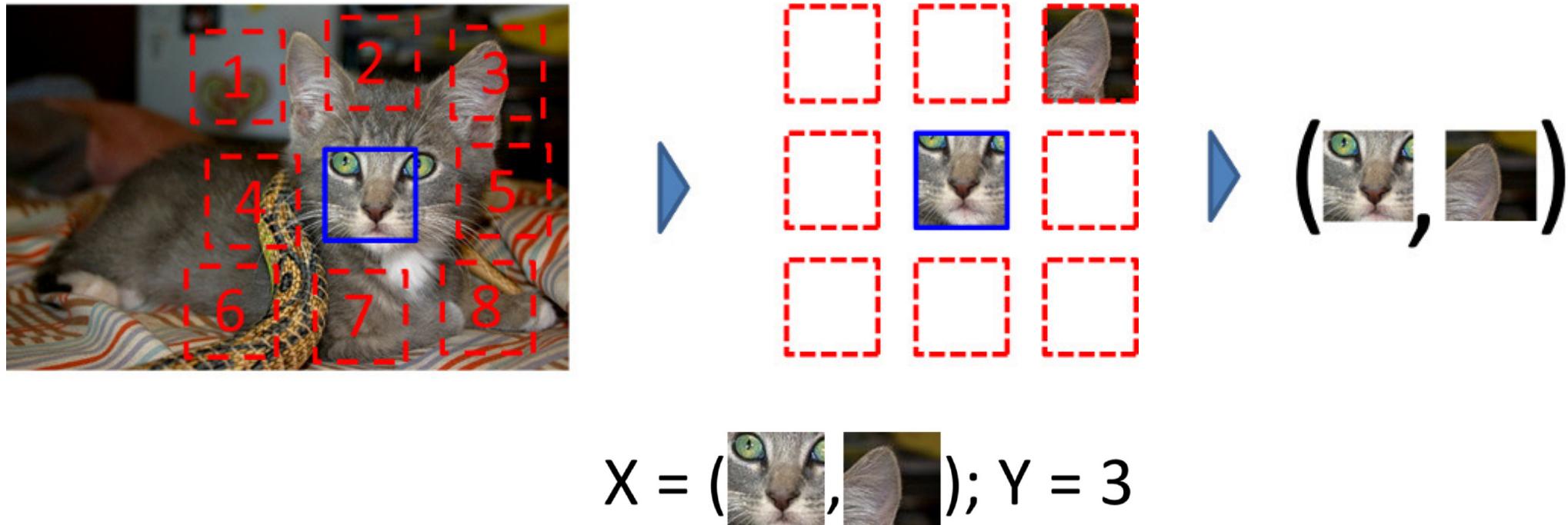
- Overview
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OUTLINE

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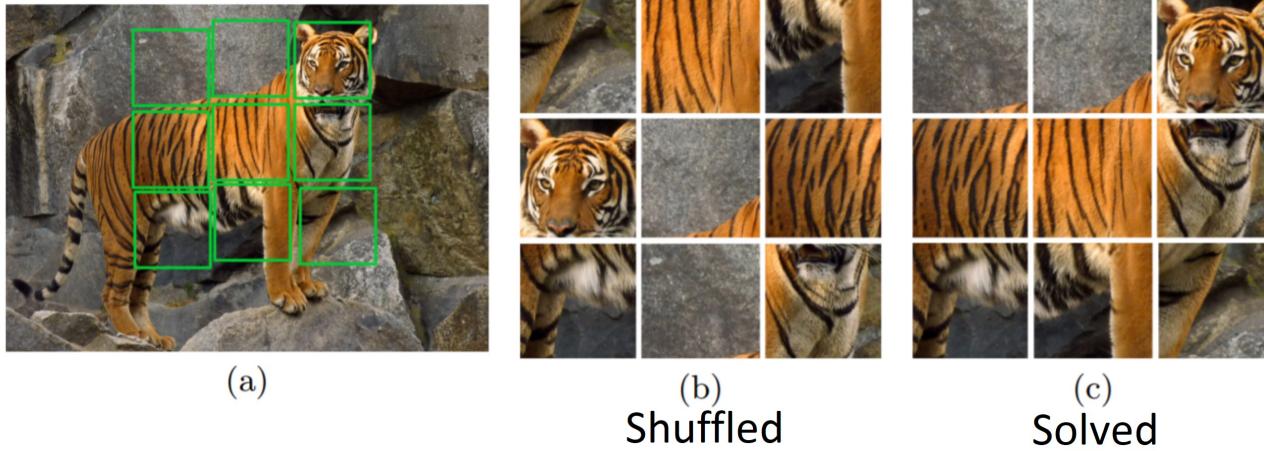
What is self-supervised learning?

- A form of unsupervised learning where the data provides the supervision
- In general, withhold some part of the data, and task the network with predicting it
- An example of **pretext task: Relative positioning**
 - Train network to predict relative position of two regions in the same image

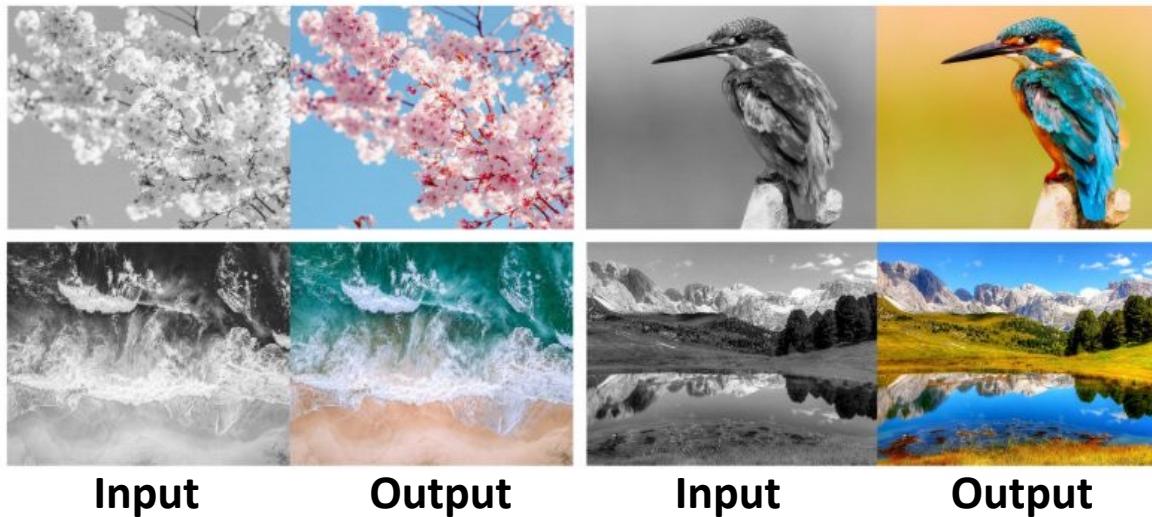


What is self-supervised learning?

- Pretext task: Jigsaw puzzle

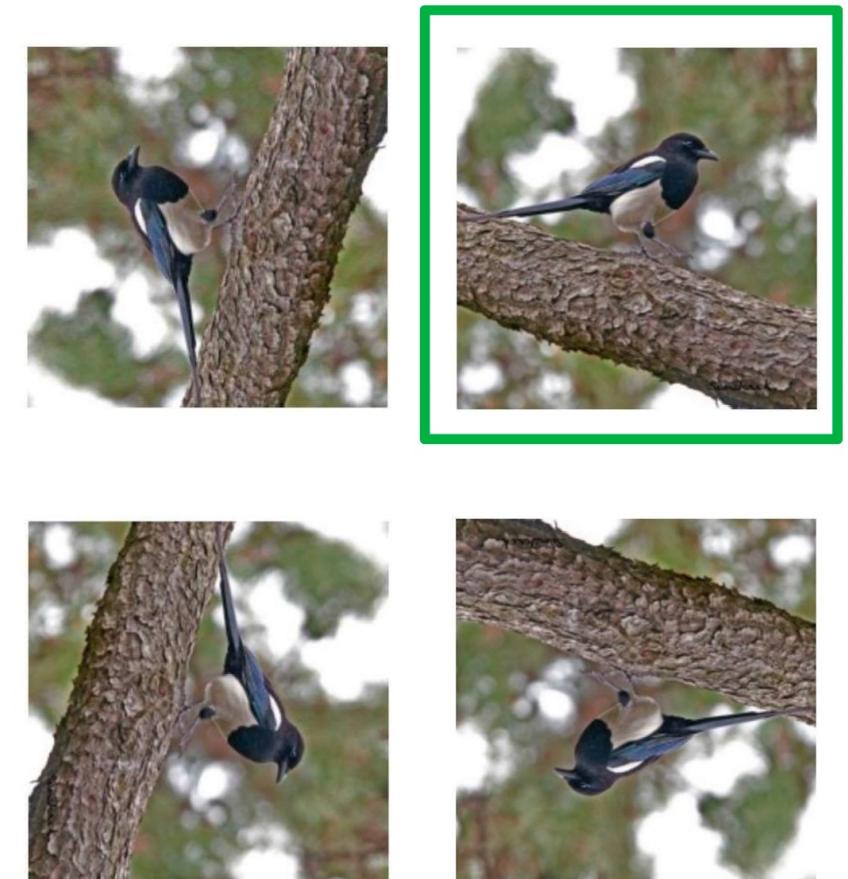


- Pretext task : Colorization



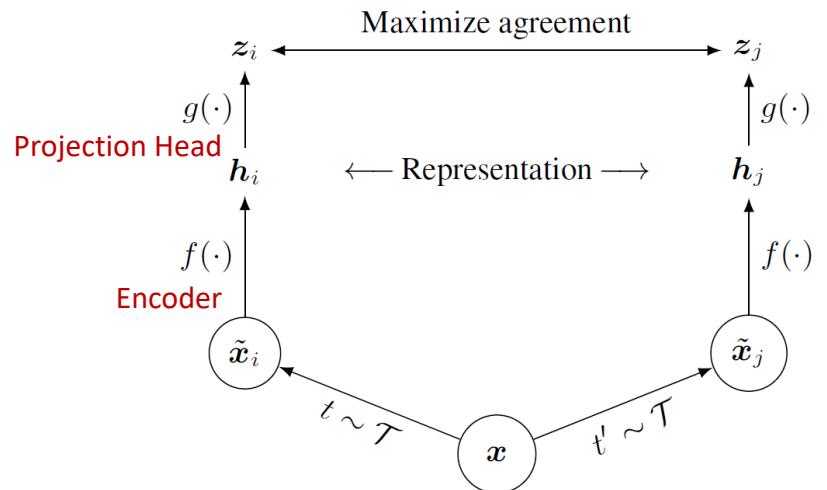
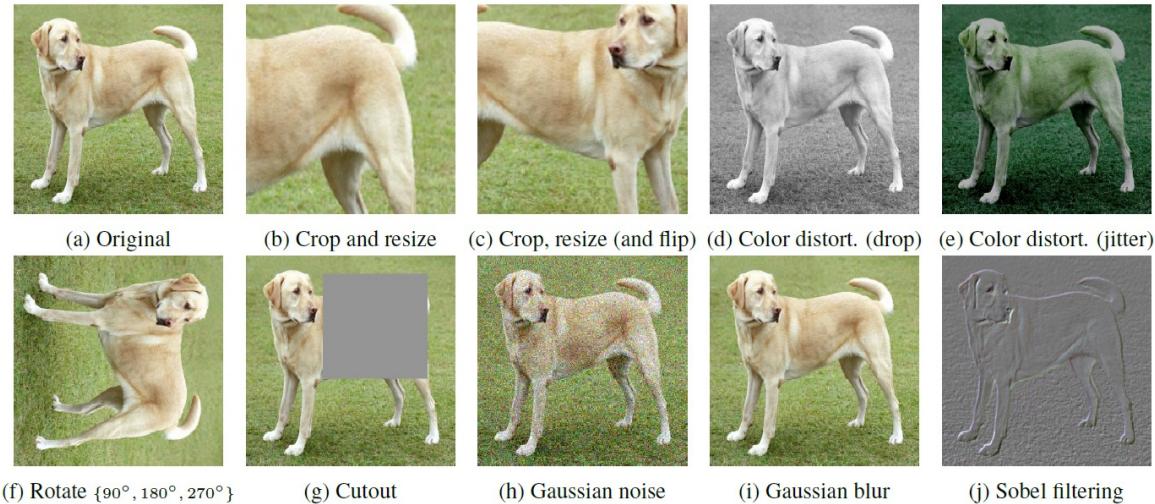
- Pretext task : Rotation

- Which one has the correct rotation?



(Figure credit) Self-Supervised Learning, Andrew Zisserman

The Contrastive Learning Paradigm



Algorithm

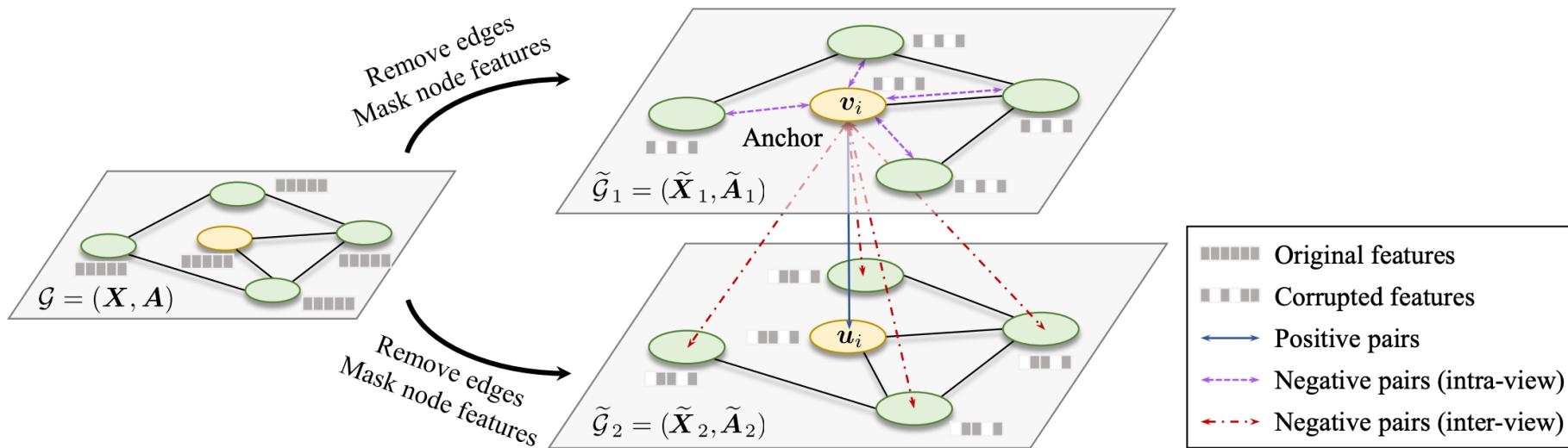
- 1) Sample mini batch of N examples
- 2) Create $2N$ data points via Data Augmentation
- 3) Given a positive pair, treat other $2(N - 1)$ points as negative examples
- **Instance Discrimination!**

$$\ell_{i,j} = -\log \frac{\exp(\text{sim}(z_i, z_j)/\tau)}{\sum_{k=1}^{2N} \mathbb{1}_{[k \neq i]} \exp(\text{sim}(z_i, z_k)/\tau)}$$

Reduce: Dist. between representations of different augmented views of the same image (Positive)
Increase: Dist. between representations of augmented views from different images (Negative)

Deep Graph Contrastive Representation Learning (GRACE)

- Pull the representation of the same node in the two augmented graphs
- Push apart representations of every other node



$$\ell(\mathbf{u}_i, \mathbf{v}_i) = \log \frac{e^{\theta(\mathbf{u}_i, \mathbf{v}_i)/\tau}}{\underbrace{e^{\theta(\mathbf{u}_i, \mathbf{v}_i)/\tau}}_{\text{the positive pair}} + \underbrace{\sum_{k=1}^N \mathbf{1}_{[k \neq i]} e^{\theta(\mathbf{u}_i, \mathbf{v}_k)/\tau}}_{\text{inter-view negative pairs}} + \underbrace{\sum_{k=1}^N \mathbf{1}_{[k \neq i]} e^{\theta(\mathbf{u}_i, \mathbf{u}_k)/\tau}}_{\text{intra-view negative pairs}}},$$

Shortcomings of Contrastive Methods

- 1) Requires negative samples → **Sampling bias**
 - Treat different image as negative even if they share the semantics
- 2) Requires careful augmentation

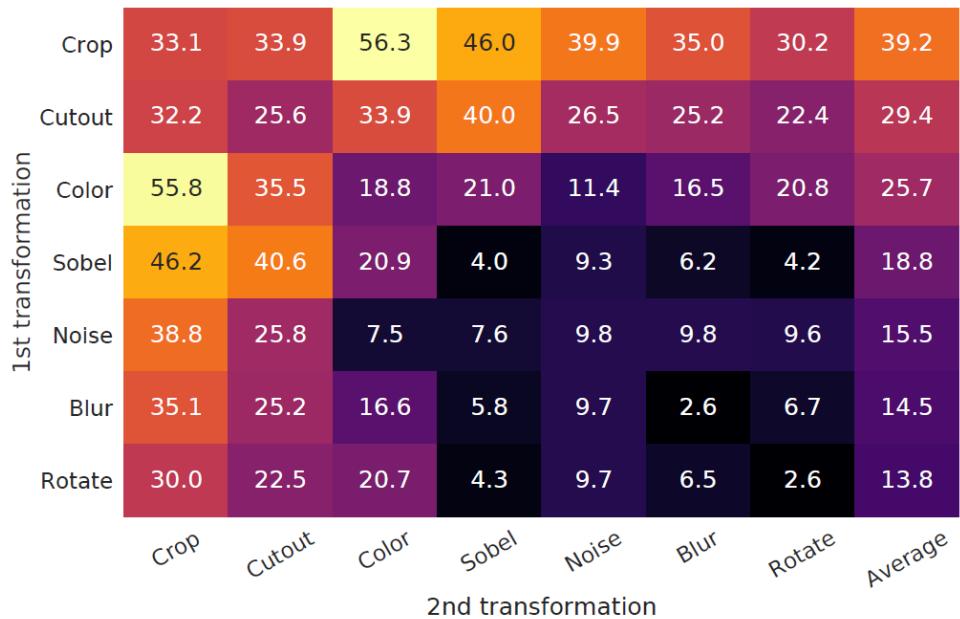
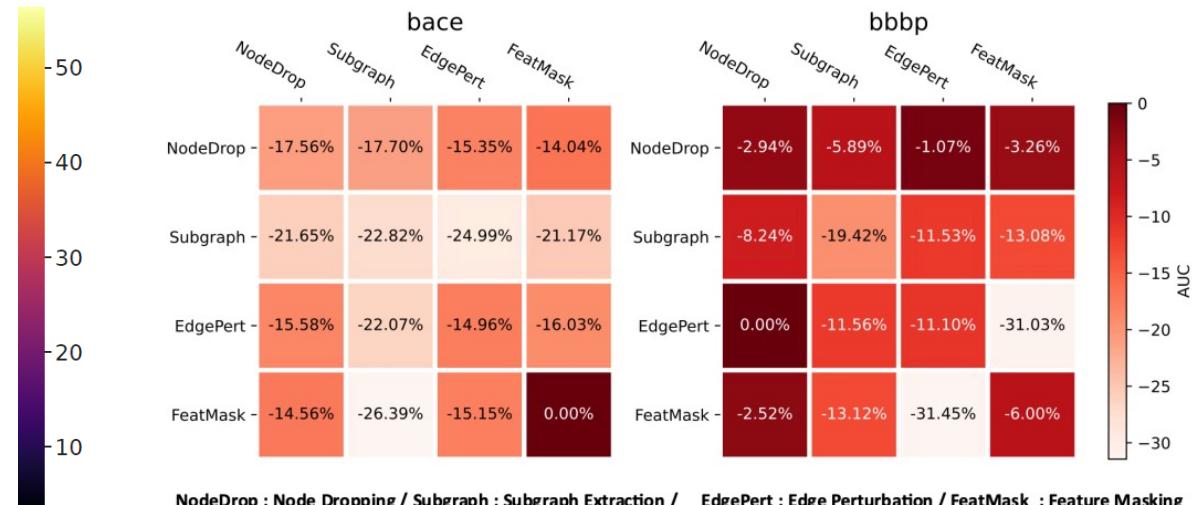


Image classification



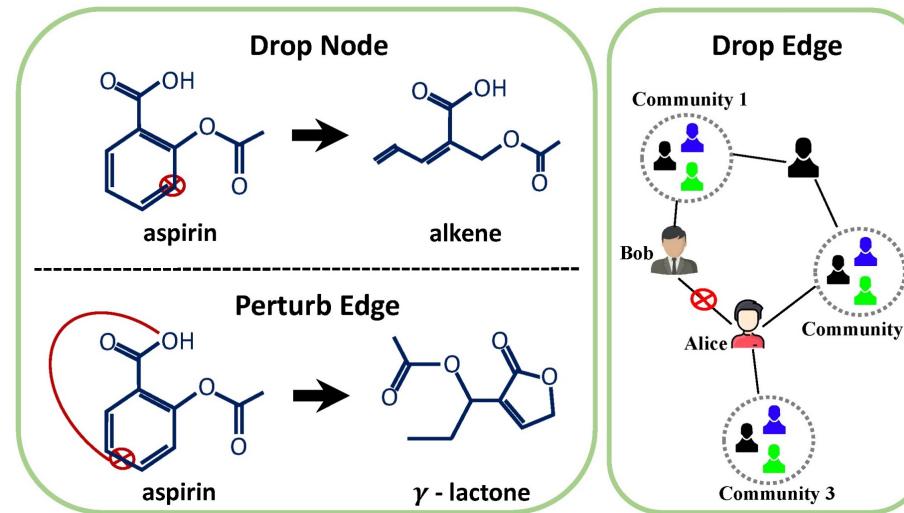
Graph classification

Motivation: Is Augmentation Appropriate for Graph-structured Data?

- Image's underlying semantic is hardly changed after augmentation



- However in the case of graphs, we cannot ascertain whether the augmented graph would be positively related to original graph



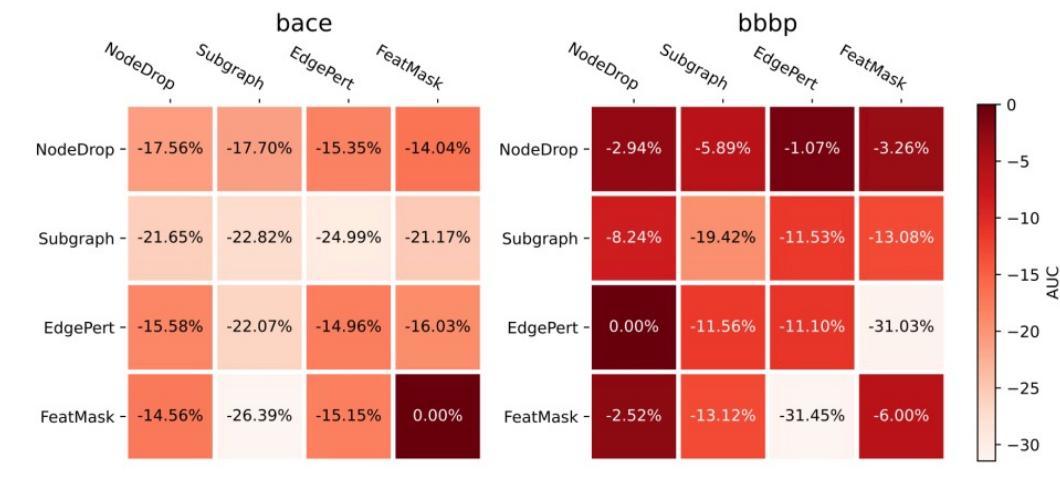
Because graphs contain not only the semantic but also the **structural information**

Motivation: Is Augmentation Appropriate for Graph-structured Data?

- Performance sensitivity according to hyperparameters for augmentations

		Comp.	Photo	CS	Physics
Node	BGRL	-4.00%	-1.06%	-0.20%	-0.69%
Classi.	GCA	-19.18%	-5.48%	-0.27%	OOM
Node	BGRL	-11.57%	-13.30%	-0.78%	-6.46%
Clust.	GCA	-26.28%	-23.27%	-1.64%	OOM

Node-level task



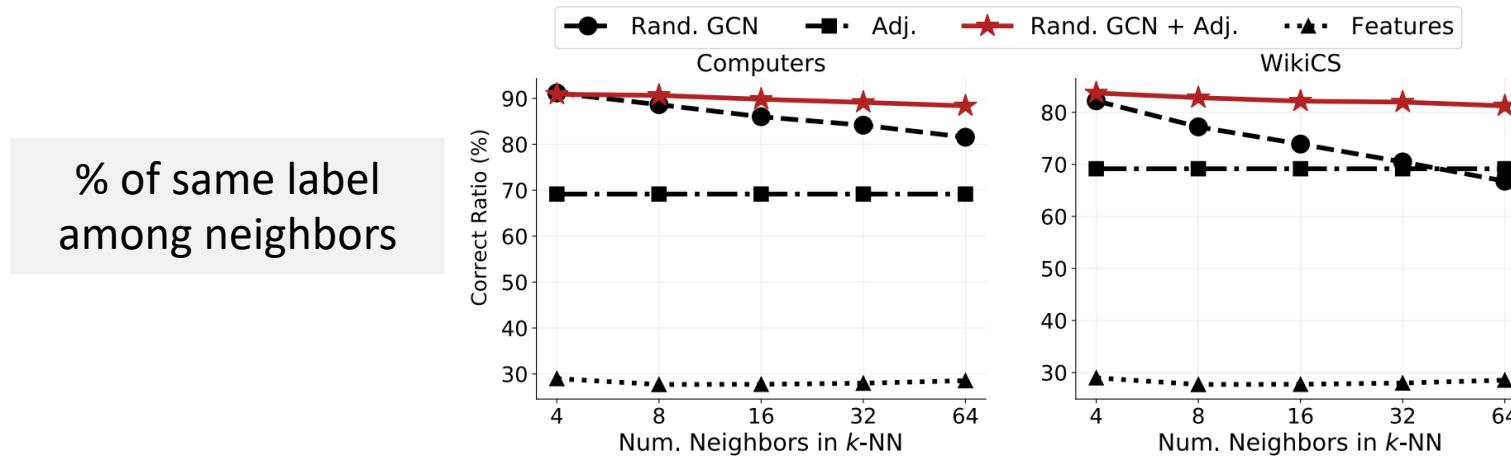
Graph-level task

- The quality of the learned representations relies on the **choice of augmentation scheme**
 - Performance on various downstream tasks varies greatly according to the choice of augmentation hyperparameters

We need more stable and general framework for generating alternative view of the original graph
without relying on augmentation

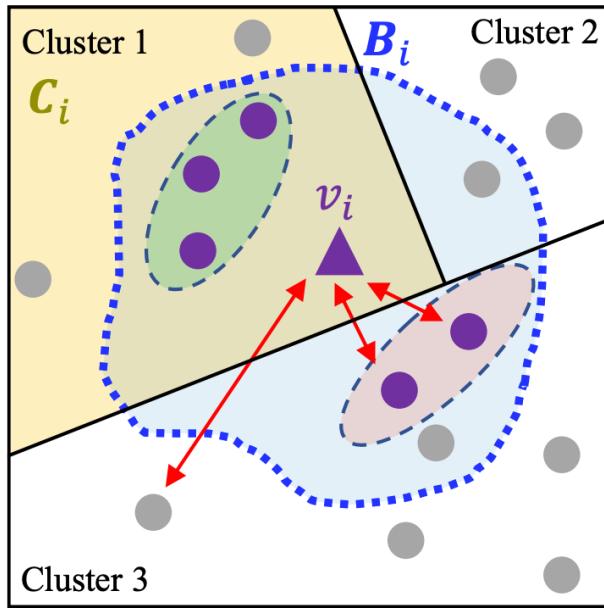
Augmentation-Free Graph Representation Learning

- Instead of creating two arbitrarily augmented views of a graph,
 - Use the original graph as-is as one view, and generate another view by discovering nodes that can serve as positive samples via **k-nearest neighbor search in embedding space**
- However, naively selected positive samples with k-NN includes false positives
 - More than 10% of false negatives



We need to filter out false positives regarding local and global perspective!

Capturing Local and Global Semantics



- B_i : Set of k-NNs of query v_i
- N_i : Set of adjacent nodes of query v_i
- C_i : Set of nodes that are in the same cluster with query v_i

- ▲ Query Node (v_i)
- Node ($\mathcal{V} \setminus v_i$)
- Nearest Neighbors (B_i)
- ↔ Adjacency (N_i)
- Same cluster as v_i (C_i)
- Local Positive ($B_i \cap N_i$)
- Global Positive ($B_i \cap C_i$)
- Real Positive (P_i)

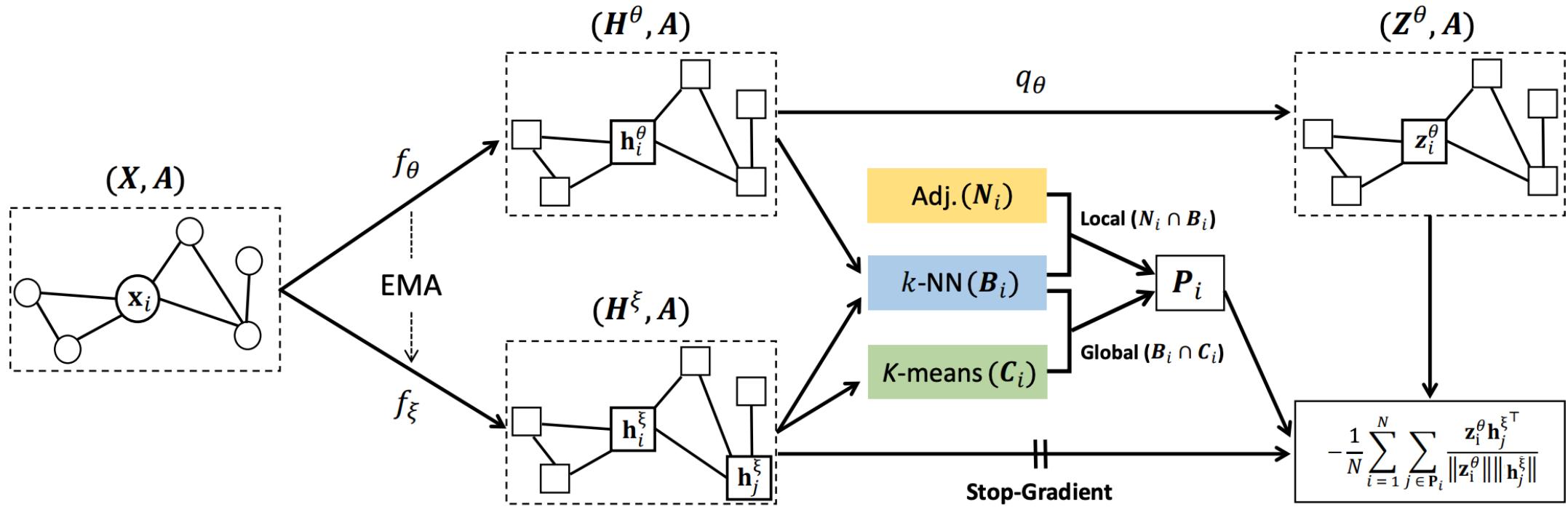
- Obtain real positives for v_i

$$\mathbf{P}_i = (\mathbf{B}_i \cap \mathbf{N}_i) \cup (\mathbf{B}_i \cap \mathbf{C}_i)$$

- Minimize the cosine distance between query and real positives \mathbf{P}_i

$$\mathcal{L}_{\theta, \xi} = -\frac{1}{N} \sum_{i=1}^N \sum_{v_j \in \mathbf{P}_i} \frac{\mathbf{z}_i^\theta \mathbf{h}_j^{\xi \top}}{\|\mathbf{z}_i^\theta\| \|\mathbf{h}_j^\xi\|}$$

Overall Architecture of AFGRL



Experiments

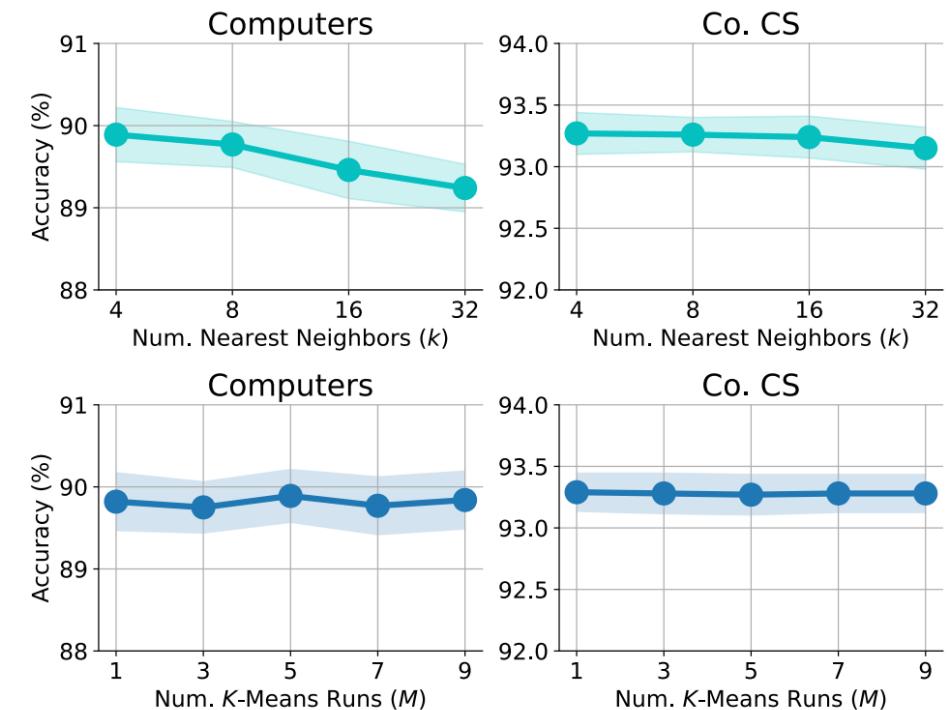
- Task: Node classification

	WikiCS	Computers	Photo	Co.CS	Co.Physics
Sup. GCN	77.19 ± 0.12	86.51 ± 0.54	92.42 ± 0.22	93.03 ± 0.31	95.65 ± 0.16
Raw feats.	71.98 ± 0.00	73.81 ± 0.00	78.53 ± 0.00	90.37 ± 0.00	93.58 ± 0.00
node2vec	71.79 ± 0.05	84.39 ± 0.08	89.67 ± 0.12	85.08 ± 0.03	91.19 ± 0.04
DeepWalk	74.35 ± 0.06	85.68 ± 0.06	89.44 ± 0.11	84.61 ± 0.22	91.77 ± 0.15
DW + feats.	77.21 ± 0.03	86.28 ± 0.07	90.05 ± 0.08	87.70 ± 0.04	94.90 ± 0.09
DGI	75.35 ± 0.14	83.95 ± 0.47	91.61 ± 0.22	92.15 ± 0.63	94.51 ± 0.52
GMI	74.85 ± 0.08	82.21 ± 0.31	90.68 ± 0.17	OOM	OOM
MVGRL	77.52 ± 0.08	87.52 ± 0.11	91.74 ± 0.07	92.11 ± 0.12	95.33 ± 0.03
GRACE	77.97 ± 0.63	86.50 ± 0.33	92.46 ± 0.18	92.17 ± 0.04	OOM
GCA	77.94 ± 0.67	87.32 ± 0.50	92.39 ± 0.33	92.84 ± 0.15	OOM
BGRI	76.86 ± 0.74	89.69 ± 0.37	93.07 ± 0.38	92.59 ± 0.14	95.48 ± 0.08
AFGRL	77.62 ± 0.49	89.88 ± 0.33	93.22 ± 0.28	93.27 ± 0.17	95.69 ± 0.10

Recall...

		Comp.	Photo	CS	Physics
Node	BGRL	-4.00%	-1.06%	-0.20%	-0.69%
Classi.	GCA	-19.18%	-5.48%	-0.27%	OOM
Node	BGRL	-11.57%	-13.30%	-0.78%	-6.46%
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AFGRL outperforms SOTA baselines



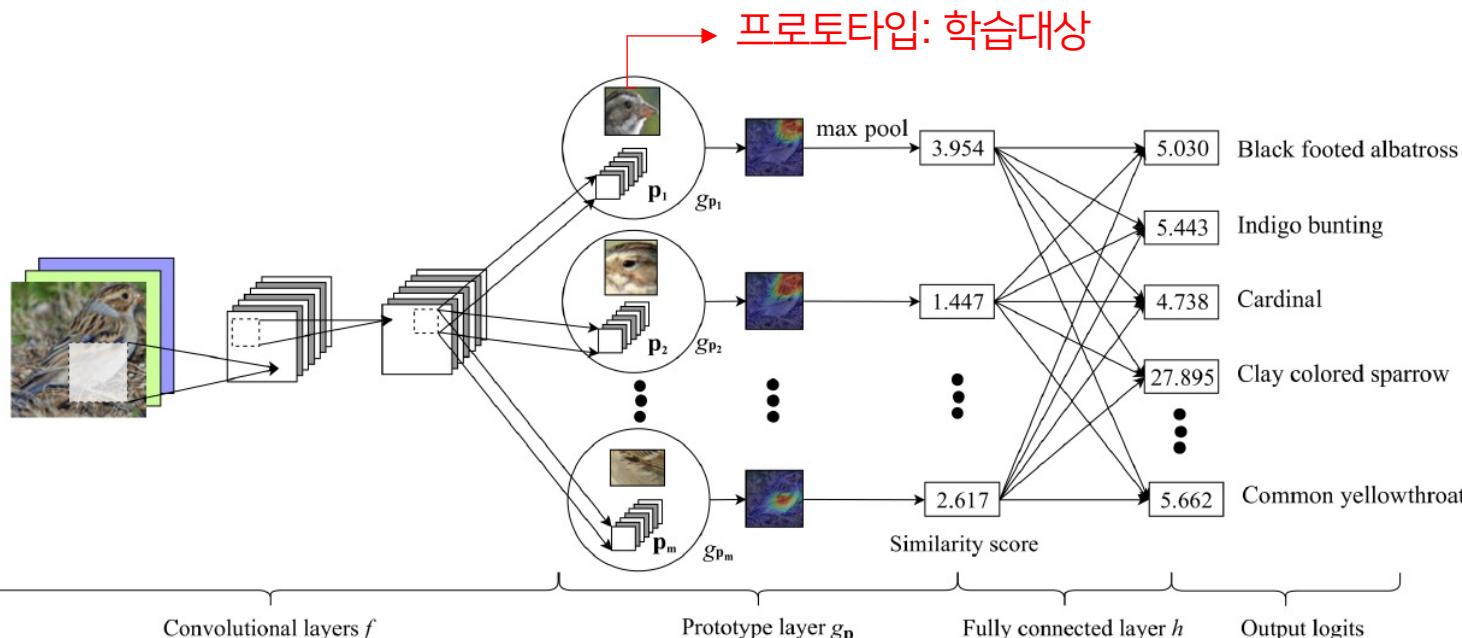
AFGRL is stable over hyperparameters
→ Can be easily trained compared with other augmentation-based methods.

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INTRODUCTION

- Explainable AI
 - Reasoning process : 모델이 해당 결정을 내리기 위해 수행한 과정을 제시
 - 모델이 올바른 예측 또는 잘못된 예측을 한 이유에 대한 과정을 시각화하고 분석



- 학습된 Prototype을 활용하여 reasoning process 제시
 - 각 class에 대해 고정된 수의 prototype을 할당

1. 입력 이미지로부터 feature 추출
2. 각 class에 대한 prototype과 유사성 점수를 계산
3. 계산한 유사성 점수는 fully connected layer를 통해 output logit값을 생성

INTRODUCTION

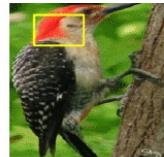
- Reasoning process : 모델이 해당 결정을 내리기 위해 수행한 과정을 제시

새의 종을 분류



붉은배 딱따구리

프로토타입



유사성
점수

$$6.499 \times 1.180 = 7.669$$



$$4.392 \times 1.127 = 4.950$$



$$3.890 \times 1.108 = 4.310$$

⋮ ⋮ ⋮ ⋮ ⋮ ⋮

붉은배 딱따구리에 대한 총 점수

: 32.736

붉은 벼슬 딱따구리

프로토타입

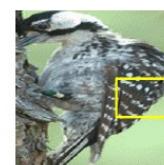


유사성
점수

$$2.452 \times 1.046 = 2.565$$



$$2.125 \times 1.091 = 2.318$$



$$1.945 \times 1.069 = 2.079$$

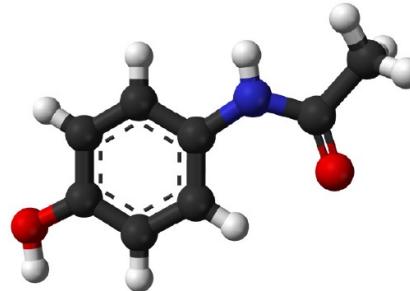
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붉은 벼슬 딱따구리에 대한 총 점수

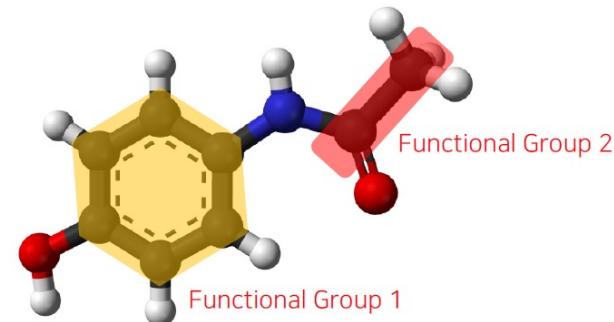
: 16.886

INTRODUCTION

- Graph data
 - Node와 Edge로 구성, 개체 간의 관계를 표현하기 위해 사용
 - Graph Classification : Graph가 어떤 레이블에 속하는지 예측
 - Application : 분자 특성 예측, 약물 발견, 장면 이해
- 분자 구조의 Graph data
 - 분자는 원자 간의 결합으로 구성되어 있어 Graph로 표현하기 적합
 - Functional Group : 분자 전체의 특성을 결정 짓는 서브그래프

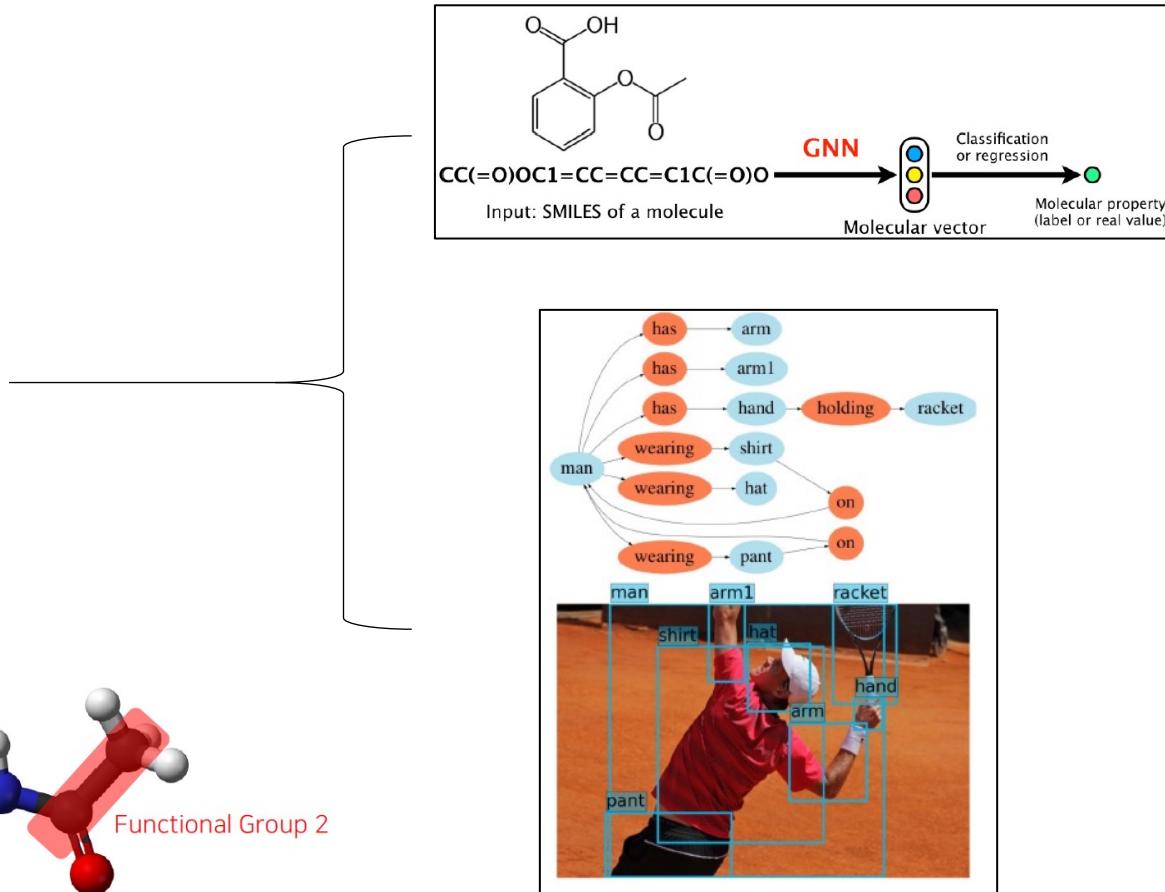


분자 = Graph



Functional Group = Subgraph

→ 중요한 subgraph를 탐지하기 위해 정보 이론 기반 방식이 제안됨



INTRODUCTION

- Graph Information Bottleneck

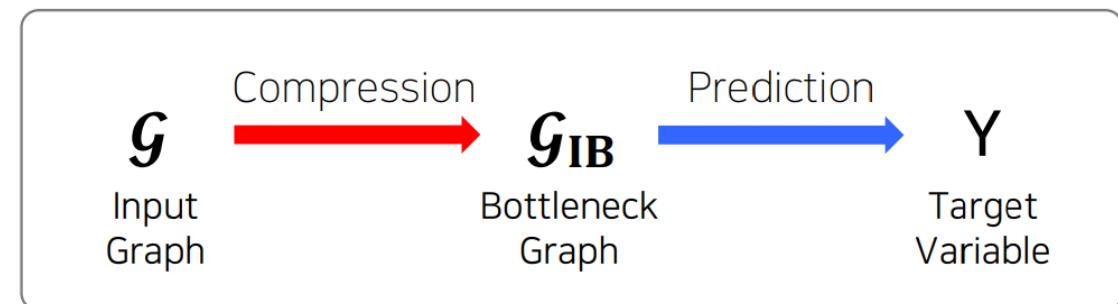
- Original graph \mathcal{G} 를 bottleneck graph \mathcal{G}_{IB} 로 압축하는 동시에 prediction Y 와 관련된 정보를 유지하는 subgraph를 탐지



- 목표 : Label Y 를 예측하는 데 중요한 Bottleneck graph \mathcal{G}_{IB} 을 찾는다.
- Mutual Information(I) : 특정 variable이 다른 variable에 영향을 미치는 정도

$$\min_{\mathcal{G}_{IB}} -I(Y, \mathcal{G}_{IB}) + \beta I(\mathcal{G}; \mathcal{G}_{IB})$$

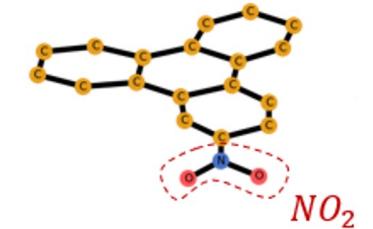
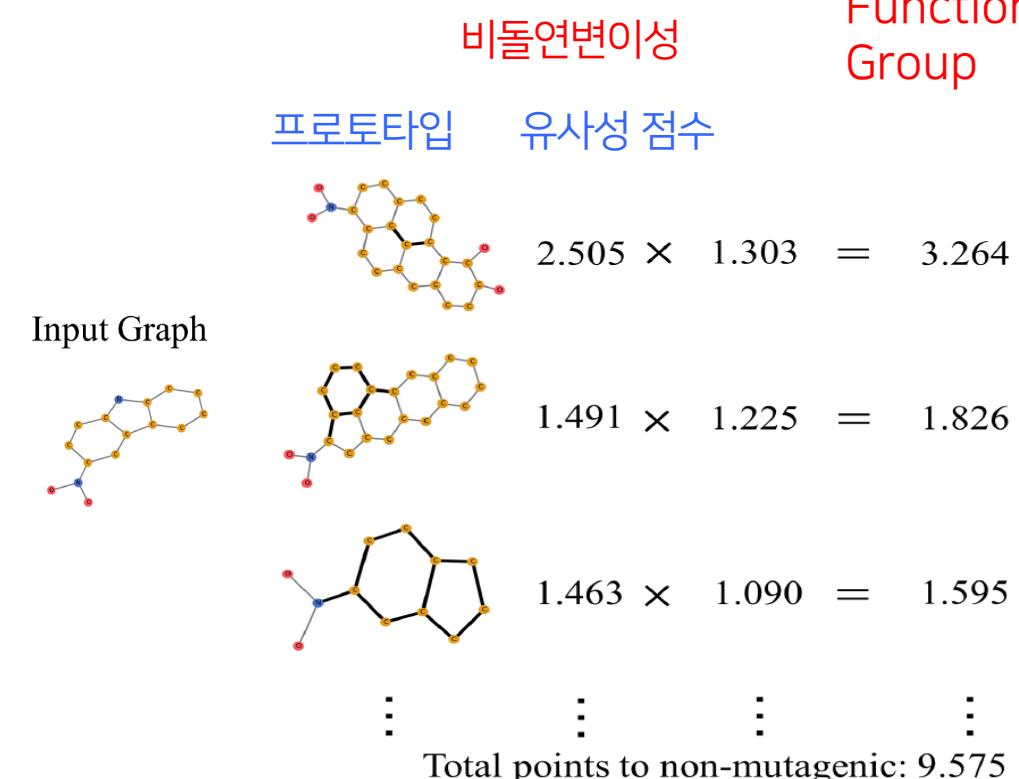
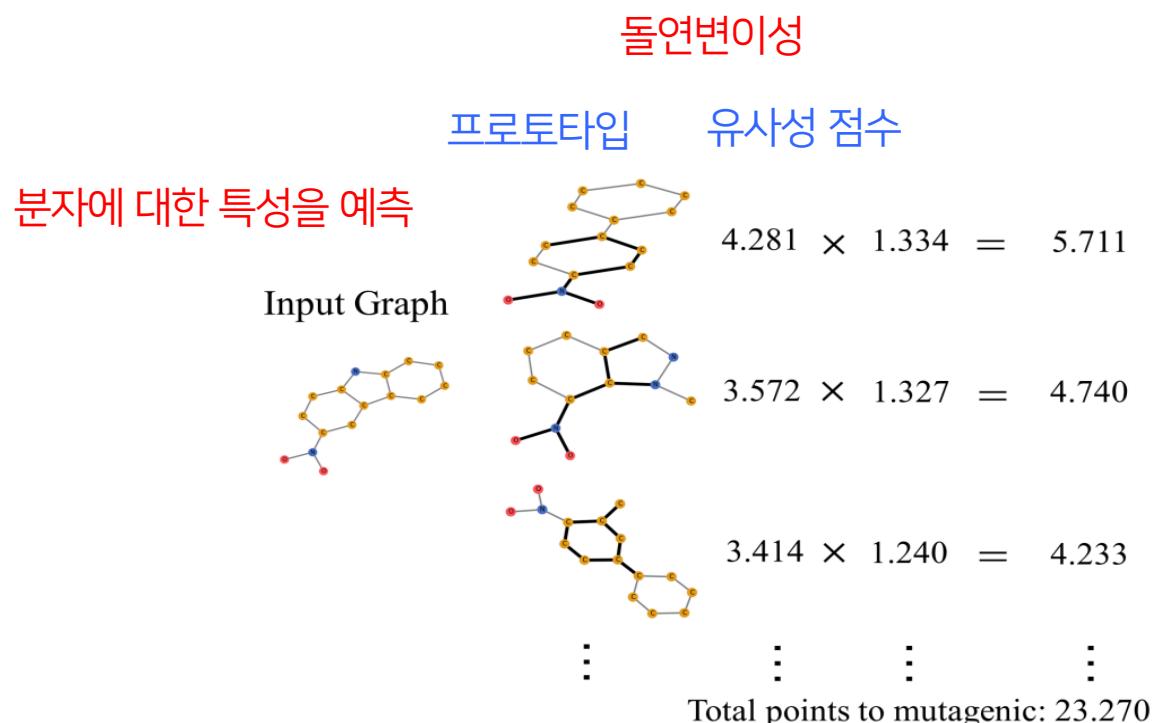
Prediction Compression



INTRODUCTION

- ProtGNN

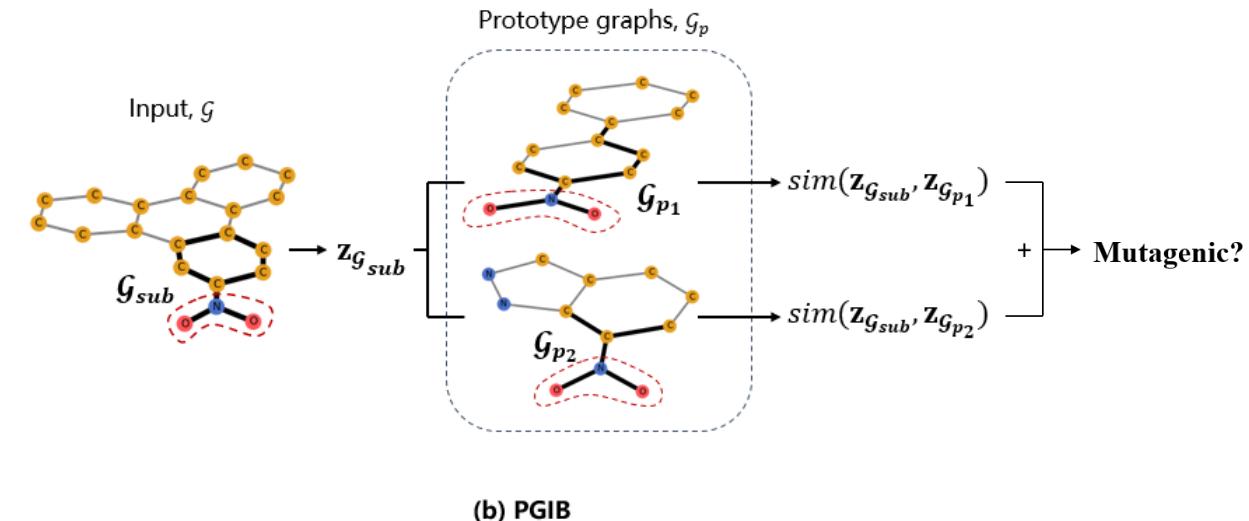
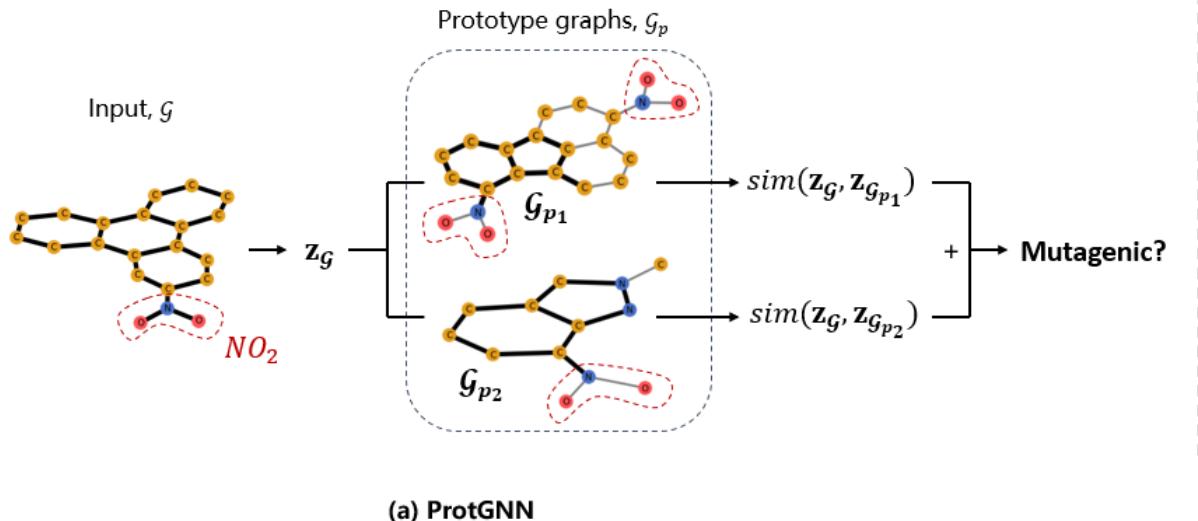
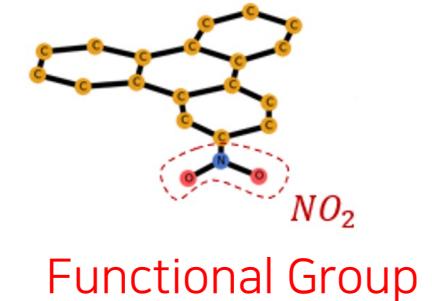
- 그래프 데이터에서 Prototype을 활용하여 reasoning process를 제시한 모델
- 모델이 특정 그래프를 해당 레이블로 예측하게 된 과정을 시각화



INTRODUCTION

- ProtGNN

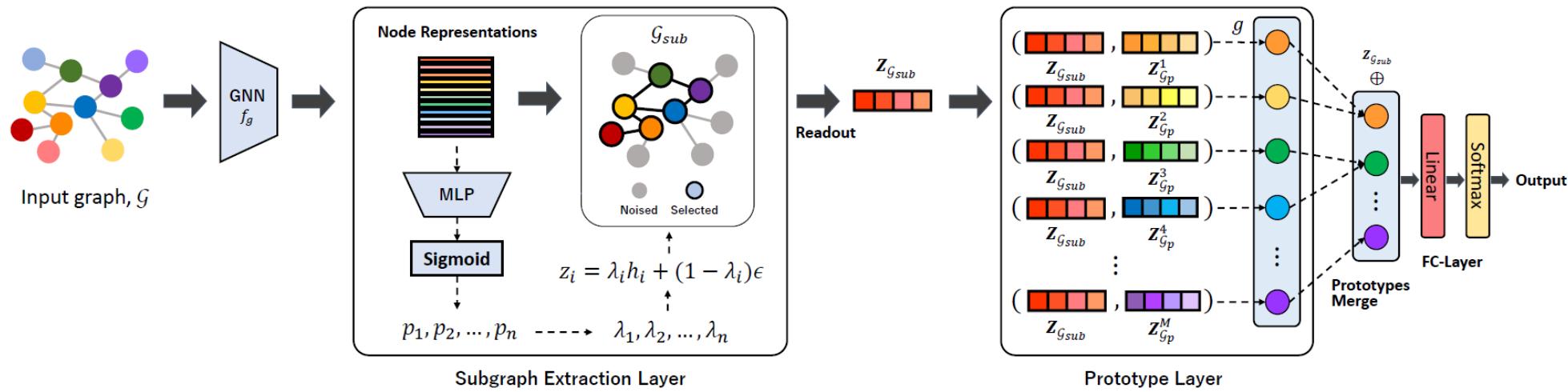
- 학습 과정에서 prototype이 중요한 subgraph를 포착하지 못하거나 불필요한 정보를 포함
- **NO₂**(Functional group)를 제외한 다른 공통의 subgraph에 대해서 높은 similarity score를 계산
→ Prototype이 예측에 중요한 결정 근거를 충분히 포함하도록 해야함.
- Information Bottleneck를 Prototype 관점에서 접근하여 prototype이 예측에 중요한 subgraph에 관한 정보를 전달받음



PROPOSED METHOD

Interpretable Prototype-based Graph Information Bottleneck

- Architecture



PGIB

- Identify core subgraphs.
- Calculate similarity scores with prototypes.
- Merge prototypes.
- Predict the label from the fully connected layer.

$$\min_{\mathcal{G}_{sub}} -I(Y; \mathcal{G}_{sub}) + \beta I(\mathcal{G}; \mathcal{G}_{sub})$$



$$\min_{\mathcal{G}_{sub}} -I(Y; \mathcal{G}_{sub}, \mathcal{G}_p) + I(Y; \mathcal{G}_p | \mathcal{G}_{sub}) + \beta I(\mathcal{G}; \mathcal{G}_{sub})$$

Prototype Layer

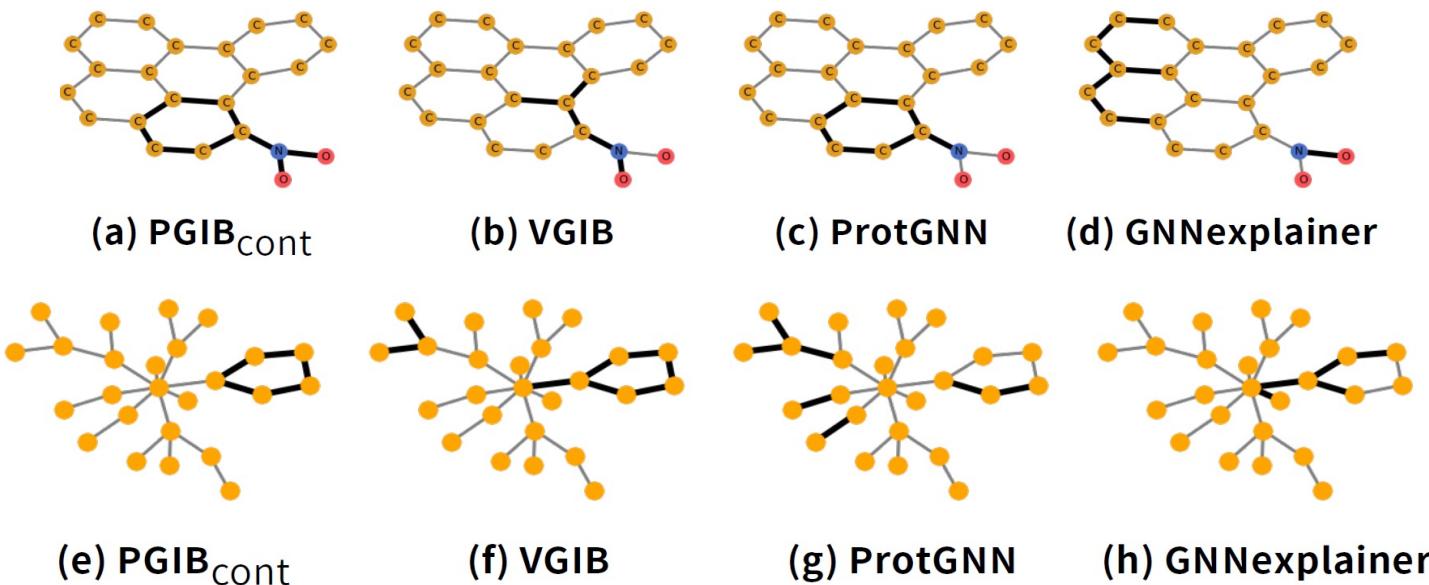
Subgraph Extraction Layer

EXPERIMENT

Graph Classification

Dataset	Methods								
	GCN	GIN	GAT	ProtGNN	GIB	VGIB	GSAT	PGIB	PGIB_{cont}
MUTAG	74.50±7.89	80.50±7.89	73.50±7.43	80.50±9.07	79.00±6.24	81.00±6.63	80.88±8.94	85.00±7.07	85.50±5.22
PROTEINS	72.83±4.23	70.30±4.84	71.35±4.85	73.83±4.22	75.25±5.92	73.66±3.32	69.64±4.71	77.14±2.19	77.50±2.42
NCI1	73.16±3.49	75.04±2.08	66.05±1.03	74.13±2.10	64.65±6.78	63.75±3.37	68.13±2.64	77.65±2.20	78.25±2.13
DD	72.53±4.51	72.04±3.62	70.81±4.33	69.15±4.33	72.61±8.26	72.77±5.63	71.93±2.74	73.36±1.80	73.70±2.14

Graph Interpretation



- 제안된 모델의 그래프 분류 성능이 여러 최신 모델들을 능가함.
- 탐지된 subgraph가 분류성능을 향상시키는데 기여

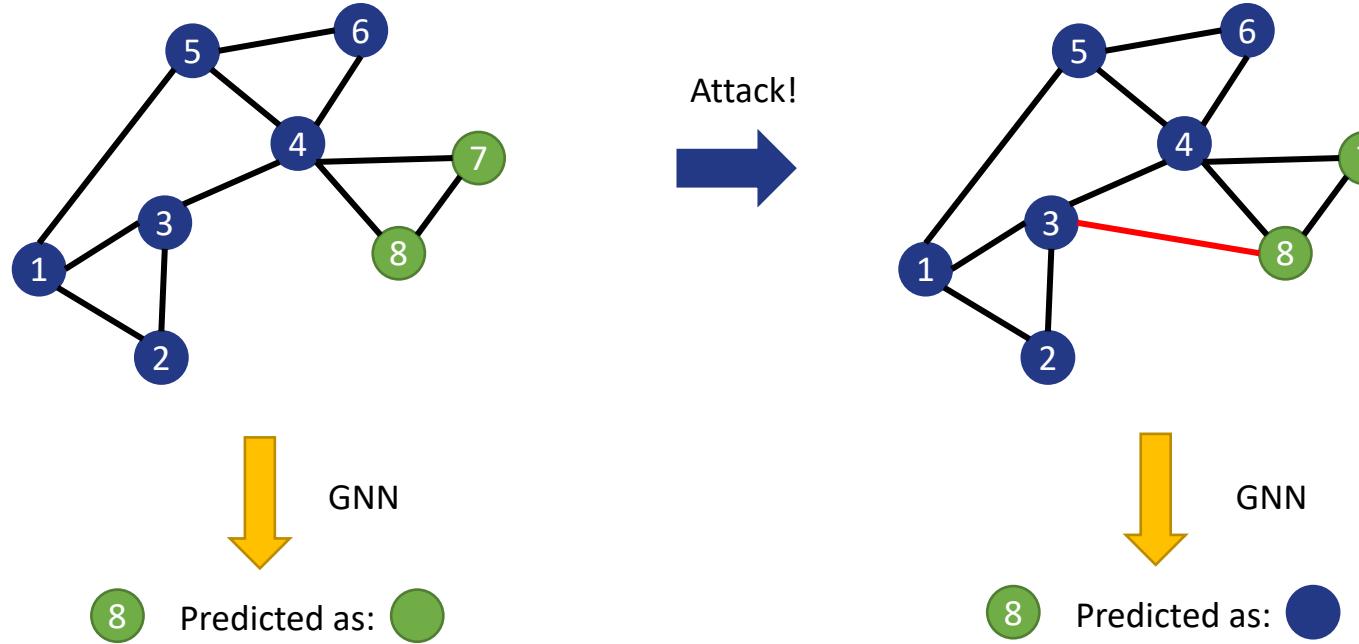
- 제안된 모델에 의해서 탐지된 subgraph가 functional group을 올바르게 포착함

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Background

Adversarial Attacks on Graph Structures



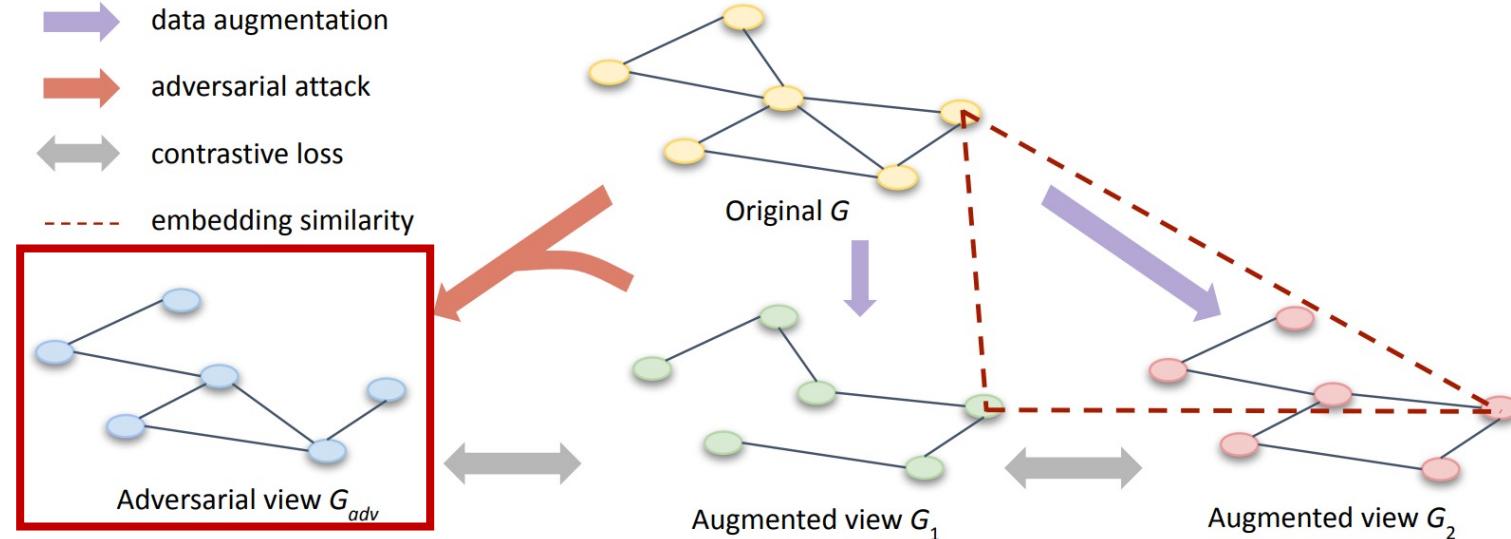
- Graph Neural Networks are vulnerable to adversarial attacks on graph structures.
- Unsupervised GRL models are also vulnerable to such attacks.

→ *Leads to the requirements of robust graph representation learning methods*

Motivation

Applying Adversarial Training (AT) to Graph Contrastive Learning (GCL)

A graph with the worst-case attack
maximizing contrastive loss



Formulation of the adversarial attack in GCL models

$$\delta_A^*, \delta_X^* = \arg \max_{\delta_A, \delta_X \in \Delta} \underset{\text{Contrastive loss}}{\mathbb{E}} [\mathcal{L}(f(A^1 + \delta_A, X^1 + \delta_X), f(A^2, X^2))]$$

$$\Delta = \{(\delta_A, \delta_X) | \|\delta_A\|_0 \leq \underline{\Delta}_A, \|\delta_X\|_0 \leq \underline{\Delta}_X\}$$

Perturbation budgets

- **Goal:** to find the optimal edges and node feature perturbations for the A^1, X^1 that maximally increase the contrastive loss.
- Since we consider unsupervised adversarial attacks, a contrastive loss is employed instead of a supervised loss.

Motivation

Characteristic of Adversarial Attacks on GCL

- If $\mathbf{z}_i^2 - \mathbf{z}_i^{atk}$ is large, δ_A is effective perturbation.
- $\mathbf{z}_i^2 - \mathbf{z}_i^{atk}$ is computed as follows:

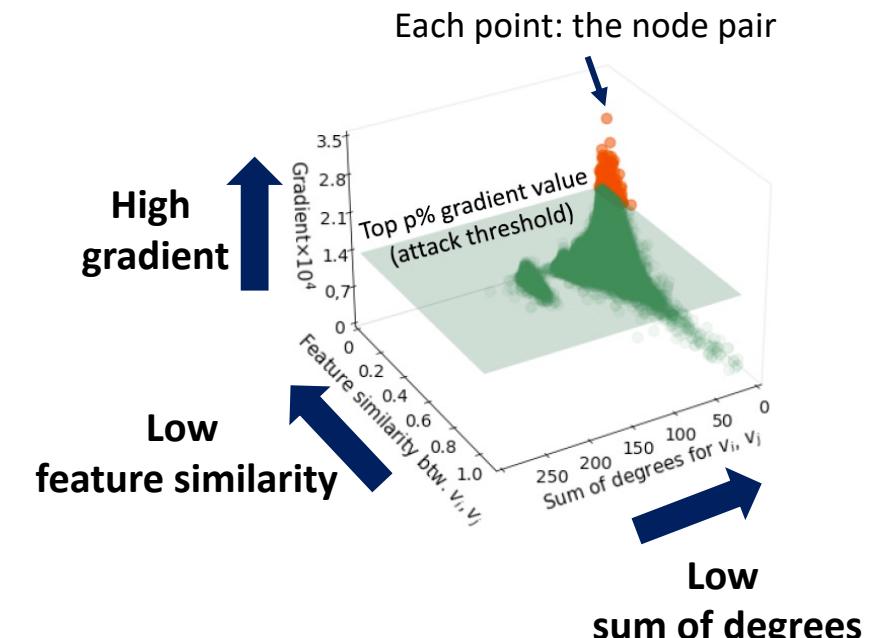
$$\begin{aligned} \mathbf{z}_i^2 - \mathbf{z}_i^{atk} &= (\mathbf{z}_i^2 - \mathbf{z}_i^1) + (\mathbf{z}_i^1 - \mathbf{z}_i^{atk}) \\ &= \mathbf{e}_i + \underbrace{\frac{1}{\sqrt{|\mathcal{N}_{A^1}^i| + 1}} \left(\sum_{j \in \mathcal{N}_{A^1}^i \cup \{i\}} \frac{\alpha \mathbf{Wx}_j}{\sqrt{|\mathcal{N}_{A^1}^i|} \sqrt{|\mathcal{N}_{A^1}^j|}} - \frac{\mathbf{Wx}_k}{\sqrt{|\mathcal{N}_{A^1}^k| + 1}} \right)}_{\text{Degree term}} \quad (4) \end{aligned}$$

Degree term Feature difference term

- $\mathbf{z}_i^2 - \mathbf{z}_i^{atk}$ becomes large when degree term \downarrow and feature diff. term \uparrow
 - The degree of v_i is small (*low-degree nodes*)
 - *The features of node v_k (i.e., \mathbf{x}_k) is dissimilar from the aggregation of neighborhood features in a clean graph.*

Assumption for simplicity

- GCL model with a 1-layer GCN w/o nonlinearity.
- Perturbs only one edge $v_i \rightarrow v_k$.
- Attacked graph $(\mathbf{A}^1 + \delta_A, \mathbf{X}^1)$
- $\mathbf{Z}^{atk} = f(\mathbf{A}^1 + \delta_A, \mathbf{X}^1)$



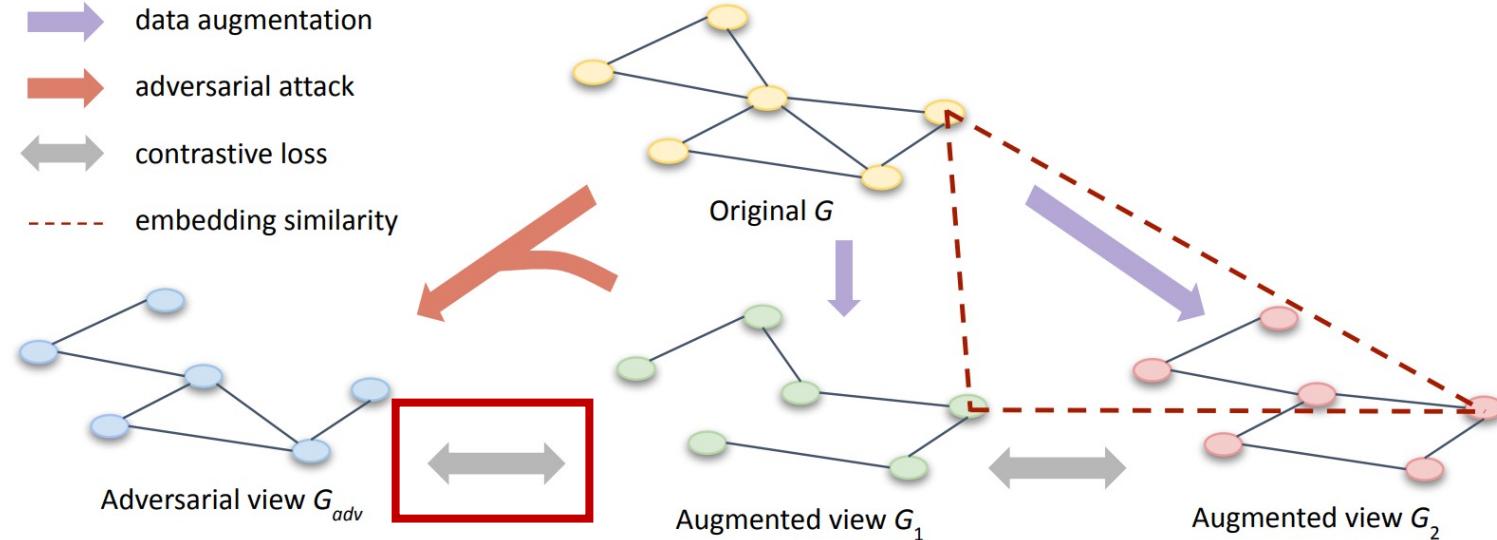
Characteristic of a generated adversarial view by contrastive loss

1. Attack the nodes that have low-degree.
2. **Connect the nodes with dissimilar feature**

Motivation

Applying Adversarial Training (AT) to Graph Contrastive Learning (GCL)

Using the attacked graph as an additional augmentation !



Formulation of Adversarial Graph Contrastive Learning (AGCL)

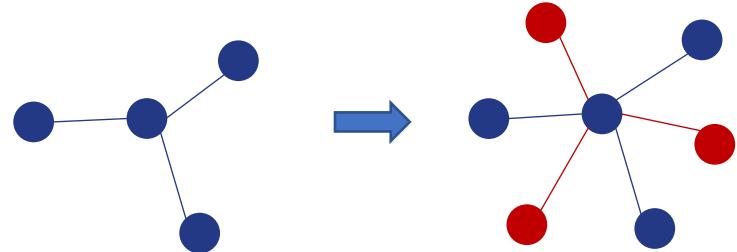
$$\min_{\Theta} \underbrace{\mathcal{L}(Z^1, Z^2)}_{\text{GCL term}} + \lambda_1 \underbrace{\mathcal{L}(Z^1, Z^{adv})}_{\text{AT term}}$$

$$Z^{adv} = f(\underbrace{A^1 + \delta_A^*, X^1 + \delta_X^*}_{\text{Adversarial graph view}})$$

- **Goal:** robust graph representation learning based on adversarial training (AT).
- **Main idea:** to force the representations in the clean graph to be close to those of the attacked graphs.
 - The adversarial graph contrastive learning model minimizes the training objective.

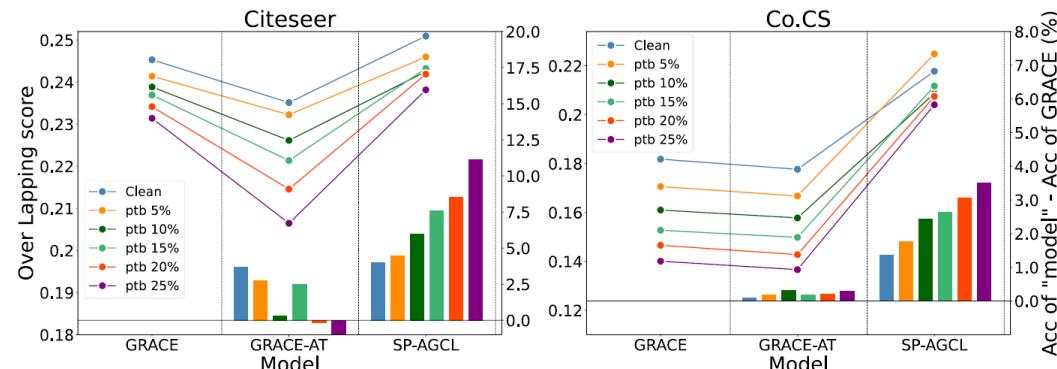
Motivation

AT fails to preserve node similarity !



- As previously demonstrated, adversarial attacks on graphs tend to connect nodes with dissimilar features.
 - The neighborhood feature distribution is changed by the adversarial attacks.
- And AGCL reduces the distance between the clean view and the adversarial view to achieve robustness.
 - Neglecting the changes in the neighborhood feature distributions in the adversarial view.

We argue that existing AGCL models obtain robustness *at the expense of losing the feature information.*



- Solid line: OL score
- Bar plot: performance improvement compared to GRACE

$$OL(A^{kNN(Z)}, A^{kNN(X)}) = \frac{|A^{kNN(Z)} \cap A^{kNN(X)}|}{|A^{kNN(X)}|}$$

- indicates how much the feature information the representations have

We observe

- GRACE-AT have higher **accuracy** than GRACE
 - They obtain robustness.
- GRACE-AT have lower **OL score** than GRACE
 - They lose the feature information.

Motivation

Node similarity preservation is crucial !

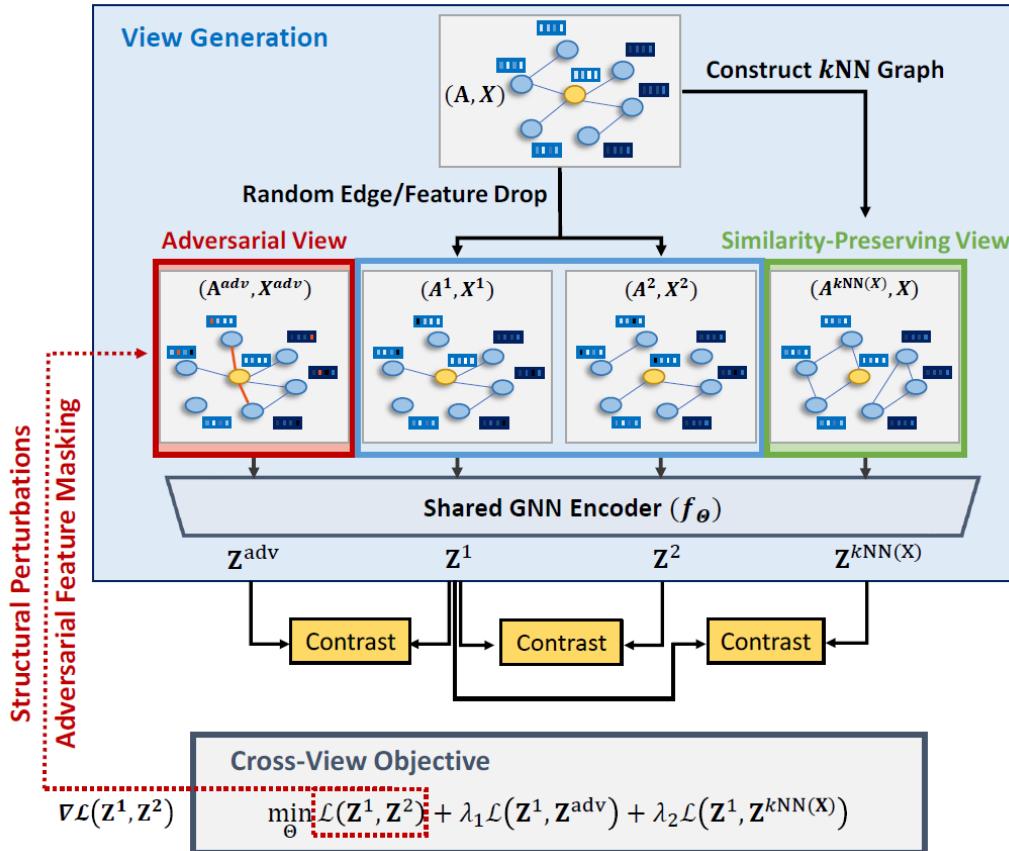
- As previously demonstrated, existing AGCL models obtain robustness *at the expense of losing the feature information.*
- However, **the node feature information is crucial for the robustness against graph structure attacks [1, 2].**

We argue that **the robustness of AGCL model can be further enhanced by fully exploiting the node feature information.**

- Moreover, preserving the node feature similarity becomes especially useful for most real-world graphs.
 - Graphs with noisy node labels
 - Graphs with heterophilous neighbors
 - Low-degree nodes
- To this end, we propose a *similarity-preserving adversarial graph contrastive learning* (SP-AGCL) framework

Proposed Method

Similarity Preserving Adversarial Graph Contrastive Learning (SP-AGCL)



View generation

- *Step 1.* Two stochastically augmented views, $(\mathbf{A}^1, \mathbf{X}^1)$ and $(\mathbf{A}^2, \mathbf{X}^2)$
 - Same as the previous GCL models

• *Step 2. Adversarial View*

- Structural perturbations

$$\frac{\partial \mathcal{L}}{\partial \mathbf{A}^1} + \frac{\partial \mathcal{L}}{\partial \mathbf{A}^2} = \mathbf{G}_A \in \mathbb{R}^{N \times N}$$

- Adversarial feature masking

$$\frac{\partial \mathcal{L}}{\partial \mathbf{X}^1} + \frac{\partial \mathcal{L}}{\partial \mathbf{X}^2} = \mathbf{G}_X \in \mathbb{R}^{N \times F}$$

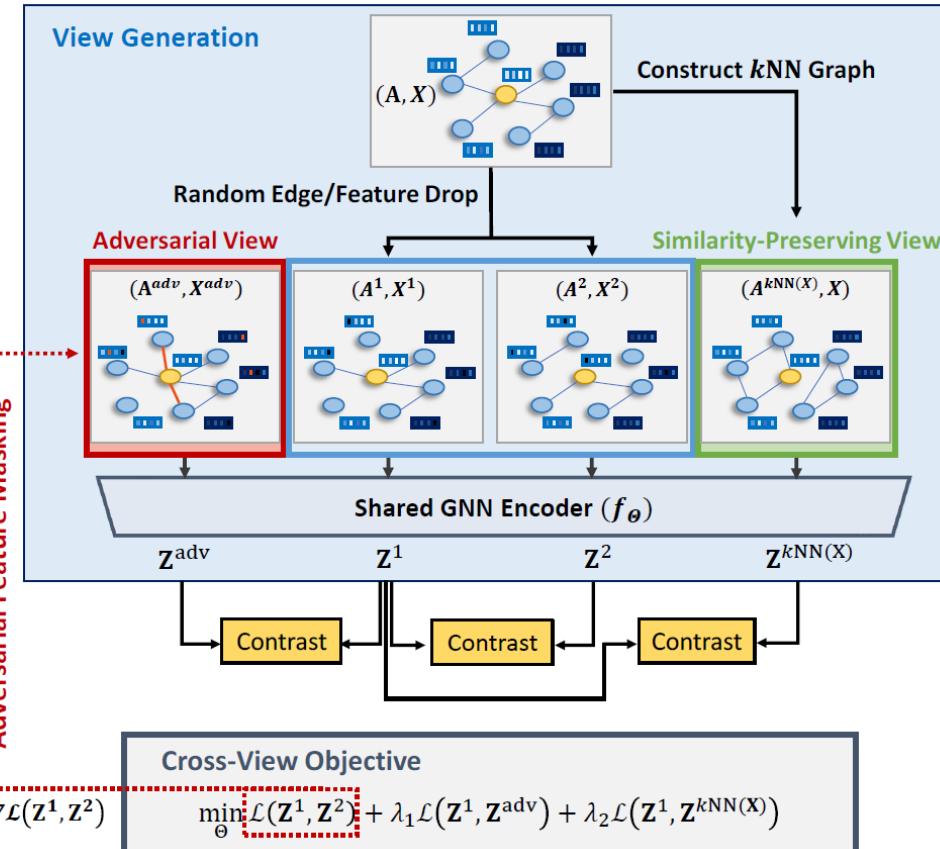
- Existing works flip the node feature
- But, it corrupts the co-occurrence/correlation statistics.
- By masking instead of flipping, we maintain them.

• *Step 3. Similarity preserving view*

- Aims to preserve the node feature similarity.
- kNN graph of node features $(\mathbf{A}^{kNN(\mathbf{X})}, \mathbf{X})$

Proposed Method

Similarity Preserving Adversarial Graph Contrastive Learning (SP-AGCL)



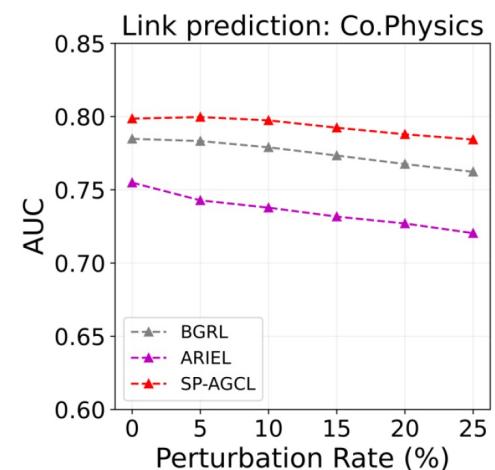
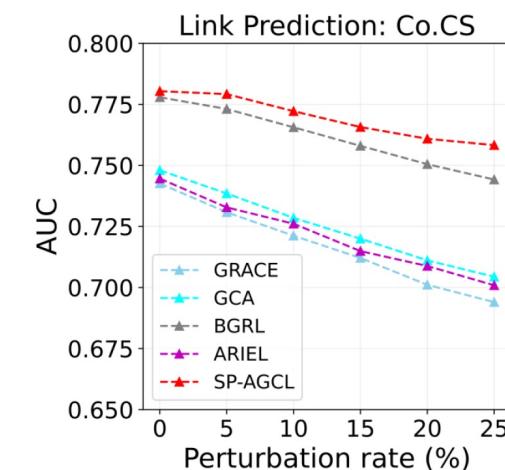
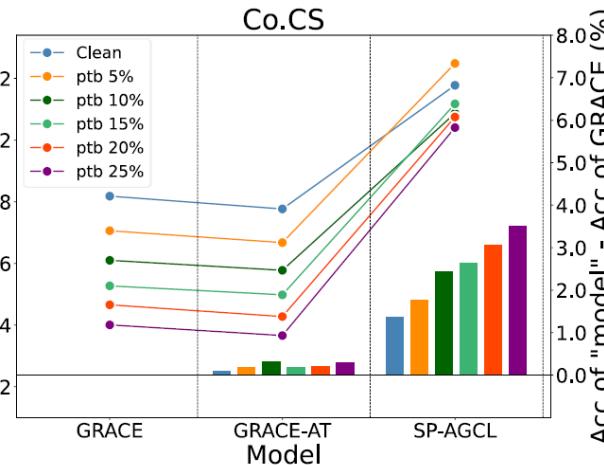
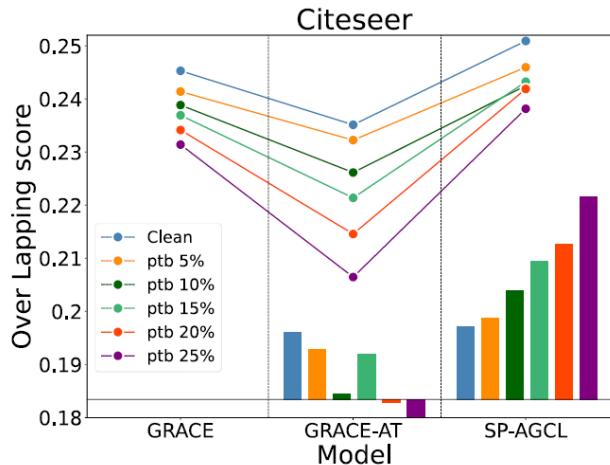
Cross-view Training for Robust GCL

$$\min_{\Theta} \frac{\mathcal{L}(Z^1, Z^2)}{\text{GCL term}} + \lambda_1 \frac{\mathcal{L}(Z^1, Z^{adv})}{\text{AT term}} + \lambda_2 \frac{\mathcal{L}(Z^1, Z^{kNN(X)})}{\text{Similarity-preserving term}}$$

The **representations of nodes with similar features are pulled together**, which in turn preserves the node feature similarity.

Experiment

Preserving Feature Similarity is beneficial !



- SP-AGCL preserves the node feature similarity, which results in the robust graph representation.
- SP-AGCL consistently predicts reliable links compared with other baselines across all the perturbation ratios.
 - Moreover, ARIEL, the sota AGCL model, shows the worst performance

Node feature information is beneficial to predicting reliable links
since nodes with similar features tend to be adjacent in many real-world graphs.

Outline

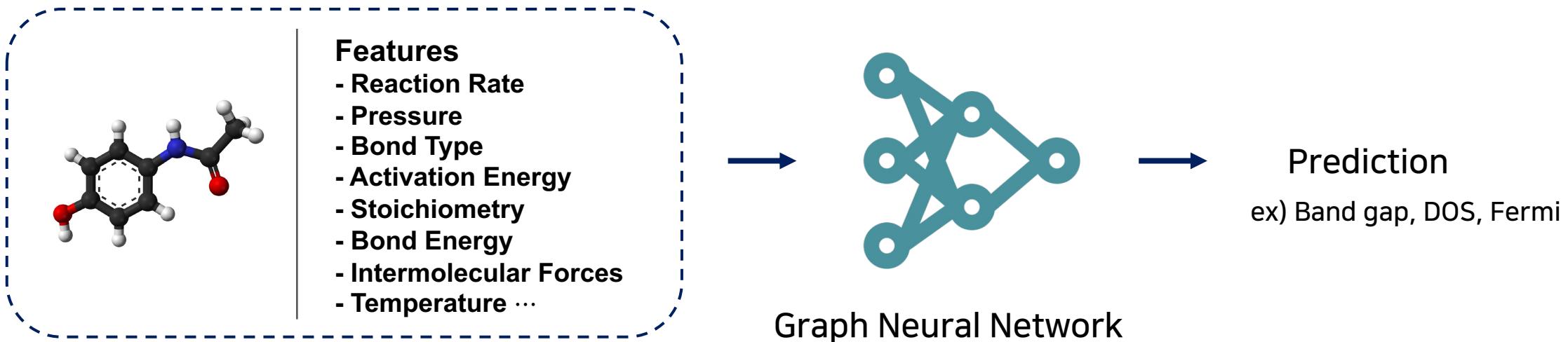
- Overview
- Random walk-based Methods
- Graph Neural Networks (GNNs)
- How to effectively train GNNs?
 - Self-supervised Learning (AAAI'22)
 - Explainable Model (NeurIPS'23)
 - Robustness (KDD'23)
- Applications of GNNs
 - GNNs for Science (화학/소재/생물) (ICML'23)
 - GNNs for Medical Data (멀티모달 의료 데이터 분석) (AAAI'23)
 - GNNs for Computer vision (Scene Understanding) (AAAI'23)

Outline

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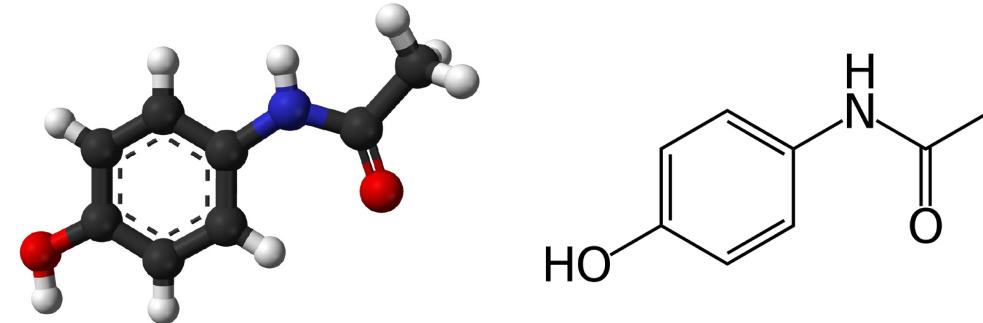
Introduction: Molecular Property Prediction

- Predict the properties of a molecule (소재 물성 예측)



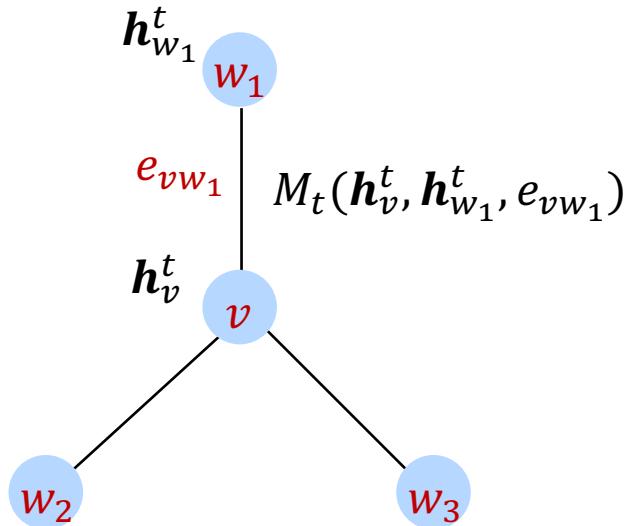
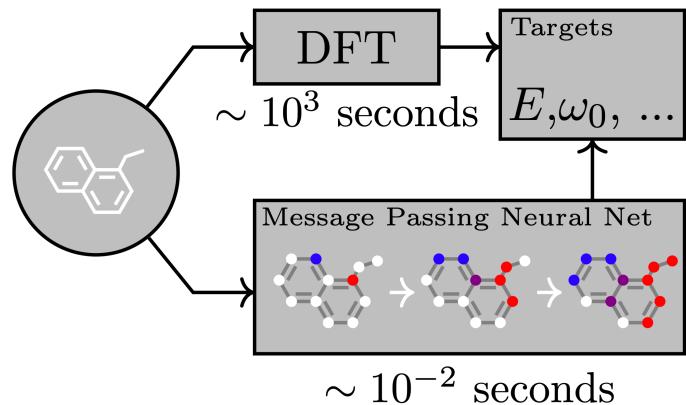
Molecular Graphs

- Molecules can be represented as a graph with node features and edge features
 - Node features: atom type, atom charges...
 - Edge features: valence bond type...



Message Passing Neural Network

- Unified various graph neural network and graph convolutional network approaches



$$m_v^{t+1} = \sum_{w \in N(v)} M_t(h_v^t, h_w^t, e_{vw})$$

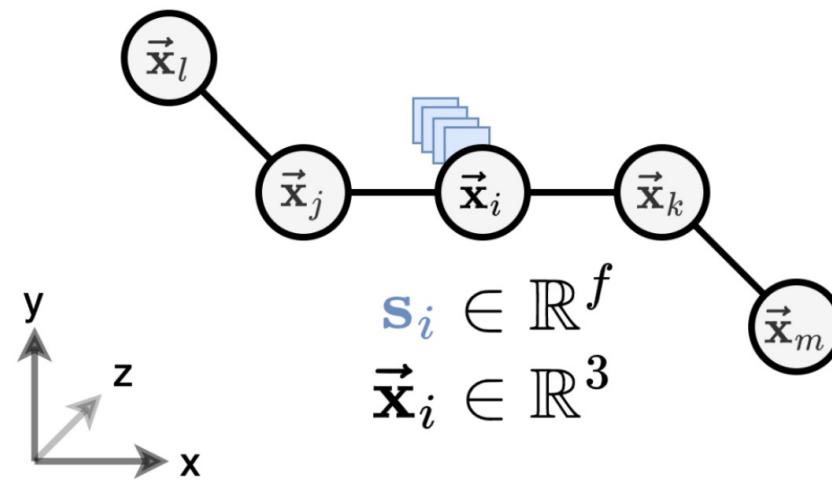
Edge embedding
Neighbor of v

$$h_v^{t+1} = U_t(h_v^t, m_v^{t+1})$$

$$\hat{y} = R(\{h_v^T \mid v \in G\})$$

Geometric Graphs

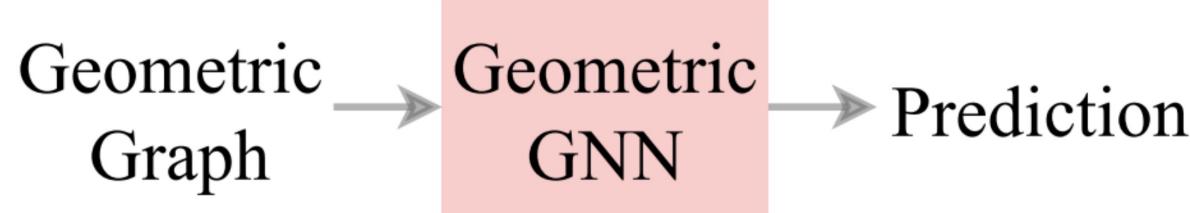
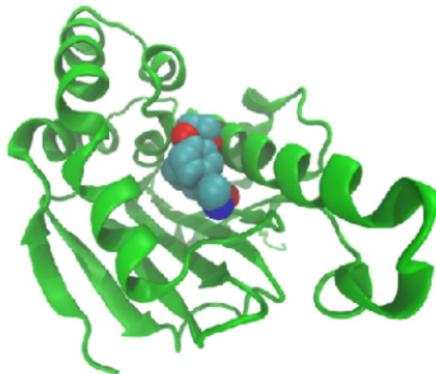
- Sometimes, we also know the 3D positions of atoms, which is actually more informative
- A geometric graph $G = (A, S, X)$ is a graph where each node is embedded in d -dimensional Euclidean space:



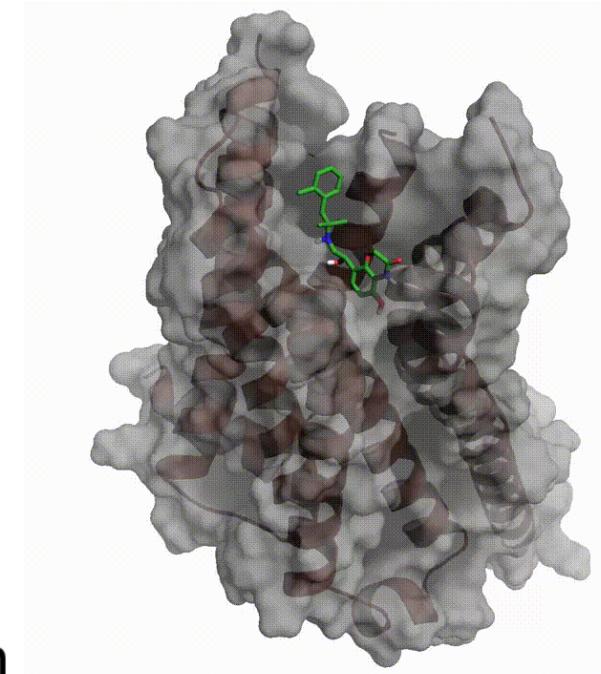
- A : an $n \times n$ adjacency matrix
- $S \in \mathbb{R}^{n \times f}$: Scalar features (atom type, atom charges, ...)
- $X \in \mathbb{R}^{n \times d}$: tensor features, e.g., coordinates

Broad Impact on Sciences

- Supervised Learning: Prediction
 - Properties prediction
 - 3D Protein-ligand interaction (binding)



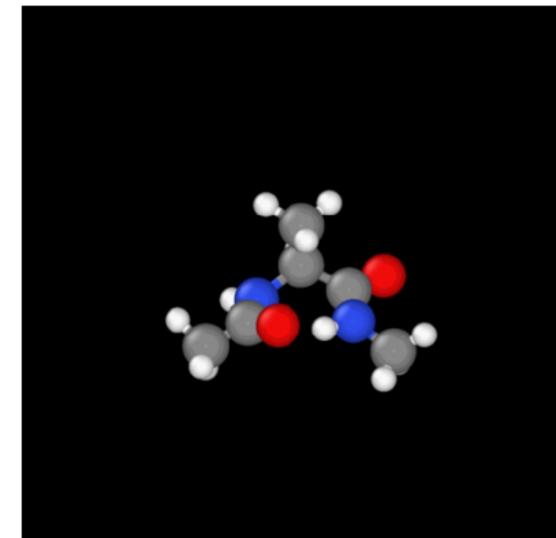
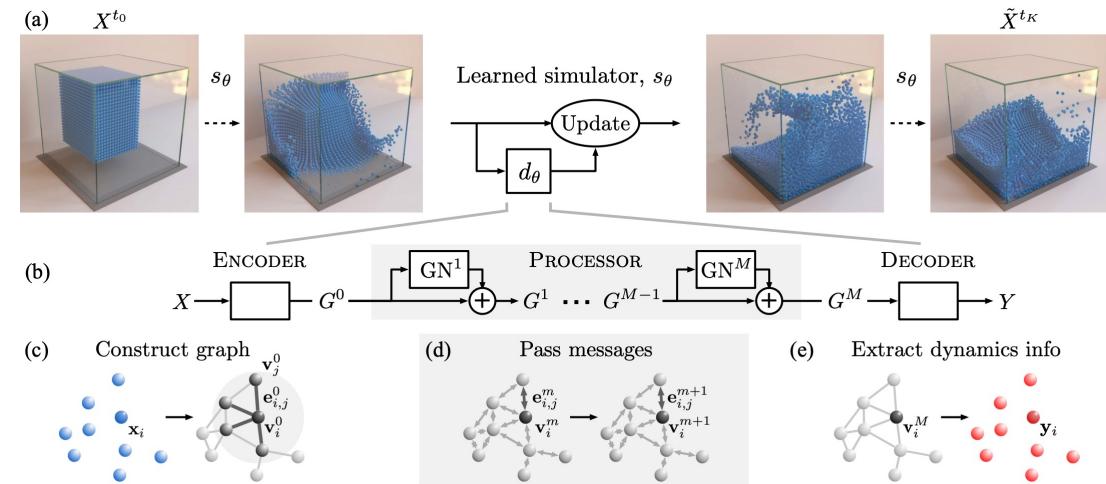
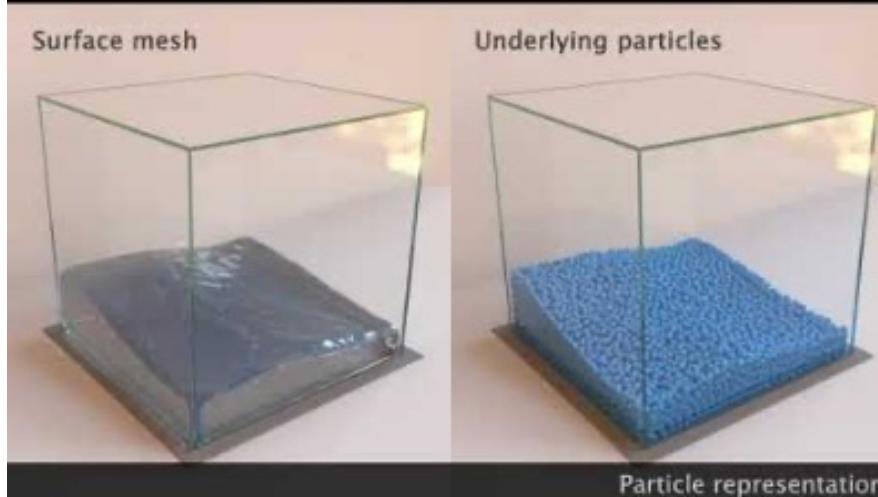
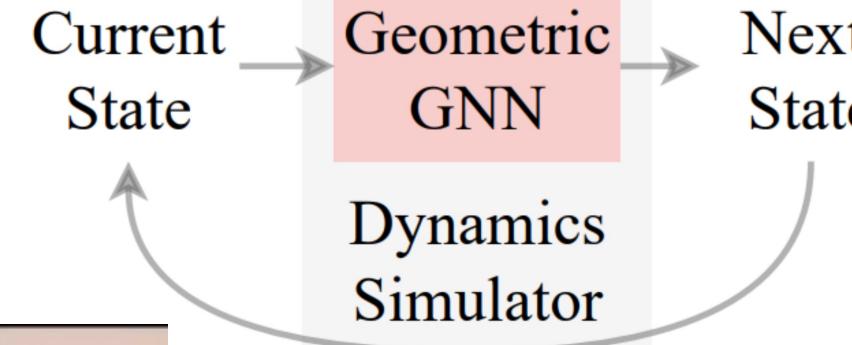
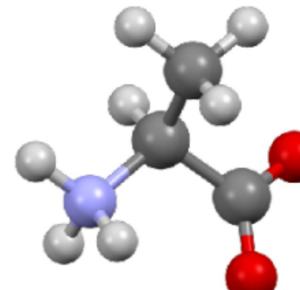
- Functional properties?
- Ligand binding affinity?
- Ligand efficacy?



Broad Impact on Sciences

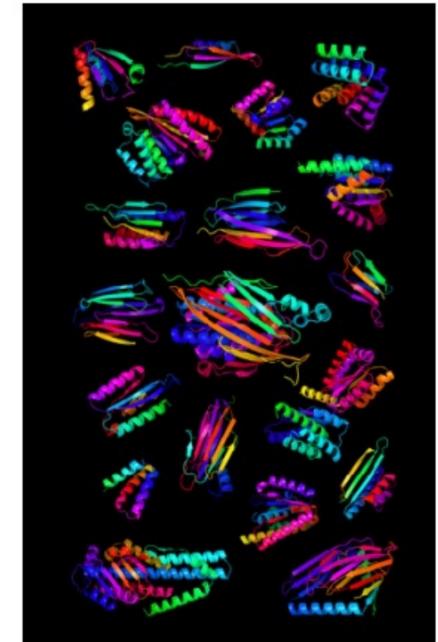
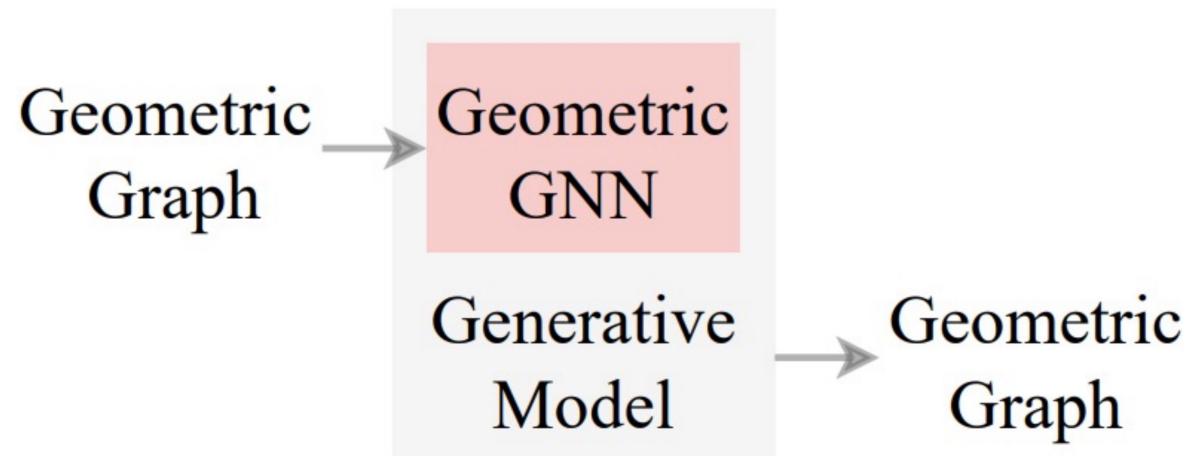
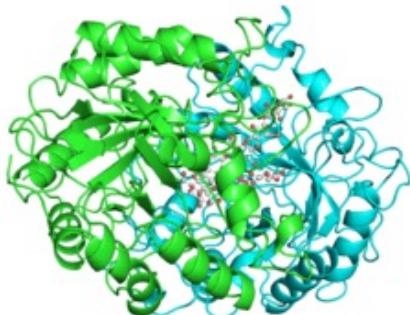
■ Supervised Learning: Structured Prediction

- Molecular Simulation



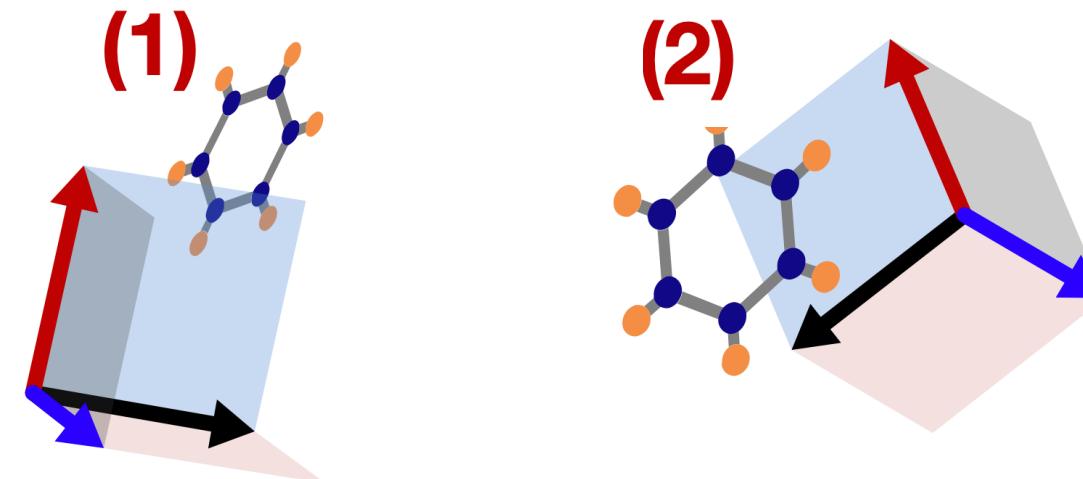
Broad Impact on Sciences

- Generative Models
 - Drug or material design



Geometric graph is more challenging than Molecular graph

- To describe geometric graphs, we use **coordinate systems**
 - (1) and (2) use different coordinate systems to describe the **same** molecular geometry.
- We can describe the transform between coordinate systems with **symmetries** of Euclidean space
 - 3D rotations, translations



However, output of traditional GNNs given (1) and (2) are completely different!
→ Enforcing symmetry is crucial (Invariant GNNs)

Schnet: Overview

■ Input

- Feature representations of n atoms $X^l = (\mathbf{x}_1^l, \dots, \mathbf{x}_n^l)$ with $\mathbf{x}_i^l \in R^F$
- At locations $R = (\mathbf{r}_1, \dots, \mathbf{r}_n)$ with $\mathbf{r}_i \in R^D$ ($D = 3$ for 3-dim coordinates)

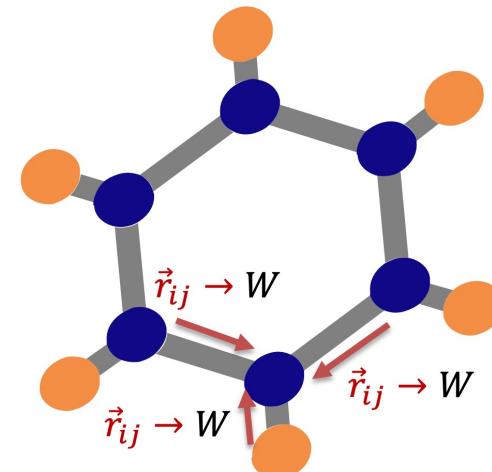
■ Output

- Molecular total energy $E(\mathbf{r}_1, \dots, \mathbf{r}_n)$

■ SchNet updates the node embeddings at the l -th layer by message passing layers

$$\mathbf{x}_i^{l+1} = (X^l * W^l)_i = \sum_j \mathbf{x}_j^l \circ W^l(\mathbf{r}_i - \mathbf{r}_j),$$

- A filter generating function $W^l: R^D \rightarrow R^F$ is determined by the relative position from neighbor atoms j to i
- \circ is the element-wise multiplication



x^l : node embeddings at l layer
 r : atomic coordinates

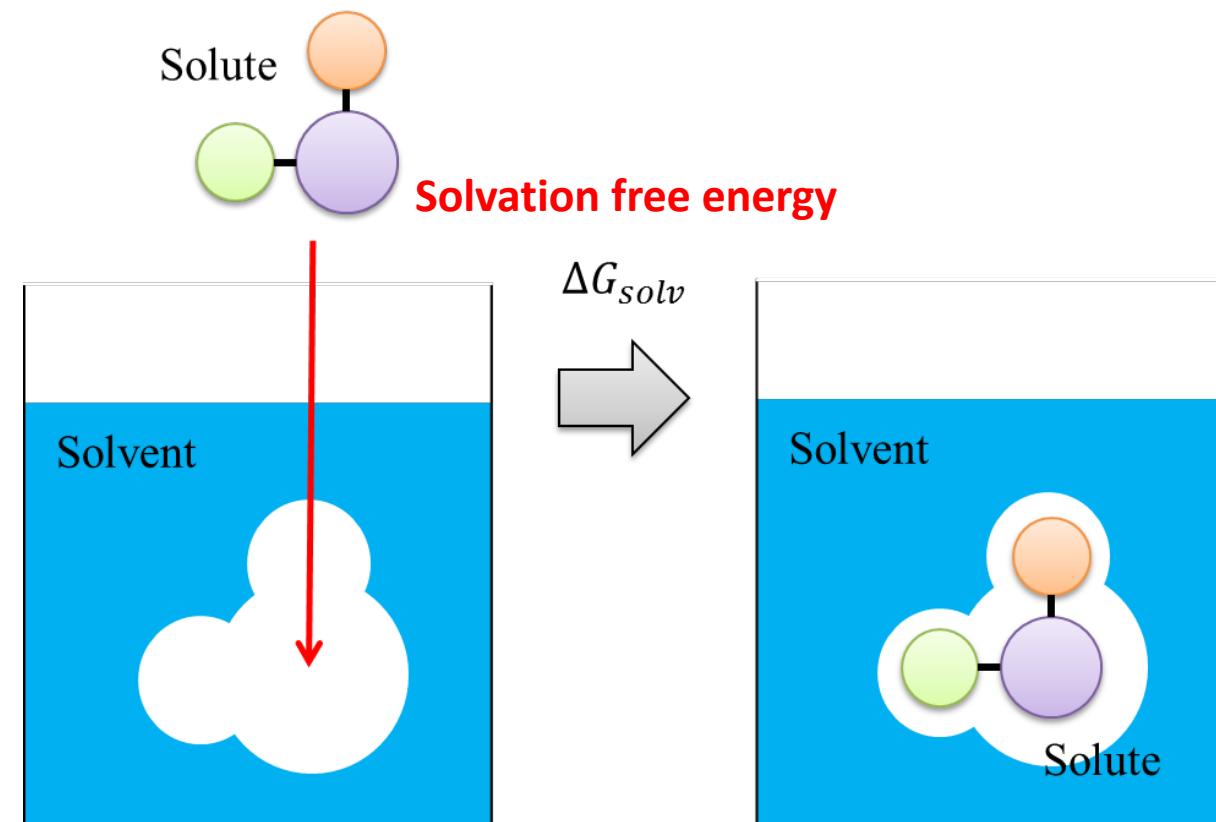
Schnet: Invariance

- W is invariant by scalarizing relative positions with relative distances ($\|r_i - r_j\| = \|r_{ij}\| = d_{ij}$)
 - $\|r_{ij}\|$ is invariant to **rotations** and **translations**
- Hence, each message passing layer W^l is invariant
 - Aggregated node embeddings $\sum_j \mathbf{x}_j^l \circ W^l(\mathbf{r}_i - \mathbf{r}_j)$ is invariant
 - **Node embeddings are invariant!**

Predicting Solvation Free Energy (용매화 자유 에너지)

▪ Solvation free energy

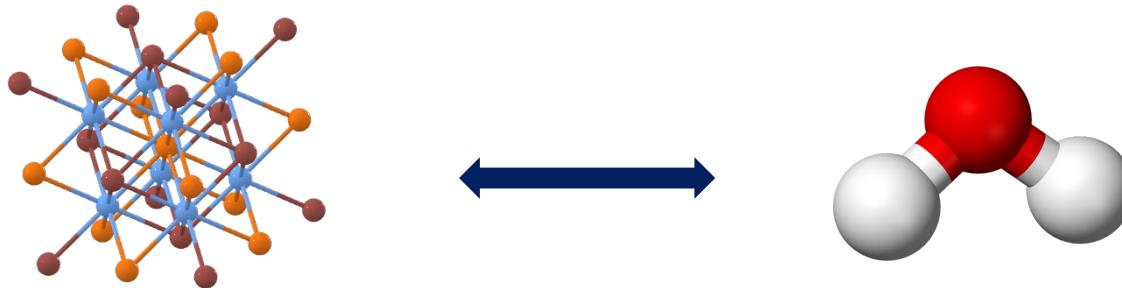
- Change in free energy for a molecule to be transferred from gas phase to a given solvent
- Quantifies **solvability** of drug molecules
 - A large negative value → high solubility
 - A lower magnitudes/positive value → poor solubility



Introduction: Relational Learning

▪ Relational Learning

- 두 개체 사이의 관계를 예측하는 기계 학습 분야
- 특히, 두 화학 물질 사이의 관계를 예측하는 것은 화학 분야에서 매우 중요함



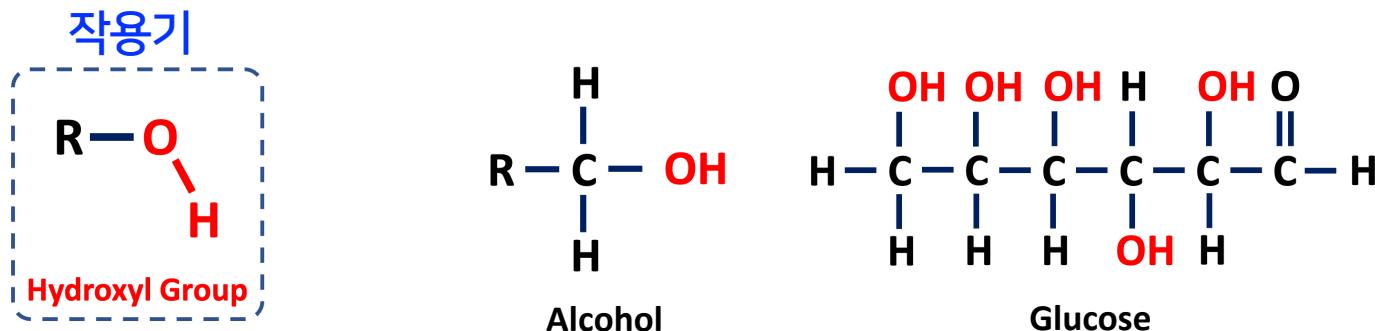
• Examples

- 발색체 (Chromophore)와 용매 (Solvent)가 반응했을 때의 광학적 성질 예측
- 용질 (Solute)과 용매 (Solvent) 가 반응했을 때의 용해도 예측
- 두 종류의 약물 (Drug) 을 동시에 섭취했을 때의 부작용 예측

Introduction: Functional Group

■ 작용기 (Functional Group)

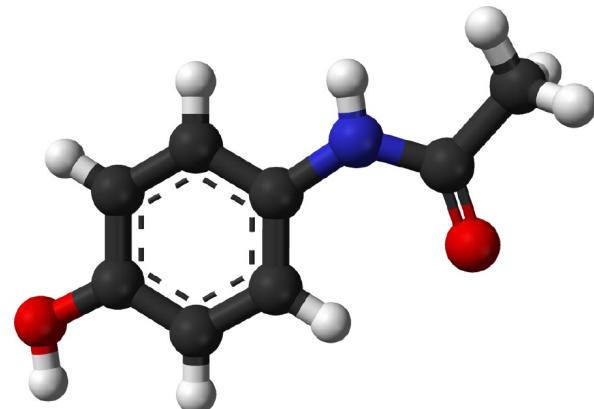
- 유기화합물의 화학 반응적 특성을 결정하는데 중요한 역할을 하는 특정 원자단 혹은 구조
- 유기화학 반응 전후에 작용기의 변환에 따라 반응물과 생성물의 물리적/화학적 성질이 변화함
- 같은 작용기를 갖는 화합물의 특성은 대부분 유사하며, 유사한 화학 반응이 일어남
- Examples
 - Hydroxyl Group 구조는 분자의 극성을 증가시키는 특성을 가지고 있음
 - → Alcohol, Glucose 같이 Hydroxyl 구조를 포함한 분자들은 공통적으로 물에 대해 용해도가 높음



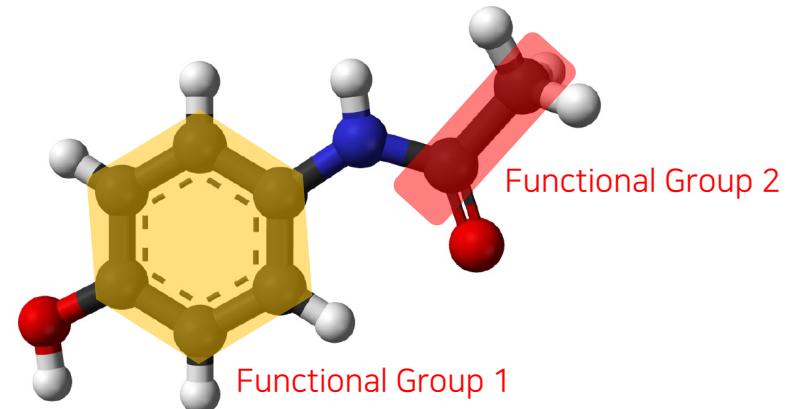
따라서, 화학 물질 사이의 상호작용을 예측하는 모델은 작용기를 기반으로 학습하는 것이 매우 중요함

Introduction: Representing Molecules as a Graph

- Molecule → Graph로 표현 가능
- Functional Group → Subgraph로 표현 가능



Molecule
(=Graph)



Functional Group
(=Subgraph)

최근 Information Theory를 기반으로 Graph 구조에서 중요한 Subgraph를 찾는 모델이 주목

Information Bottleneck

- 기계학습 관점에서 그래프에서 중요한 Subgraph를 어떻게 찾아낼 것인가?

- **Solution: Information Bottleneck Theory**

- 정보에 대한 압축 / 보존의 trade-off에 대해 Theoretical 한 방법론
- Random variable X, Y가 주어졌을 때, Y의 정보를 최대한 유지하면서 X의 정보를 최소한으로 담는 Bottleneck variable T를 학습
 - 즉, input데이터 X를 최대한 압축하면서 target 값 Y는 계속 잘 맞출 수 있는 T를 학습
 - Noise에 robust한 representation을 학습할 때 많이 사용됨

$$\min_T -I(Y;T) + \beta I(X;T)$$

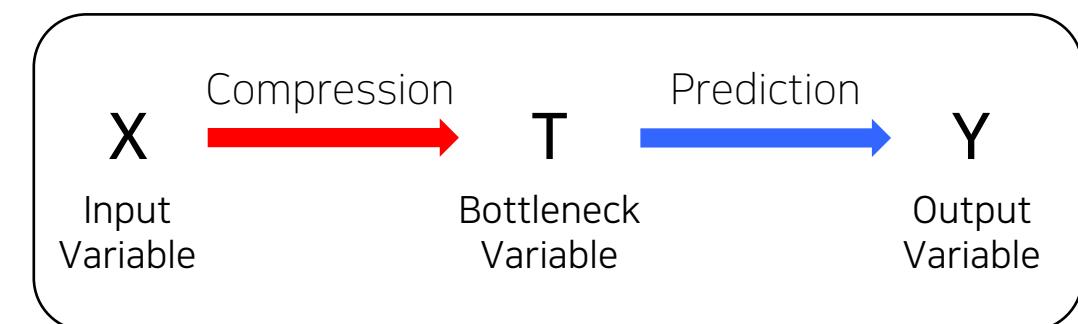
X와 T 사이의 상호 정보량 최소화
→ T가 X에 대한 정보를 최소한으로 갖도록
→ Compression

T와 Y 사이의 상호 정보량 최대화

→ T가 Y에 대한 정보를 최대한 가지고 있어야 함
→ Prediction

Information Bottleneck Objective

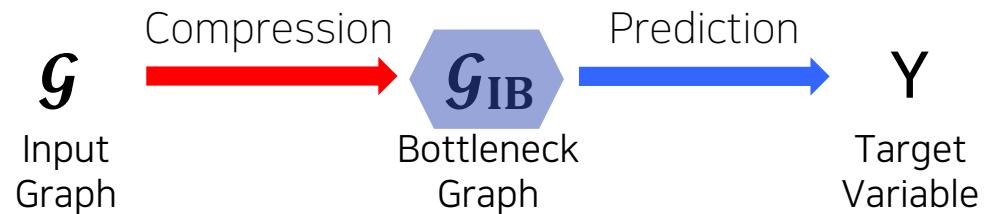
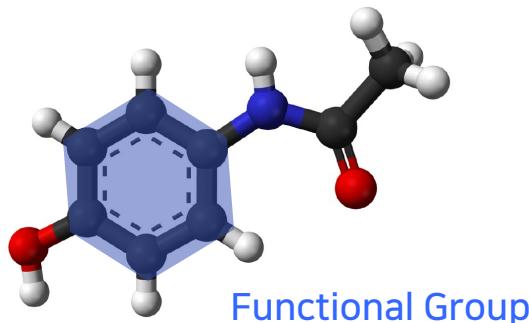
($I(X, Y)$: Mutual information between X and Y)



Graph Information Bottleneck: Overview

- Information bottleneck theory를 그래프에 어떻게 적용할 것인가?
- Information Bottleneck Graph (IB-Graph)
 - 기존 Graph 의 성질을 최대한 보존하는 Subgraph
 - Subgraph를 Bottleneck variable로 모델링
→ Target Y를 맞추는데 가장 중요한 Subgraph G_{IB} 를 찾는 문제로 Formulation

$$\mathcal{G}_{IB} = \arg \min_{\mathcal{G}_{IB}} -I(Y; \mathcal{G}_{IB}) + \beta I(\mathcal{G}; \mathcal{G}_{IB})$$



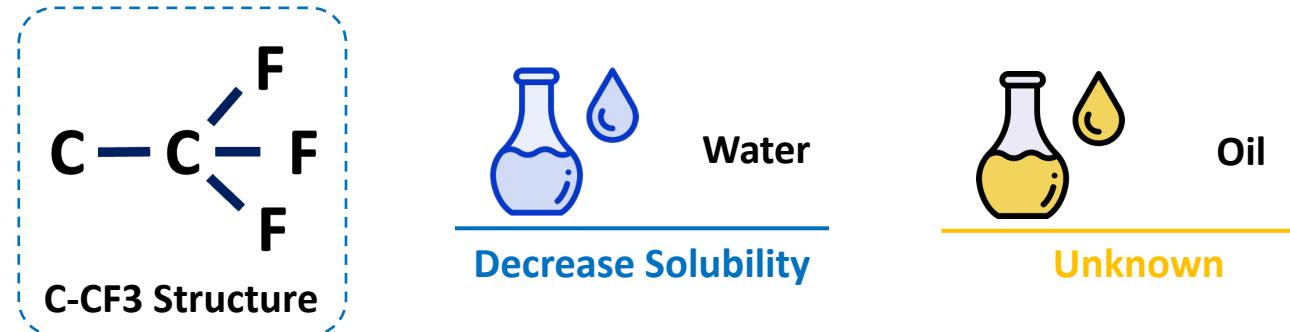
Recall: Functional Group

▪ 작용기 (Functional Group)

- 유기화합물의 화학 반응적 특성을 결정하는데 중요한 역할을 하는 특정 원자단 혹은 구조
- 같은 작용기를 갖는 화합물의 특성은 대부분 유사하며, 유사한 화학 반응이 일어남

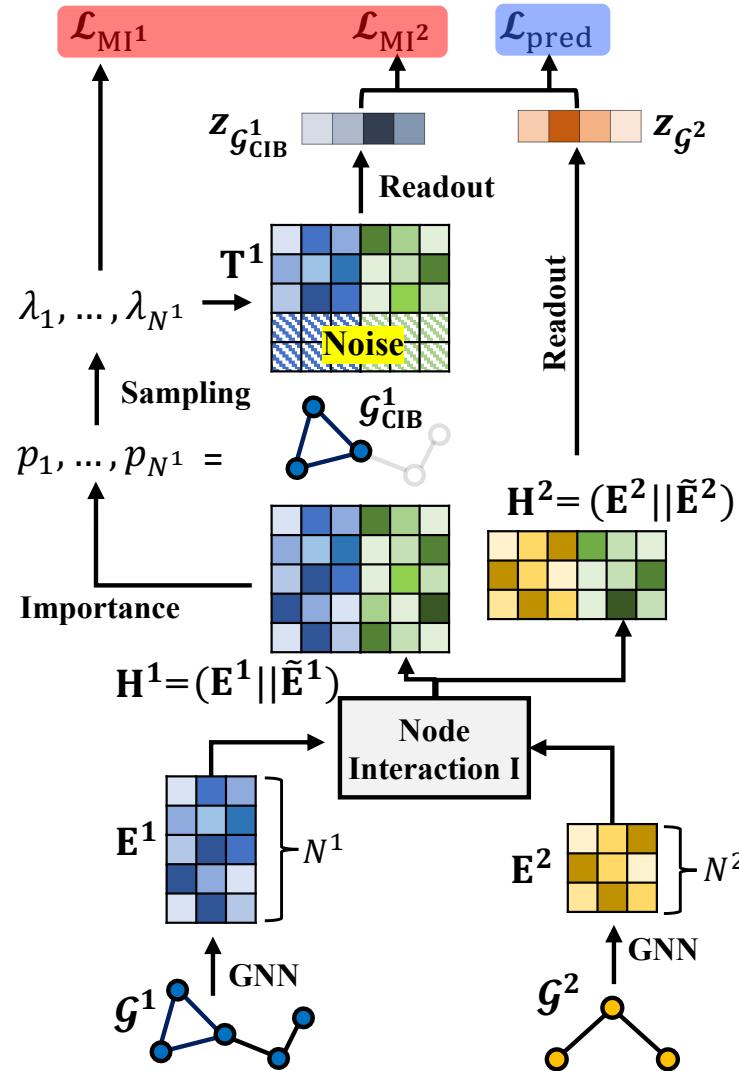
▪ 한편, 용질 (Chromophore)이 어떤 용매 (Solvent)와 반응하는지에 따라서 중요하게 작용하는 작용기가 다름

- Examples: C-CF₃ 구조는 분자의 물에 대한 용해도를 낮추는 역할을 함
 - 하지만, C-CF₃ 구조가 분자의 기름에 대한 용해도에 미치는 영향에 대해서는 알려진 바가 없음
 - 따라서, 화학 반응에서 중요한 작용기를 추출할 때 용매의 종류를 고려할 필요성이 있음



기존의 Information Bottleneck이론으로는 Material Science 지식을 제대로 모델링할 수 없음

Proposed Method: Conditional Graph Information Bottleneck



$$\min -I(Y; \mathcal{G}_{CIB}^1 | \mathcal{G}^2) + \beta I(\mathcal{G}^1; \mathcal{G}_{CIB}^1 | \mathcal{G}^2)$$

Mutual Information 의 Chain Rule에 따라 Conditional Mutual Information 을 분해 후 각 term 의 upper bound 유도

$$-I(Y; \mathcal{G}_{CIB}^1 | \mathcal{G}^2) = -I(Y; \mathcal{G}_{CIB}^1, \mathcal{G}^2) + I(Y; \mathcal{G}^2)$$

$$-I(Y; \mathcal{G}_{CIB}^1, \mathcal{G}^2) \leq \mathbb{E}_{\mathcal{G}_{CIB}^1, \mathcal{G}^2, Y} [-\log p_\theta(Y | \mathcal{G}_{CIB}^1, \mathcal{G}^2)]$$

Prediction Loss

$$I(\mathcal{G}^1; \mathcal{G}_{CIB}^1 | \mathcal{G}^2) = I(\mathcal{G}_{CIB}^1; \mathcal{G}^1, \mathcal{G}^2) - I(\mathcal{G}_{CIB}^1; \mathcal{G}^2)$$

$$I(\mathcal{G}_{CIB}^1; \mathcal{G}^1, \mathcal{G}^2) \leq \mathbb{E}_{\mathcal{G}^1, \mathcal{G}^2} \left[-\frac{1}{2} \log A + \frac{1}{2N^1} A + \frac{1}{2N^1} B^2 \right]$$

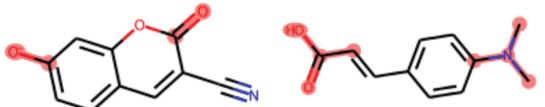
$$:= \mathcal{L}_{MI^1}(\mathcal{G}_{CIB}^1, \mathcal{G}^1, \mathcal{G}^2)$$

Compression Loss

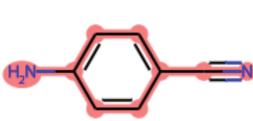
$$-I(\mathcal{G}_{CIB}^1; \mathcal{G}^2) \leq \mathbb{E}_{\mathcal{G}_{CIB}^1, \mathcal{G}^2} [-\log p_\xi(\mathcal{G}^2 | \mathcal{G}_{CIB}^1)]$$

$$:= \mathcal{L}_{MI^2}(\mathcal{G}_{CIB}^1, \mathcal{G}^2)$$

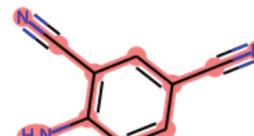
Result: Qualitative analysis



(a) Reaction with ordinary solvent



(b) Reaction with liquid oxygen solvent

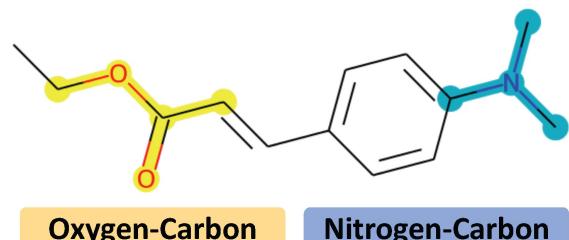


(a) 일반적인 solvent와 반응

화학 구조의 끝 부분을 중요시 여기는 기존의 화학 지식과 align

(b) 단순한 solvent와 반응 (ex. 액체 산소)

chromophore의 전체 부분이 반응한다고 여기는 기존의 화학 지식과 align



Oxygen-Carbon

Nitrogen-Carbon

(c) Chromophore: EDAC

(c) 다양한 solvent와 반응 (Trans-ethyl p-(dimethylamino)cinnamate (EDAC))

solvent의 화학적 극성에 따라 같은 Chromophore에서도 중요한 부분을 다르게 탐지함

- Case1) Benzene solvent

Nitrogen-carbon 구조와 같은 극성이 낮은 구조가 중요함
→ Benzene같은 무극성 solvent와 반응하기 때문

- Case2) Ethanol, THF, 1-hexanol, 1-butanol solvent

Oxygen-carbon 구조와 같은 극성이 높은 구조가 중요함
→ Ethanol과 THF 같은 극성 solvent와 반응하기 때문
→ 일반적으로 1-hexanol과 1-butanol은 무극성 solvent로 분류되지만
-OH 구조로 인해 local 한 극성을 띠는 특성을 가짐

Outline

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Problem definition

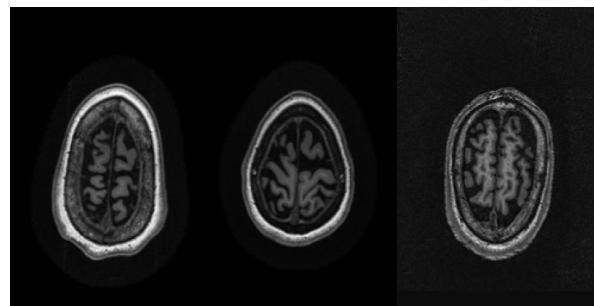
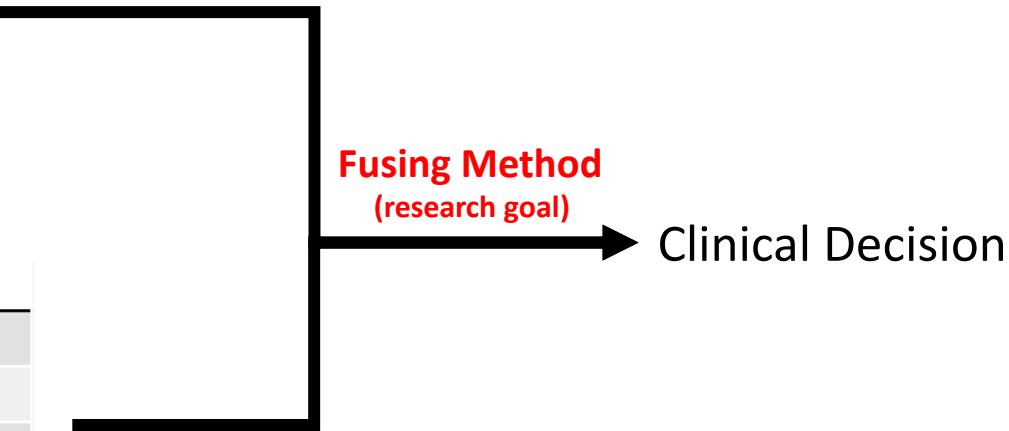


Image data

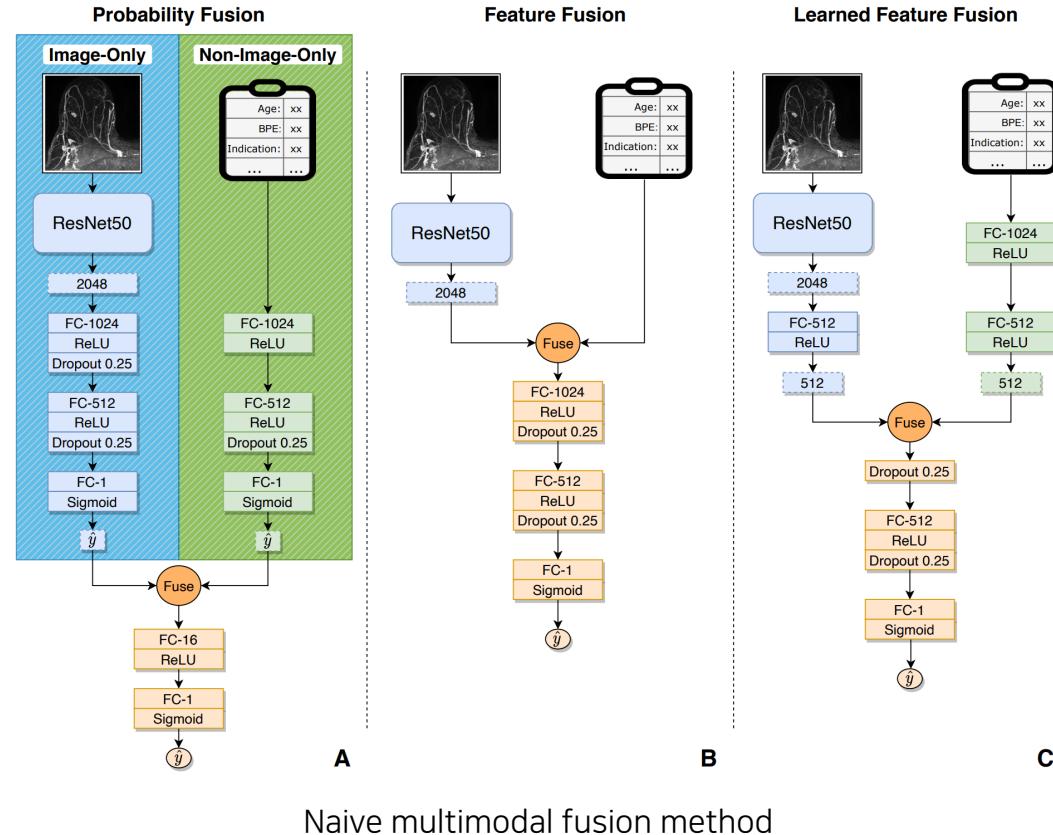
Patient	Age	DIG	TRA	ADA	MMS	City	APOE	RAV
1	42	34	-4	18.6	28	2	0	6
2	73	25	-15.8	31	20	9	1	4
3	80	38	-1.5	14.6	29	9	0	4
4	29	25	-9.4	21.3	27	4	1	4
5	65	34	-10.8	25.6	25	7	0	5

Non-image data



- Integrate the multi-modal medical data (image and non-image data) for more accurate clinical decisions
- Capture important information from various aspects of the given data

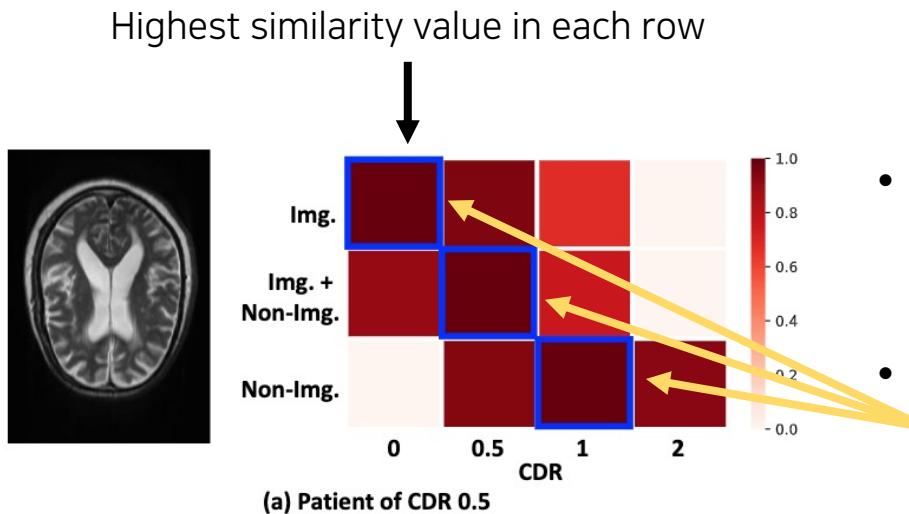
Previous works



- Previous work demonstrates incorporating non-image data with images can significantly improve predictive performance
- But naïve integration of the modalities **cannot fully benefit from the complementary relationship** between the modalities.

Motivation

Medical data is multi-modal in nature

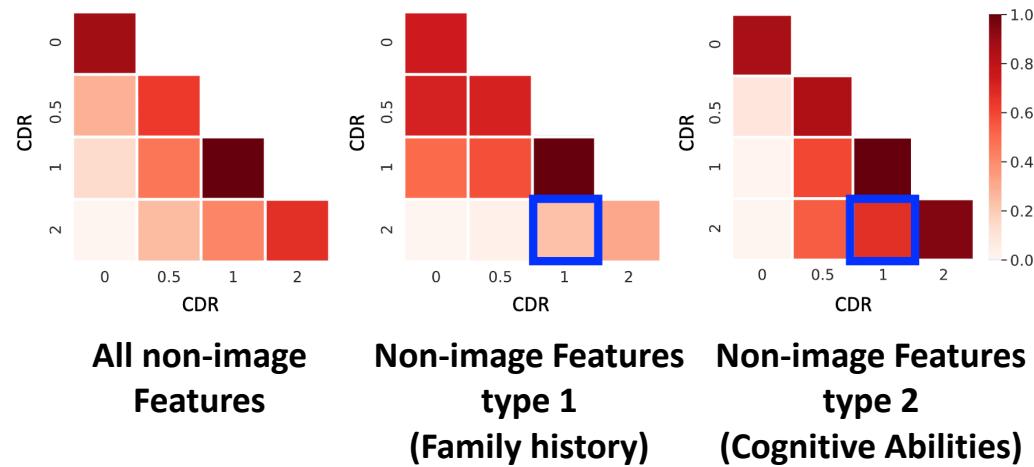


- Routine clinical visits of a patient produce not only image data, but also non-image data (i.e., clinical information).
- Multiple modalities of medical data provide different and complementary views of the same patient.

Integration of diverse and complementary views from medical data can make more informed and accurate clinical decisions.

Motivation

Diverse aspects of Clinical data provides rich information on patients



- Patients who suffer from the same disease share highly similar non-image data compared with those in different classes
- But diverse aspects of non-image data induce complex similarity relationships between patients

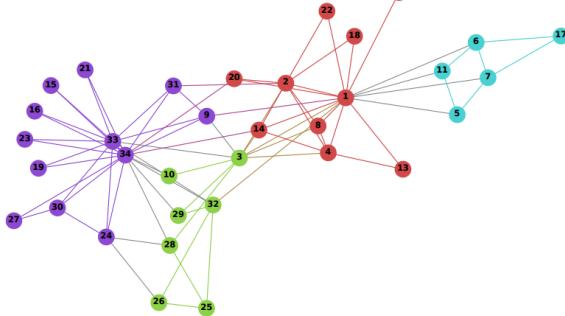
Example

- Type 1 feature fail to find connection between CDR 1 and 2
- Type 2 feature can make preemptive clinical decision on CDR 1

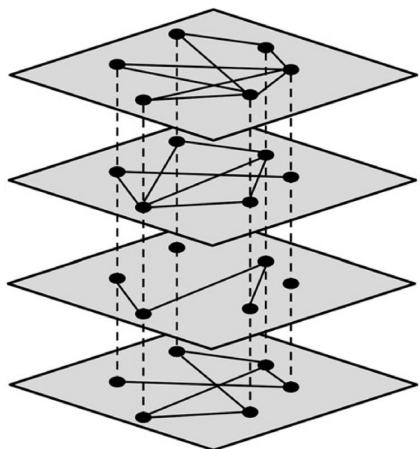
RQ1: How to incorporate complex similarity relationship between patients?

Methodology

How can we capture the relationship between patients



Single Graph

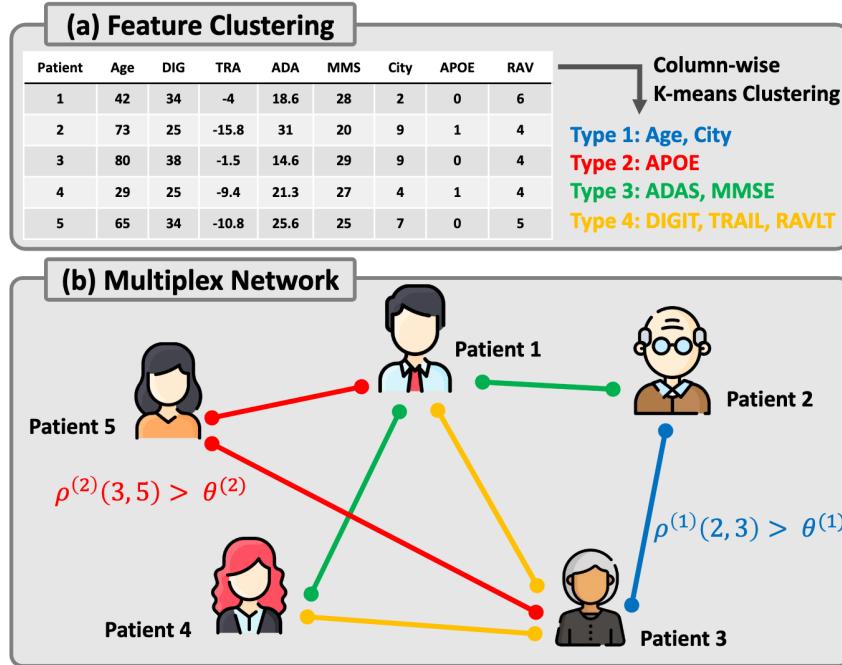


Multiplex Network

- Single graph may miss the inherent complex relationships between patients
 - Single graph considers all relationships as the same
 - Whereas in reality, different relationships can have different characteristics and properties
- Multiplex network considers the **multiple relationships** between nodes as different layers in the network
 - Each layer represents a specific type of relationship
 - Multiplex network can capture the complex relationships between patients, as it acknowledges the heterogeneity of relationships in the network

Methodology

How can we capture the inherent complex relationship between patients ?



$$\mathbf{C} \in \mathbb{R}^{|\mathcal{V}| \times F_{non-img}} \xrightarrow{\text{Column-wise K-means}} \mathbf{c}^{(1)}, \mathbf{c}^{(2)}, \dots, \mathbf{c}^{(r)}, \dots, \mathbf{c}^{(K)}$$

$$\mathbf{A}^{(r)}(i, j) = \begin{cases} 1 & \text{if } \rho^{(r)}(i, j) > \theta^{(r)}, \\ 0 & \text{otherwise} \end{cases}$$

$$\rho^{(r)}(i, j) = \frac{\mathbf{c}_i^{(r)} \cdot \mathbf{c}_j^{(r)}}{\|\mathbf{c}_i^{(r)}\| \cdot \|\mathbf{c}_j^{(r)}\|}$$

- Using Column-wise K-means Clustering, divide non-image data into non-overlapping $|\mathcal{R}|$ types of features
- Using cosine-similarity and threshold for each type of non-image feature, construct multiplex network

Methodology

Overall Frameworks

Overall Frameworks of HetMed

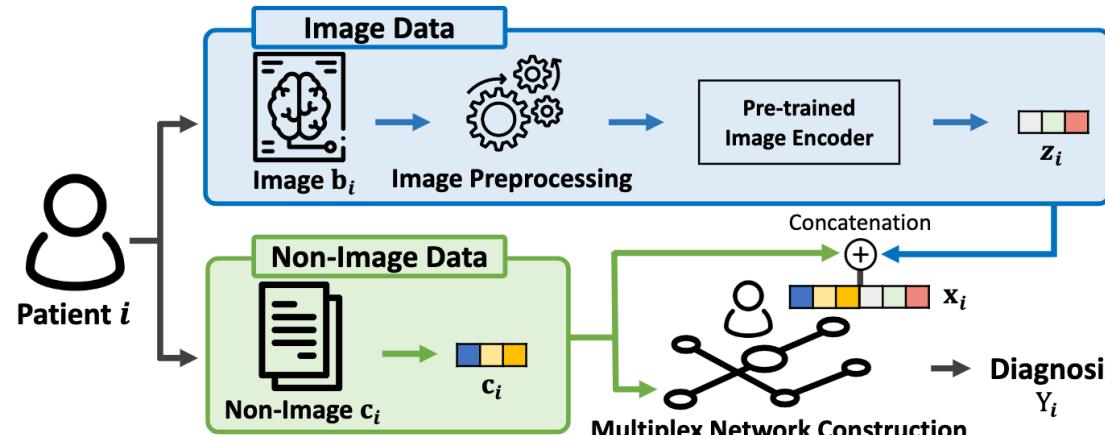


Image Data

- Image Preprocessing (atlas transform)
- Learn the image encoder (pre-trained with non-medical image)
- Extract embeddings of images

Non-image Data

- Column wise K-means Clustering
- Construct multiplex network via cosine similarity of each patient

Learn Multiplex Network

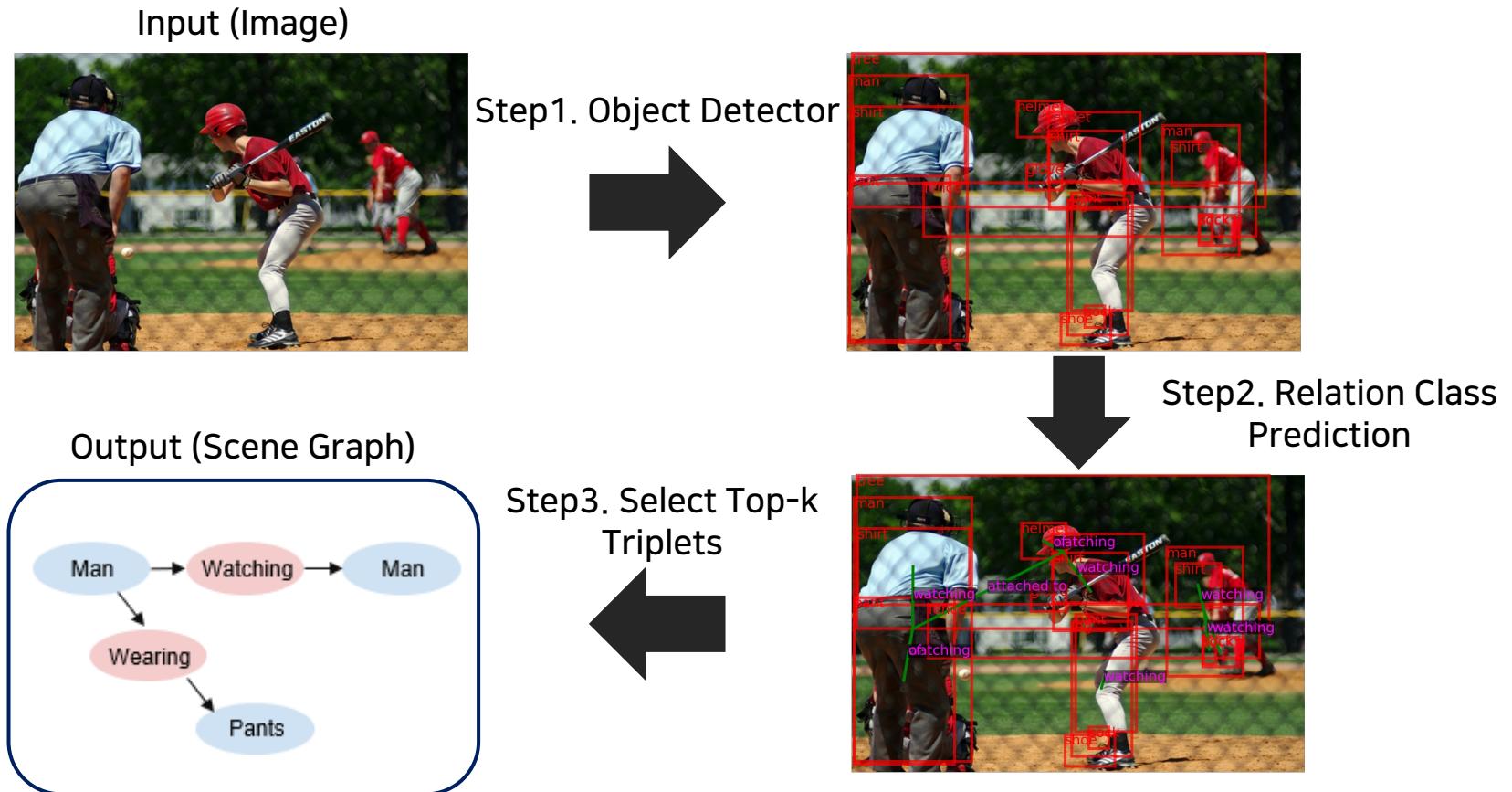
- Node feature = Concatenate (image embedding, non-image data)
- Learn consensus embedding
- Diagnosis

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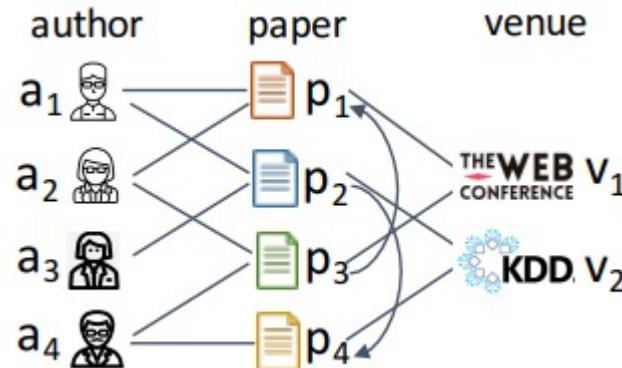
Scene graph generation (SGG)

- SGG aims to represent observable **knowledges** in an image in the form of a graph
 - The Knowledges include 1) object information and 2) their relation information
 - E.g., Object information: *man, horse, glasses, ...*
 - Relation information between objects: *feeding, wearing, ...*

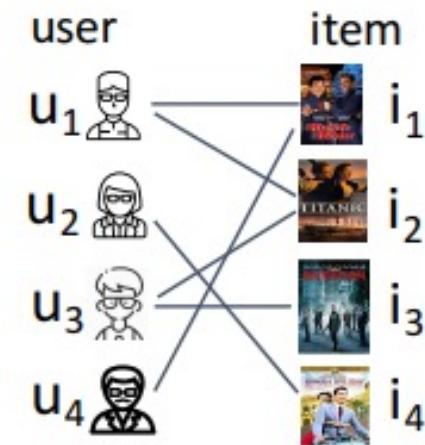


Heterogeneous graph

- Heterogeneous graph is a graph-structured data with more than one type of nodes or edges
 - By considering associations between multiple types of nodes or edges, many works demonstrate that **considering the heterogeneity of nodes/edges are helpful for learning the representations with the semantic information.**



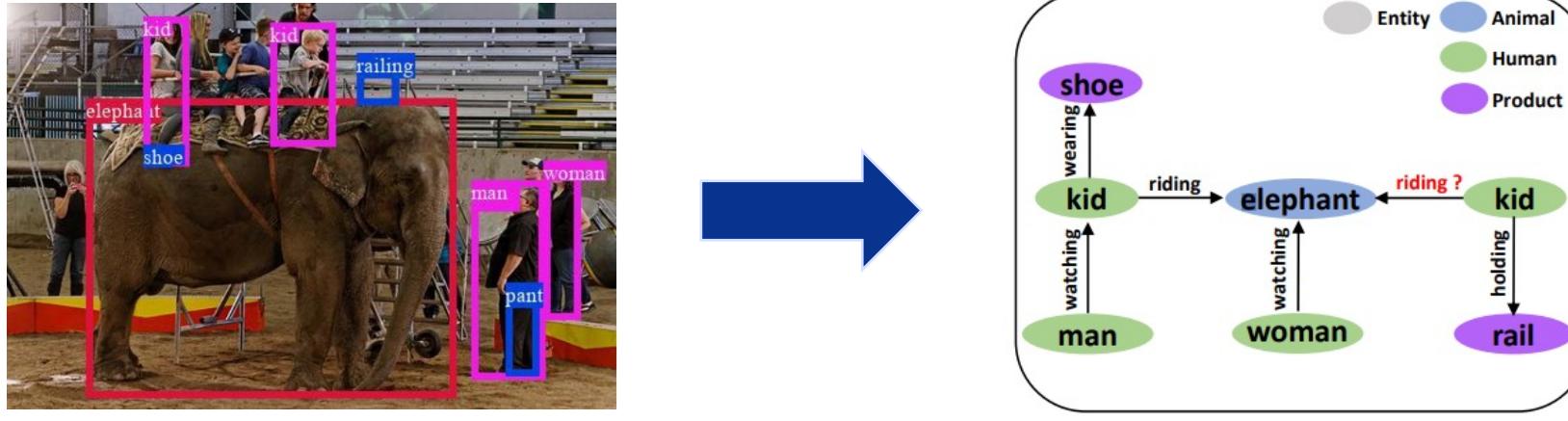
[Academic Graph]



[Review Graph]

Previous works

- In the literature of SGG, it's important to capture the context of neighborhood
 - Considering *<kid, holding, rail>* and *<woman, watching, elephant>* is helpful for predicting *<kid, riding, elephant>*
 - Compared with when kid and elephant are considered independently

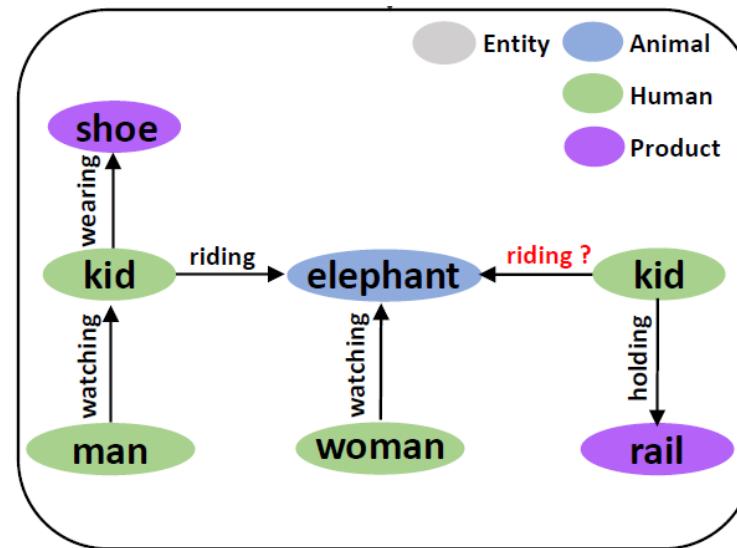


- Context-aware SGG employs RNN, GNN, …, Transformer to aggregate features of neighboring objects.

Limitations of previous works

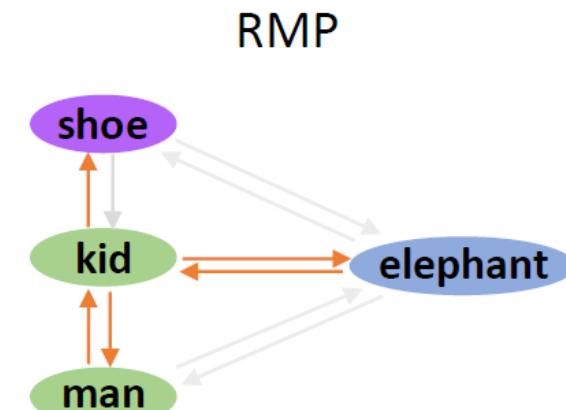
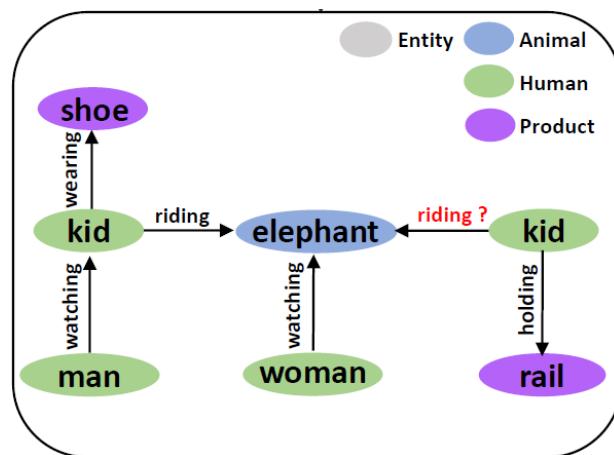
- Previous works consider the scene graph as **homogeneous graph**

- The assumption of homogeneity restricts the context-awareness of the visual relations between objects.
 - Since it **neglects the fact that predicates highly dependent on the objects where the predicates are associated.**
 - For example, when we consider **<kid, riding, elephant>**, we know the opposite triplet **<elephant, riding, kid>** is not likely to appear.
 - Because it is usually “Human” that rides “Animal”.



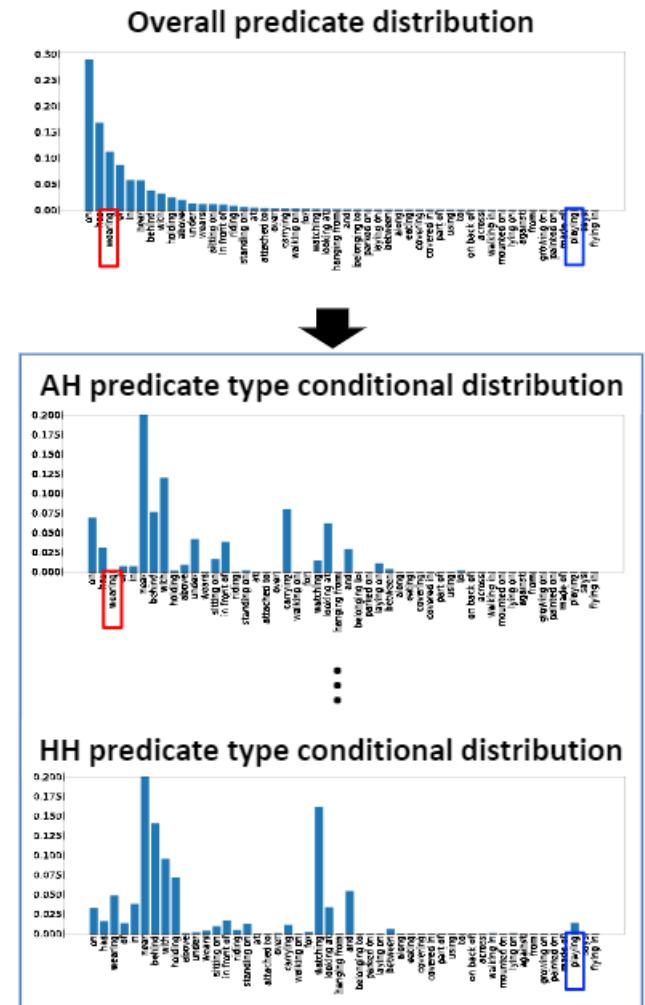
Our Goal

- We propose the Heterogeneous scene graph generation (HetSGG) framework
 - HetSGG generates a scene graph with relation-aware context
 - Consider both object types (e.g., Human, Animal, Product) & relation types (e.g., Human-Animal, Human-Human, ⋯).
 - We propose a novel message-passing called relation aware message-passing (RMP)
 - Naturally captures the semantic between “Human” and “Animal” to predict <*kid, riding, elephant*>



Relieving long-tailedness

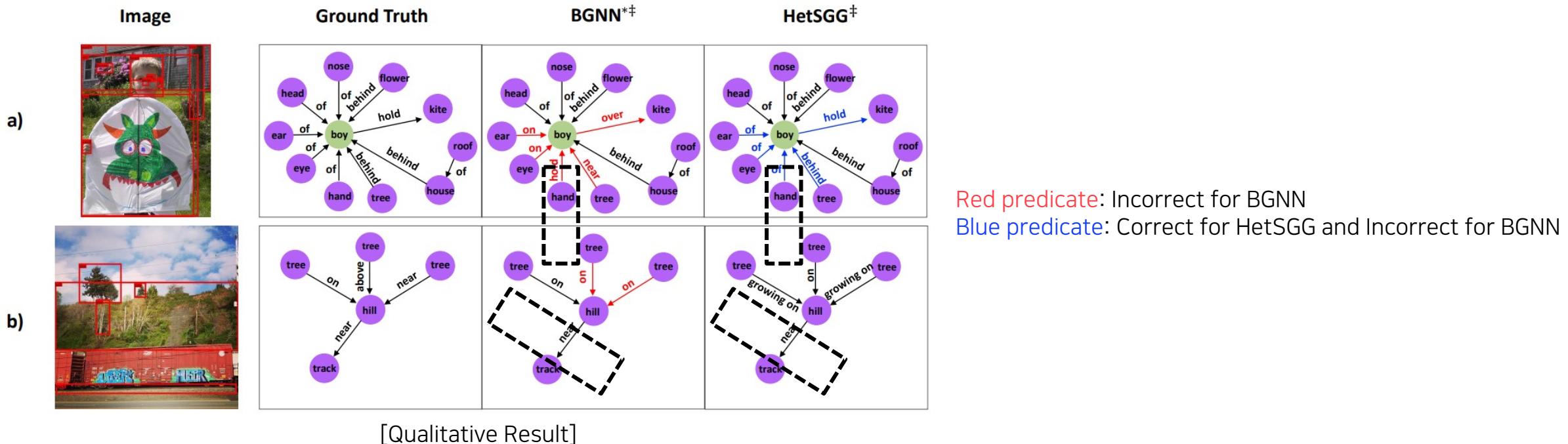
- Overall predicate distribution is **long-tailed**
 - Problem: Model primarily predicts the meaningless predicate (i.e., on, has)
- Observation of the reformulated distribution in condition of predicate types
 - Animal-Human(AH): **head** predicate (e.g., “**wearing**”) in overall distribution becomes **tail** predicate in AH distribution
 - Human-Human(HH): **tail** predicate (e.g., “**playing**”) makes up a small proportion of the overall distribution, but the proportion improves in HH distribution



We expect the long-tailed problem is naturally alleviated in the formulation of heterogeneous graph distinguishing the relation type

Experiment: Qualitative results

- a) BGNN predicts “hand hold boy”, but HetSGG predicts “hand of boy”
 - HetSGG predicts the correct predicate by filtering the non-sense semantic relation, such as “hand hold boy”
- b) BGNN predicts “tree on hill”, but HetSGG predicts the fine-grained predicate (i.e., growing on)
 - HetSGG alleviates the long-tailed predicate distribution, thus predicts the fine-grained predicate



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