



Graph Representation Learning (GRL)

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Networks/Graphs



Social Networks

Facebook

Twitter

...



Biological Networks

Brain

Protein-Protein

Drug-Target

...



Product Networks

User-Item

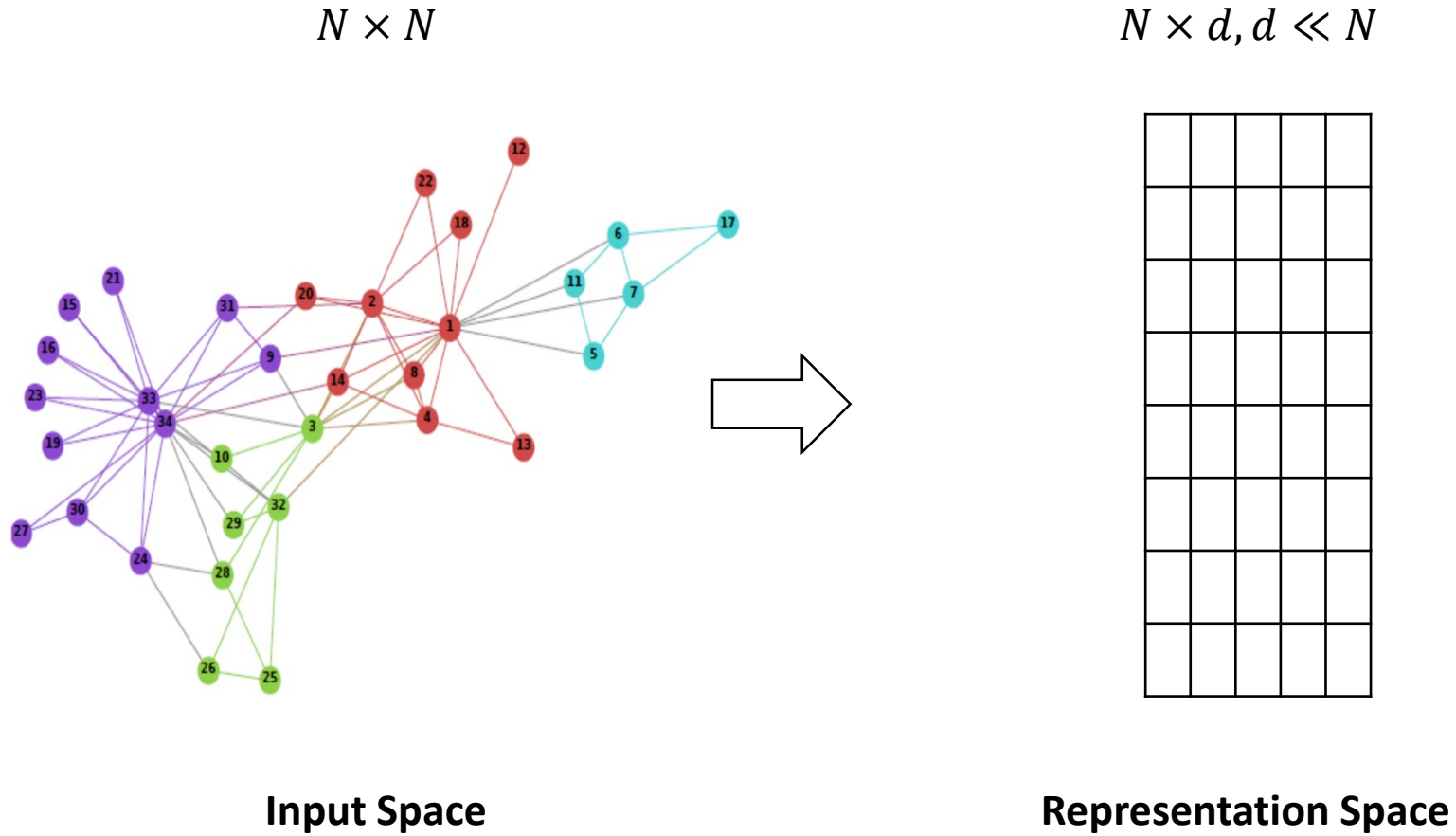
Product-Product

..

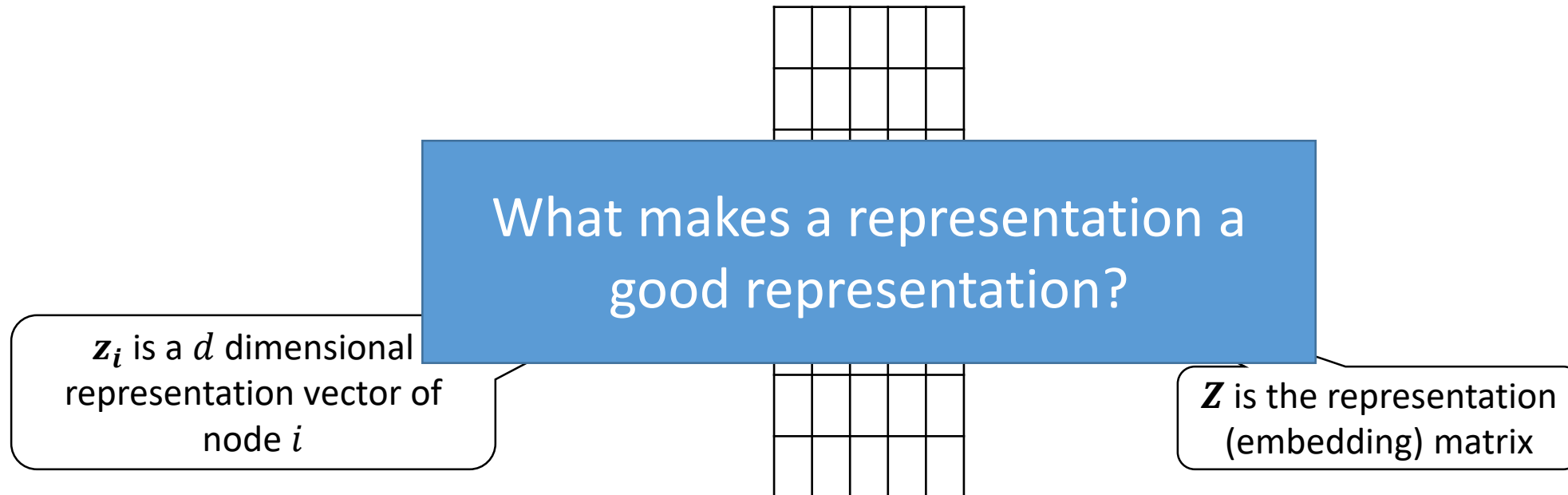
Basic Notations

- $G = (V, E)$
 - $N = |V|, M = |E|$
- $A \in \{0, 1\}^{N \times N}$ – Adjacency matrix notation
 - $A[i, j] = \begin{cases} 1, & (i, j) \in E \\ 0, & (i, j) \notin E \end{cases}$
- $X \in \mathbb{R}^{N \times D}$ – Node feature matrix
 - $X[i] = \mathbf{x}_i$ – Node i 's feature vector

What is GRL?

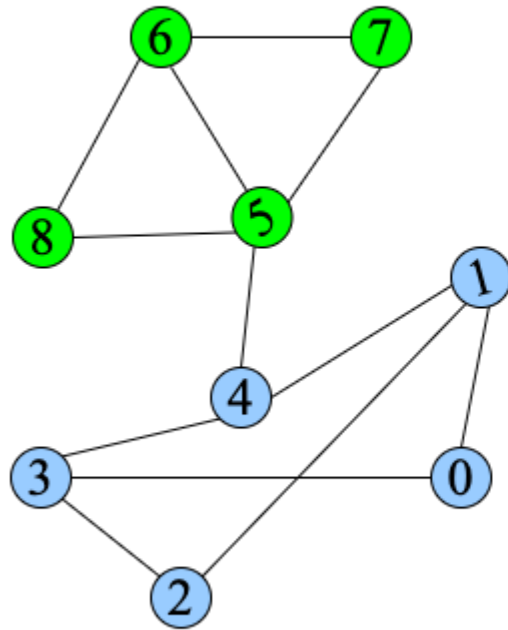


What is GRL?



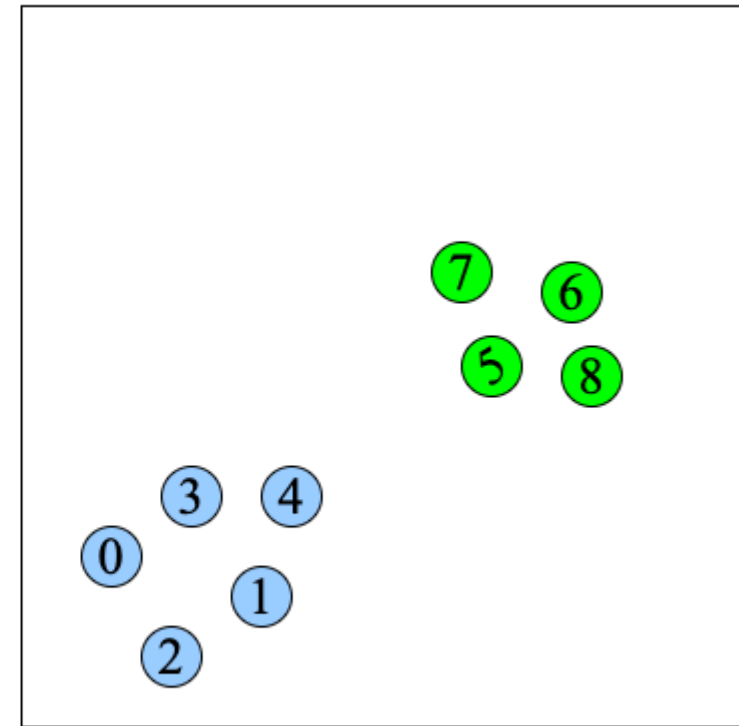
Quality of Representation

- Preserve a certain property of the graph



Input (graph) space

$$f: V \rightarrow \mathbb{R}^d$$



Embedding (representation) space

Quality of Embedding

- We model the property we want to preserve using a similarity function in the input space

$$sim_g: V \times V \rightarrow \mathbb{R}$$

- Adjacency Matrix as a similarity function

$$sim_g(i, j) = \begin{cases} 1, & \text{if } A[i, j] = 1 \\ 0, & \text{if } A[i, j] = 0 \end{cases}$$

- Common Neighborhood

- $sim_g(u, v) = |\mathcal{N}_u \cap \mathcal{N}_v|$
- $\mathcal{N}_i = \{j: (i, j) \in E\}$

- Rooted Page Rank

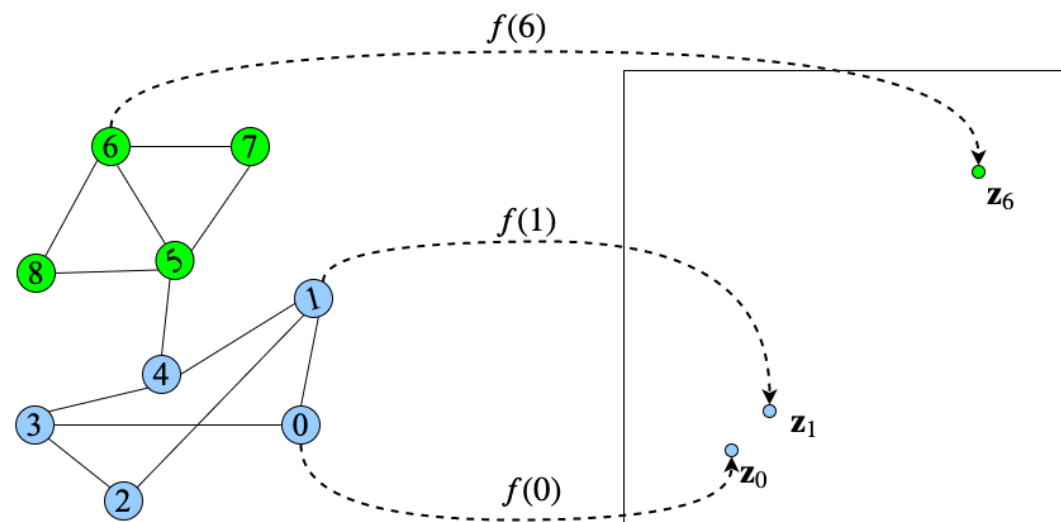
- $sim_g(u, v) = p$
- p is the probability of reaching node v via a random walk starting from node u

- Generally, we use $\mathbf{W} \in [0, 1]^{N \times N}$
 $\mathbf{W}[i, j] = sim_g(i, j)$

Learning

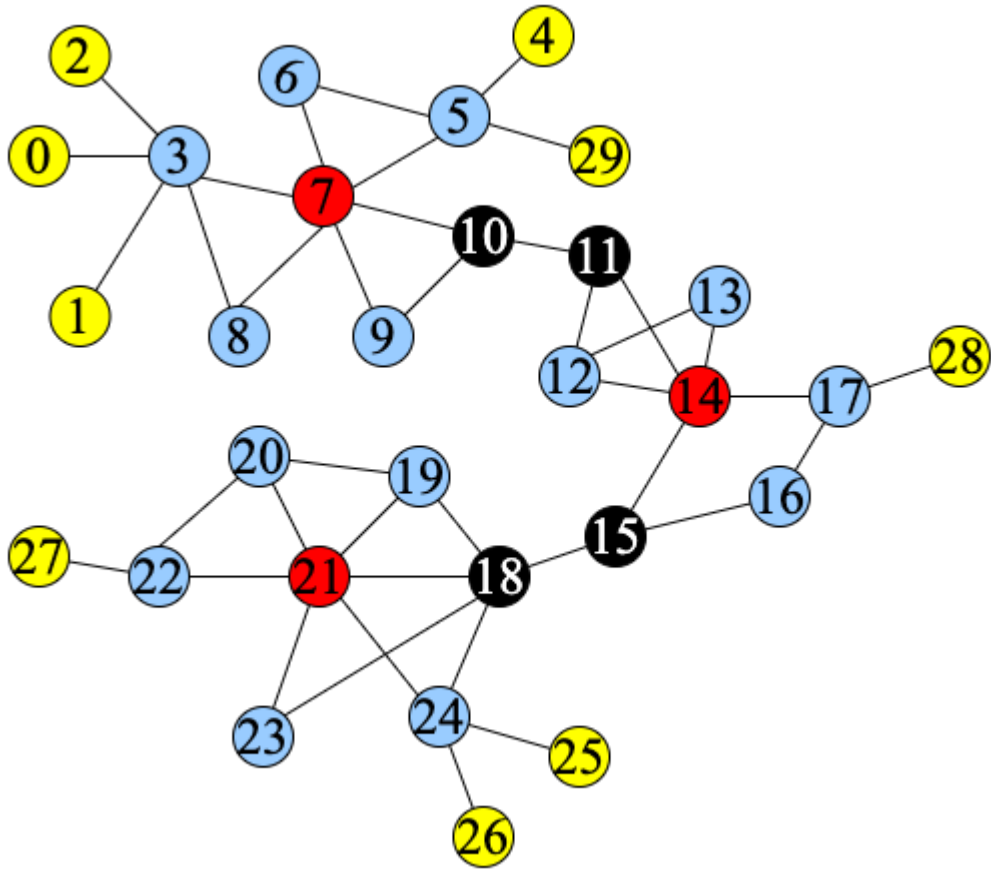
- Given a similarity function sim_g in the input space
- Learn a representation $\mathbf{z}_u = f(u) \in \mathbb{R}^d, u \in V$ that preserves sim_g
 - Similarity in the representation space, $sim_r: \mathbb{R}^d \times \mathbb{R}^d \rightarrow \mathbb{R}$
 - sim_r Should approximate sim_g

$$sim_r(f(u), f(v)) \approx sim_g(u, v)$$



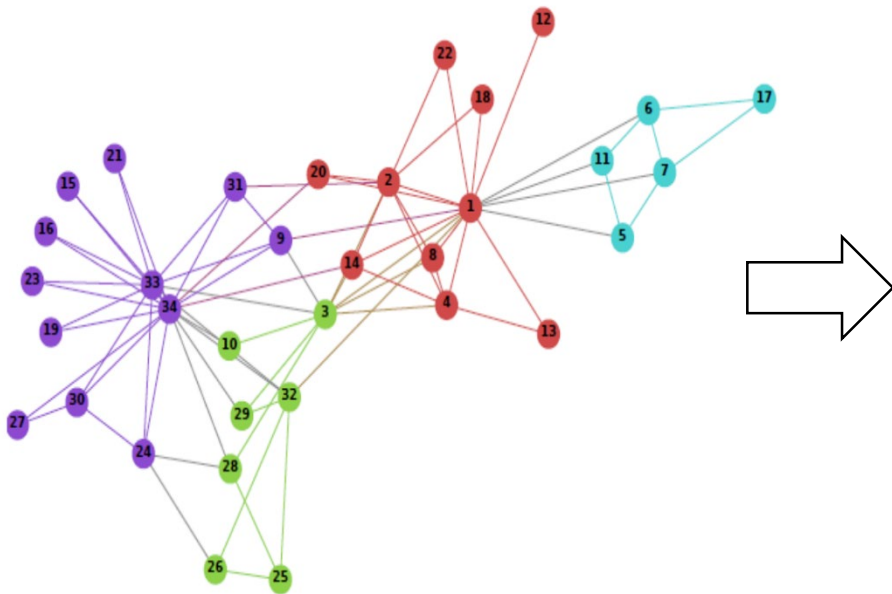
$$sim_r(f(0), f(1)) = sim_r(\mathbf{z}_0, \mathbf{z}_1) \approx sim_g(0, 1)$$

Structural similarity



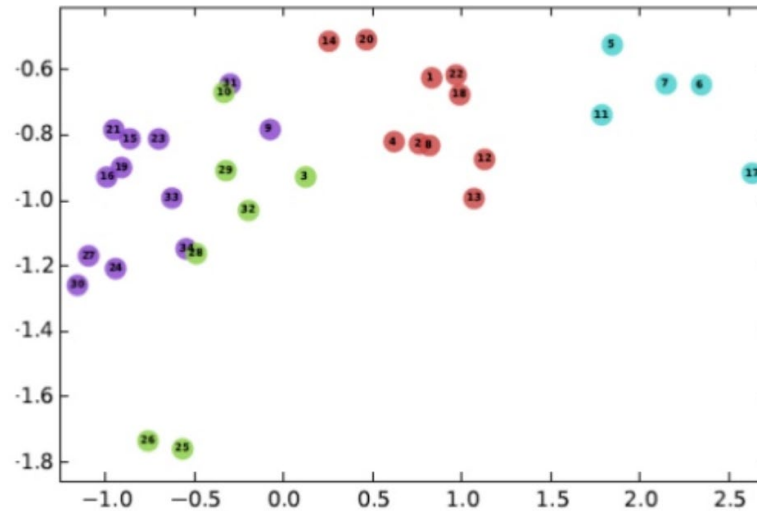
Why GRL?

$$N \times N$$



Input Space

$$N \times d, d \ll N$$



Representation Space

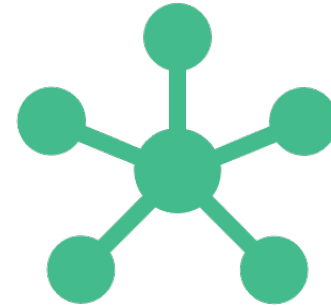
Network Analysis

- Link Prediction
- Node classification
- Clustering
- Recommendation
- ...

Learning Algorithms



Random Walk Based



Message Passing Neural Networks

Random Walk (RW) Based



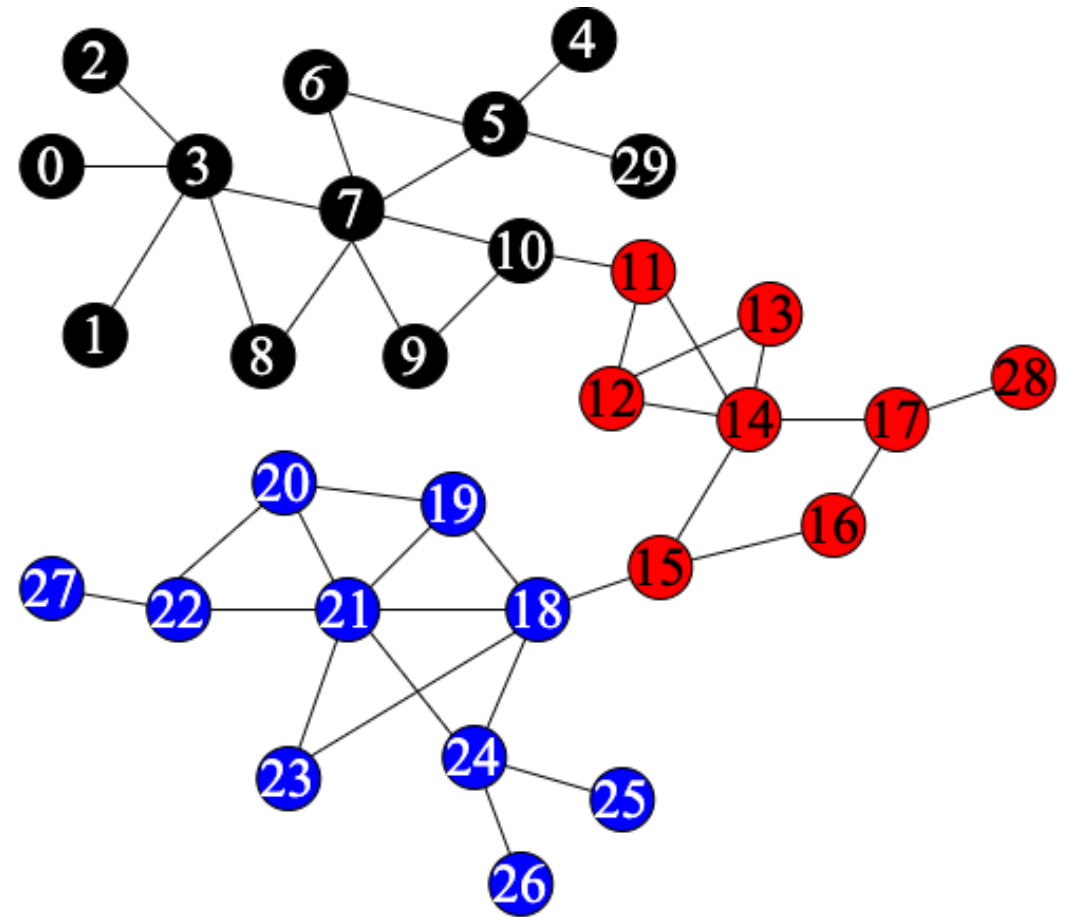
WALK SAMPLING



LEARNING

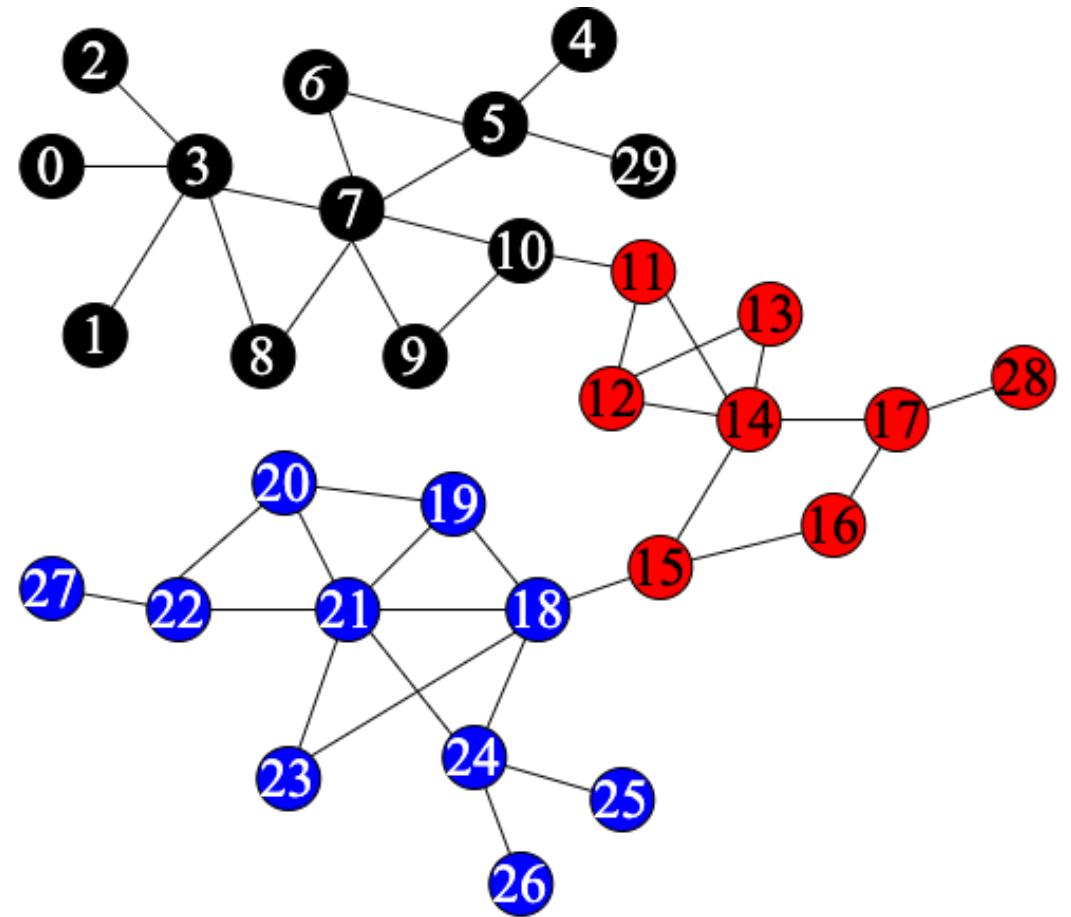
RW: Property

- Preserve Homophily: Similar nodes appear in the same neighborhood



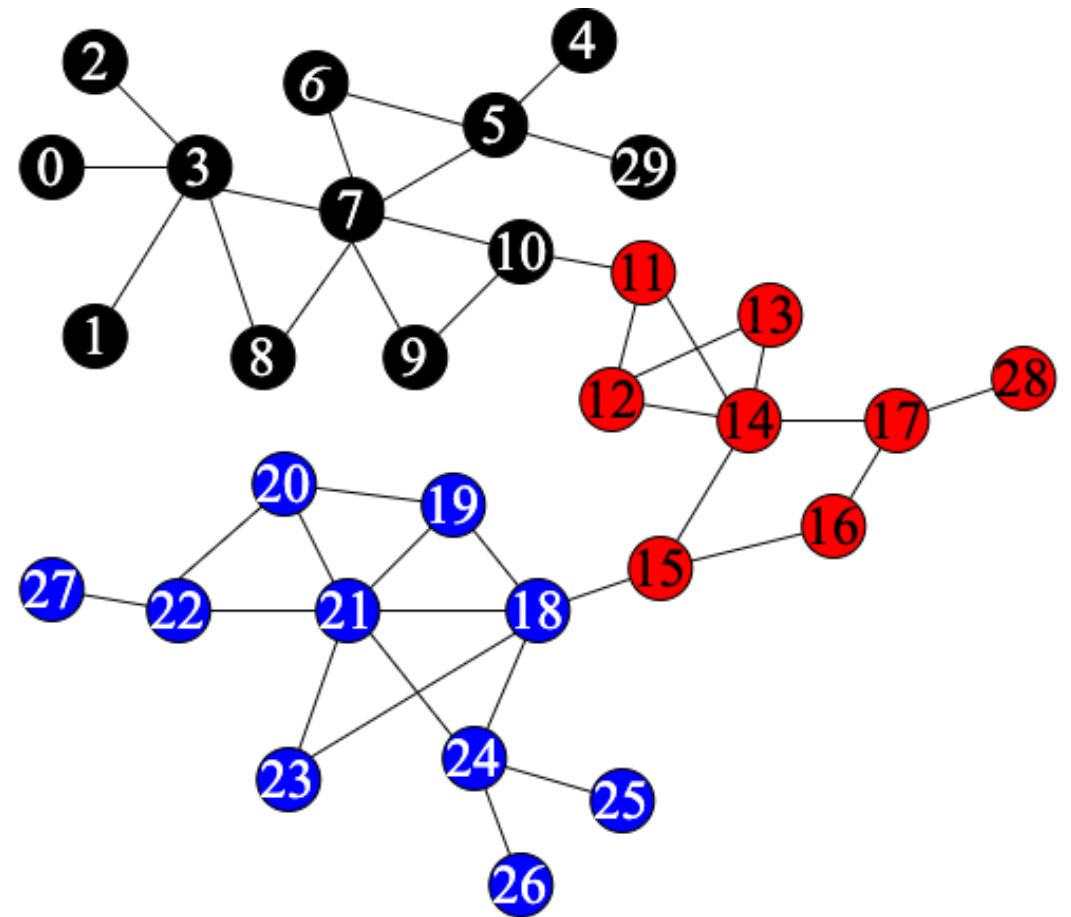
Challenge

- Graphs are arbitrary data structures
- GRL directly on top of graphs is difficult
- We need a different data structure that capture the homophily property (E.g., sequences)
 - Truncated random walks



RW Sampling

- Sample walks starting from node 21
 - 21, 24, 18, 23, 18
 - 21, 18, 19, 18, 15
 - 21, 19, 20, 22, 21
 - 21, 23, 18, 24, 21
- Sample walks starting from node 15
 - 15, 14, 12, 13, 12
 - 15, 14, 11, 14, 12
 - 15, 18, 15, 16, 17
 - 15, 14, 17, 28, 17
- ...



Learning: Algorithm

- SkipGram: Used for word representation
- Key idea: Similar words frequently appear together in similar context
- E.g.:
 - Stockholm vs. Sweden
 - Vaccine vs. AstraZeneca
 - Darwin vs. Evolution
 - Arthur Conan Doyle vs. Sherlock Holmes
- Learning:
 - Given a target word, w_t
 - Learn its embedding, \mathbf{z}_t by predicting its context words $w_c \in ctx(w_t)$

The, capital, of, Sweden, is, Stockholm

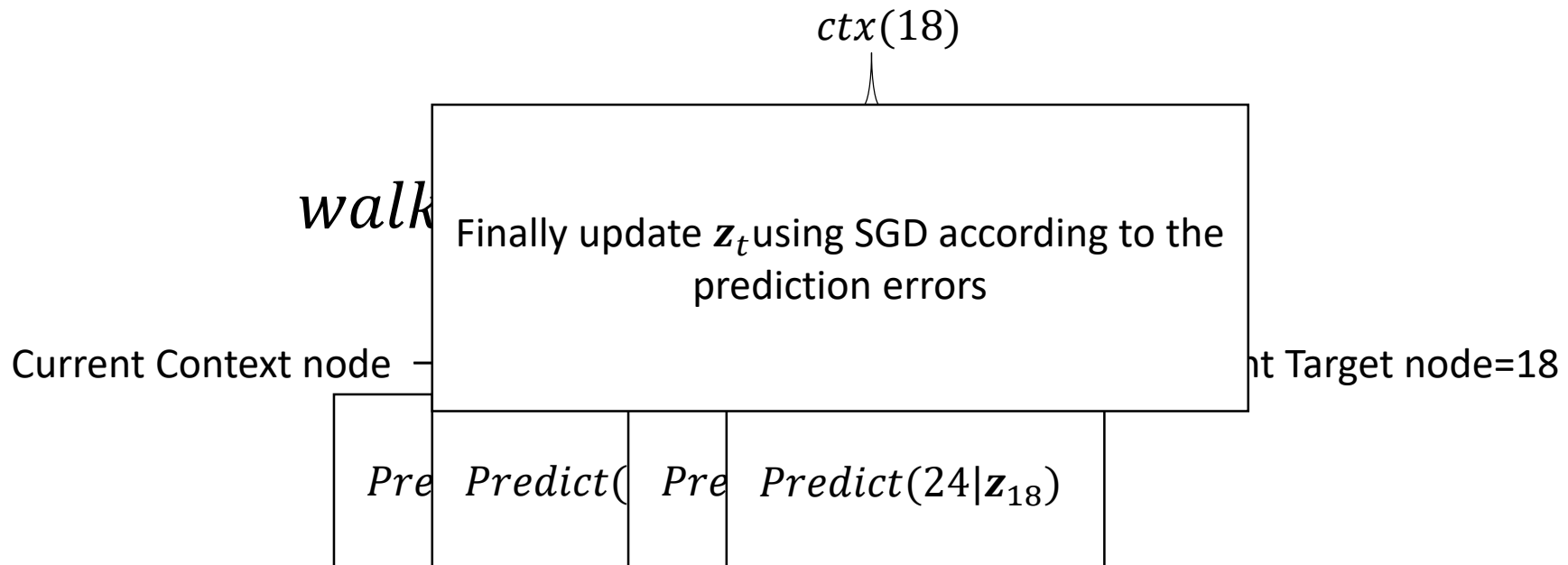
$$w_t = \textit{Sweden}$$

$$ctx(w_t) = \{\textit{capital, stockholm}\}$$

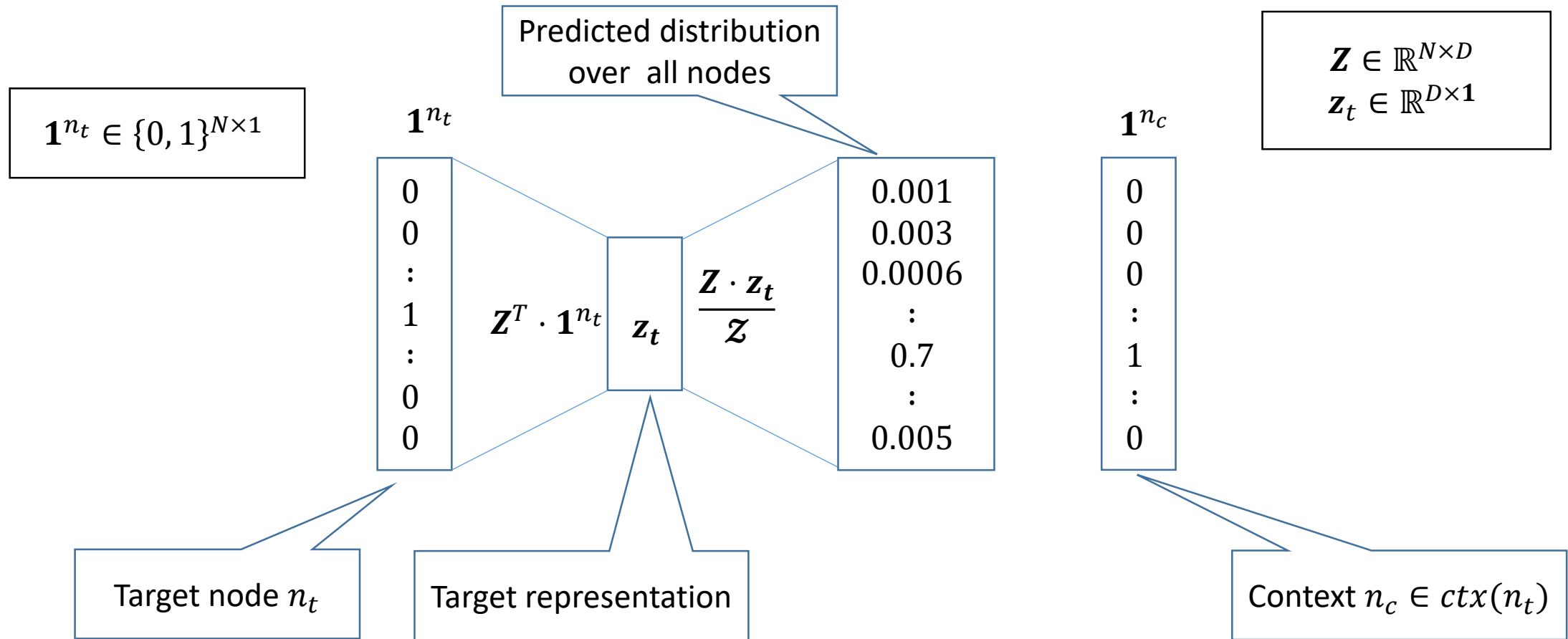
$$\max_{w_c \in ctx(w_t)} P(w_c | \mathbf{z}_t)$$

SkipGram for GRL

- Given a target node n_t from a random walk sequence
- Learn \mathbf{z}_t by predicting its context nodes $n_c \in ctx(n_t)$

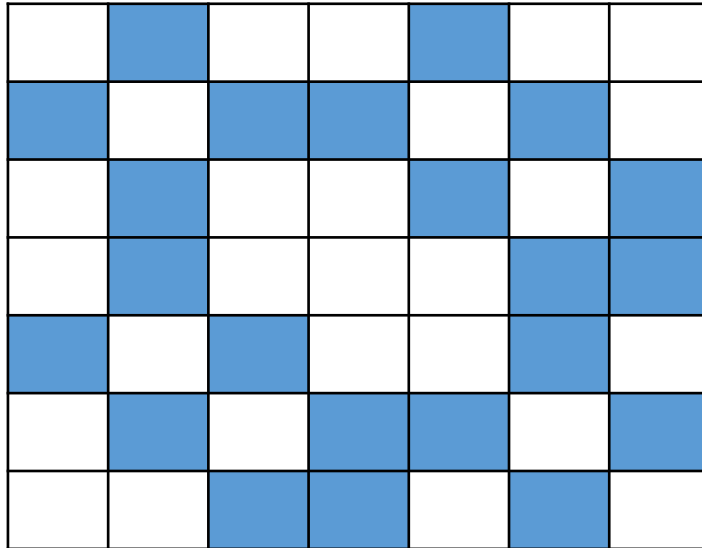


SkipGram Architecture

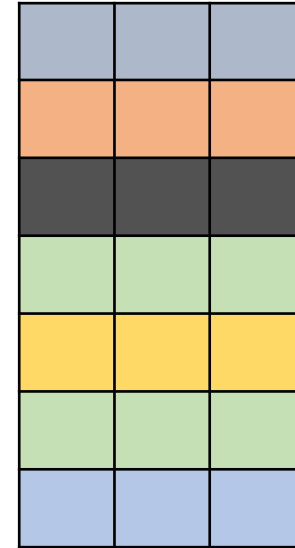


Message Passing Neural Networks (MPNN)

- Now, we consider nodes with features



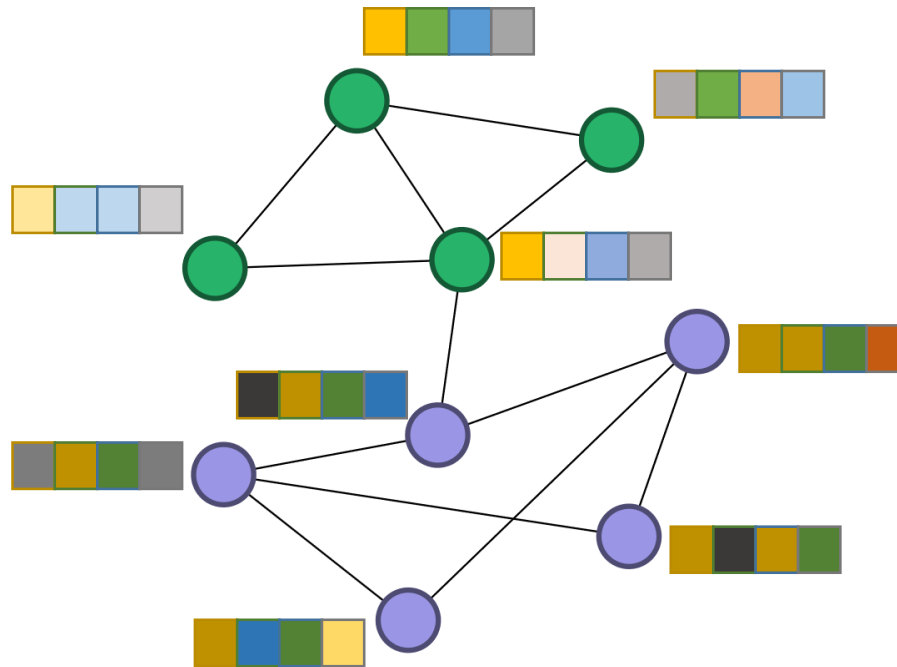
A



X

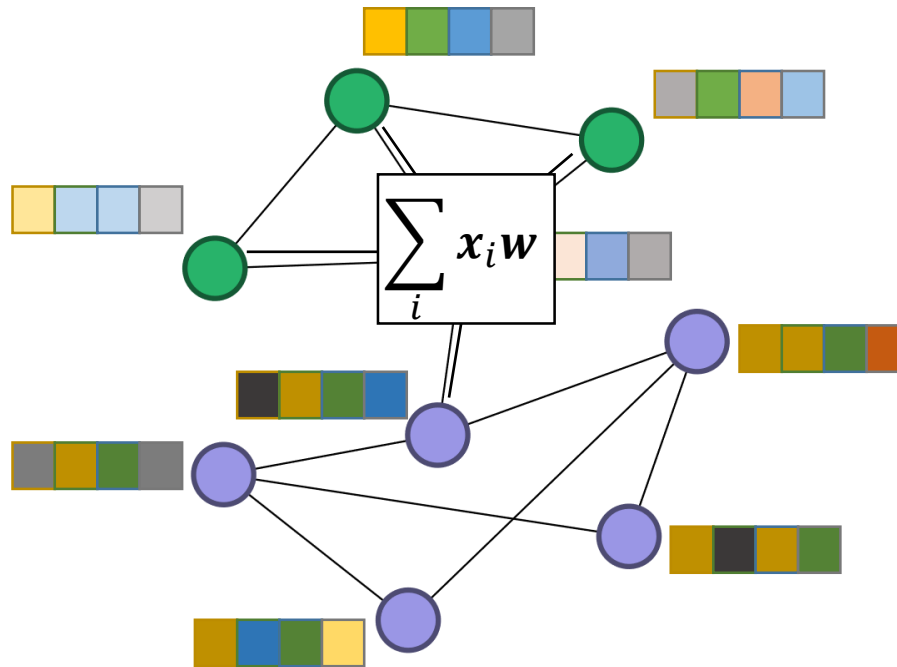
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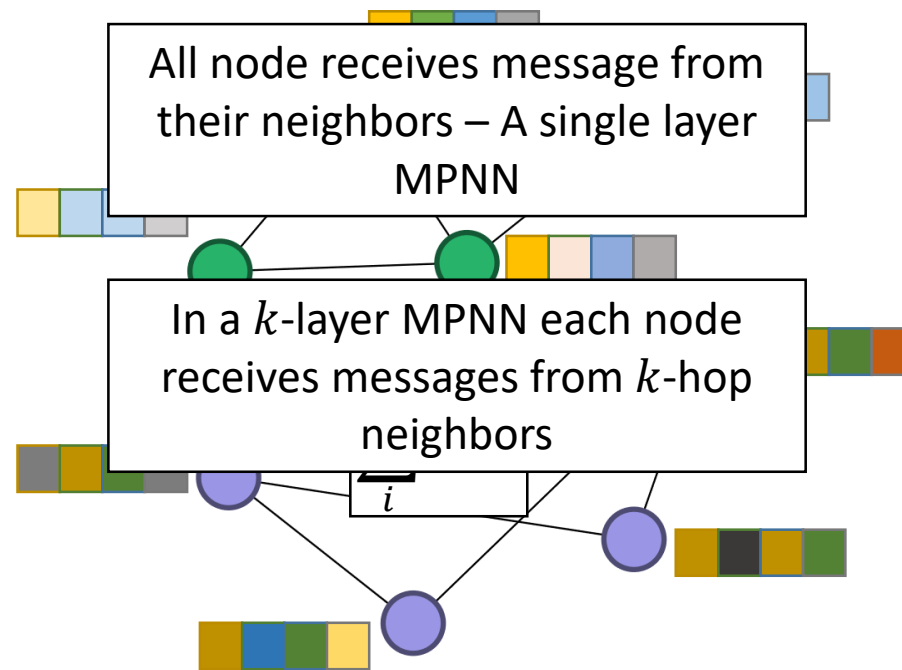
MPNN

- Key Idea: Each node sends and receives messages

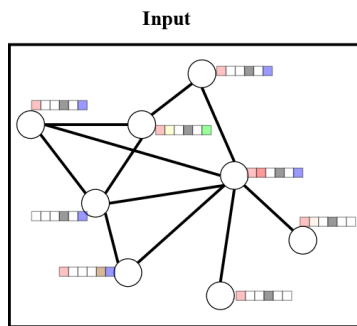


MPNN

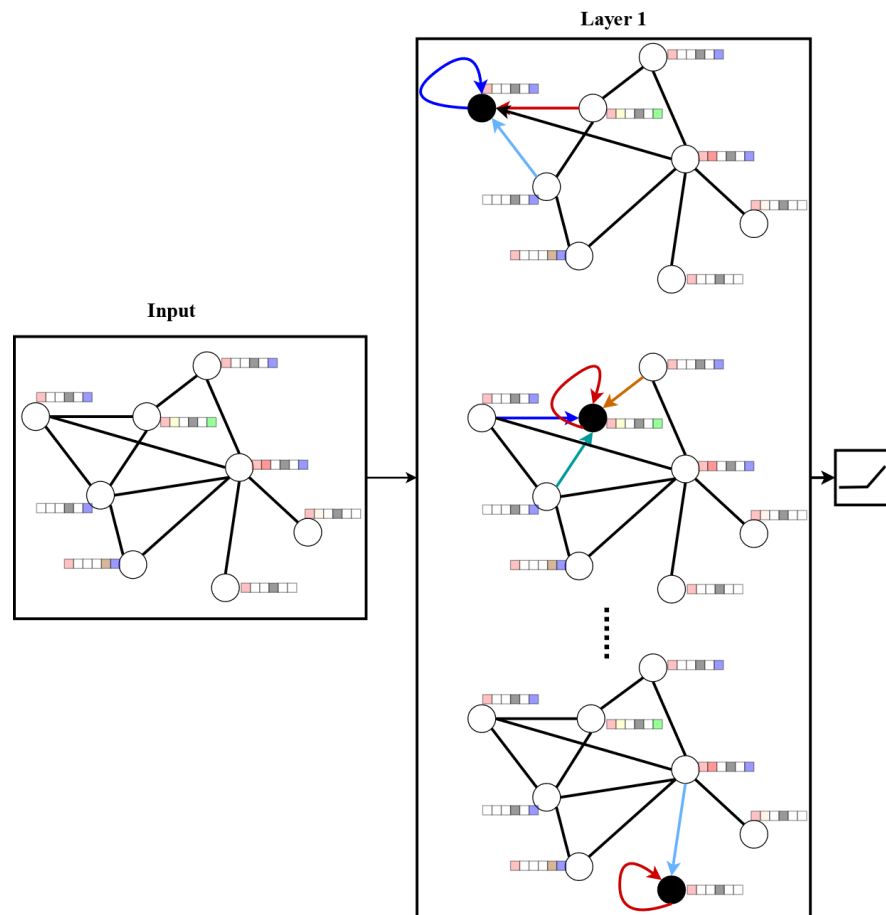
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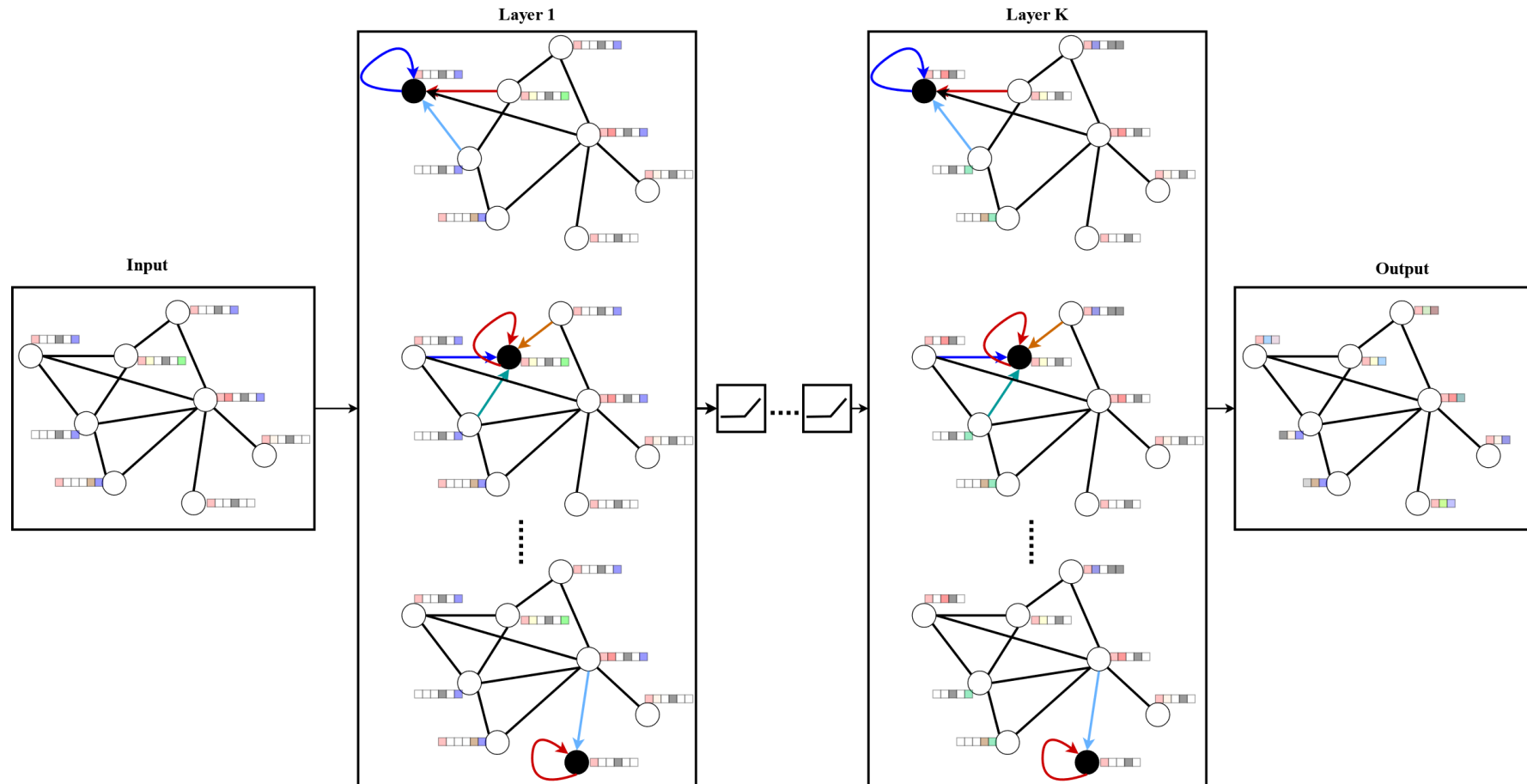
GCN



GCN



GCN



Applications of GRL

- Social Networks
 - Friendship recommendation
- Recommendation Systems
 - Content (e.g., book, movie, ...) recommendation
- Biomedical Systems
 - Drug discovery
 - Predicting functions of proteins
 - Predicting molecular properties
- ...



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