

Applications of Graph Neural Networks

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(1) Graph Convolutional Networks (GCN)

$$\mathbf{h}_{v}^{(l)} = \sigma \left(\mathbf{W}^{(l)} \sum_{u \in N(v)} \frac{\mathbf{h}_{u}^{(l-1)}}{|N(v)|} \right)$$

How to write this as Message + Aggregation?





(1) Graph Convolutional Networks (GCN)

$$\mathbf{h}_{v}^{(l)} = \sigma \left(\sum_{u \in N(v)} \mathbf{W}^{(l)} \frac{\mathbf{h}_{u}^{(l-1)}}{|N(v)|} \right)$$

Message:

• Each Neighbor:
$$\mathbf{m}_{u}^{(l)} = \frac{1}{|N(v)|} \mathbf{W}^{(l)} \mathbf{h}_{u}^{(l-1)}$$

Normalized by node degree (In the GCN paper they use a slightly different normalization)

Aggregation:

Sum over messages from neighbors, then apply activation

•
$$\mathbf{h}_{v}^{(l)} = \sigma\left(\operatorname{Sum}\left(\left\{\mathbf{m}_{u}^{(l)}, u \in N(v)\right\}\right)\right)$$



(2) GraphSAGE

$$\mathbf{h}_{v}^{(l)} = \sigma \left(\mathbf{W}^{(l)} \cdot \text{CONCAT} \left(\mathbf{h}_{v}^{(l-1)}, \text{AGG} \left(\left\{ \mathbf{h}_{u}^{(l-1)}, \forall u \in N(v) \right\} \right) \right) \right)$$

- How to write this as Message + Aggregation?
 - Message is computed within the AGG(·)
 - Two-stage aggregation

 $\mathbf{h}_{N(v)}^{(l)} \leftarrow \mathrm{AGG}\left(\left\{\mathbf{h}_{u}^{(l-1)}, \forall u \in N(v)\right\}\right)$

• Stage 2: Further aggregate over the node itself

$$\mathbf{h}_{v}^{(l)} \leftarrow \sigma\left(\mathbf{W}^{(l)} \cdot \text{CONCAT}(\mathbf{h}_{v}^{(l-1)}, \mathbf{h}_{N(v)}^{(l)})\right)$$



Mean: Take a weighted average of neighbors



 Pool: Transform neighbor vectors and apply symmetric vector function Mean(·) or Max(·)

 $AGG = Mean(\{MLP(\mathbf{h}_{u}^{(l-1)}), \forall u \in N(v)\})$

Aggregation Message computation

LSTM: Apply LSTM to reshuffled of neighbors

AGG = LSTM($[\mathbf{h}_{u}^{(l-1)}, \forall u \in \pi(N(v))]$) Aggregation



(3) Graph Attention Networks

$$\mathbf{h}_{v}^{(l)} = \sigma(\sum_{u \in N(v)} \alpha_{vu} \mathbf{W}^{(l)} \mathbf{h}_{u}^{(l-1)})$$

- In GCN / GraphSAGE
 - $\alpha_{vu} = \frac{1}{|N(v)|}$ is the weighting factor (importance) of node *u*'s message to node *v*
 - $\Rightarrow \alpha_{vu}$ is defined **explicitly** based on the structural properties of the graph (node degree)
 - ⇒ All neighbors $u \in N(v)$ are equally important to node v



(3) Graph Attention Networks

$$\mathbf{h}_{v}^{(l)} = \sigma(\sum_{u \in N(v)} \alpha_{vu} \mathbf{W}^{(l)} \mathbf{h}_{u}^{(l-1)})$$

ALLEIILION WEIGHLS

Not all node's neighbors are equally important

- Attention is inspired by cognitive attention.
- The **attention** α_{vu} focuses on the important parts of the input data and fades out the rest.
 - Idea: the NN should devote more computing power on that small but important part of the data.
 - Which part of the data is more important depends on the context and is learned through training.



Application of Graph Neural Networks

Node Classification



□Some labels of nodes are given

□ Predict the labels of Unlabelled nodes.





Application of Graph Neural Networks

Link Prediction





With existing edges (links) between nodes, predict the existence of possible links.

Link prediction can be used to explore potential interaction between proteins.



Applications of GNNs

- Structured Entity Analysis (Chemical, Biomedical)
- Subgraph Isomorphism (Database)
- Fraud Detection (e-Commerce)



Structured Entity Analysis



Structured Entity Analysis

- Many real-life structured entities can be modelled as graphs, such as chemical molecules and proteins.
- Graph neural networks can be used to analyse these structured entities.





Structured Entity Analysis

- Toxicity of chemical molecules.
- Property prediction for molecules.
- Interactions between molecules.



Interaction between molecules

- Prediction of reaction between two molecules
- Prediction of the side effect of drug pairs.





Immediate solutions

- The immediate way to investigate the interactions between two entities (molecules or drugs) is to conduct experiments in laboratories and clinics.
- However, it is time consuming and labour intensive. It is impractical to test all entity pairs.
- Therefore, computational approaches are proposed to predict structured entity interaction effectively and efficiently.



Naïve computational solution

- Model the molecules as graphs
- Directly apply the graph neural networks for the predictions.



 However, such solution ignores the interaction relation between molecules.



Other computational methods

Simplified molecular-input line-entry system (SMILES)

- Using strings to represent molecules.
- Natural Language Processing (NLP) models are applied to capture the molecule information and produce the representations for molecules
- Downstream applications are based on the representations.
- However, these models cannot capture the structural information of the molecules.



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Goals

- Capture the structural information of the molecule graphs.
- Preserve the interaction relationship between the structured entities (molecules).
- Identify the important substructures within the molecules that are key to the prediction of interactions.



Graph of Graphs

- Local graph: the molecule graph representing the chemical structure.
- (nodes: atoms, edges: chemical bonds between the atoms.)
- Global graph: the graph of interactions between the chemicals.
- (nodes: chemical molecules, edges: interaction relations between the molecules.)





Target of the model

- With the input graph, the target of the model is to conduct the link prediction on the global graph.
- The graph neural network applied on the *local graph* should be able to capture the important substructure (functional groups) in the molecules.







Multi-Resolution Graph Neural Network

- Use the concatenation of the output after different layer of GNNs to keep multi-scale (multi-hop) substructure information.
- Each layer of graph neural network captures one-hop neighbour relationship.

Local graph neural network:

The neural network used to learn representations for chemical molecules. We use *I*-layer graph convolutional network (GCN) as the local graph neural network:

$$M_{(l+1)} = GCN_l(A, M_l)$$
$$GCN_l(A, M_l) = \sigma \left(D^{-\frac{1}{2}} \tilde{A} D^{-\frac{1}{2}} M_l W_l \right)$$

Graph pooling

Graph pooling is a computational strategy to reduce the number of graph nodes; in this way, one has a unified graph-level rather than node-level representation for graph-structured data while the size and topology of an individual graph are changing.

Graph pooling examples

(1) Global mean pooling

 $\mathbf{y}_0 = \text{Mean}(\{\mathbf{h}_v \in \mathbb{R}^d, \forall v \in G\})$

• (2) Global max pooling

$$\mathbf{y}_0 = \operatorname{Max}(\{\mathbf{h}_v \in \mathbb{R}^d, \forall v \in G\})$$

(3) Global sum pooling

$$\mathbf{y}_0 = \mathbf{Sum}(\{\mathbf{h}_v \in \mathbb{R}^d, \forall v \in G\})$$

Issue: Global pooling over a (large) graph will lose information

Graph attention pooling

 The attention-based pooling method to select the most representative substructure to represent the molecule graph:

$$s_{l} = \sigma \left(\underbrace{D^{-\frac{1}{2}} \widetilde{AD}^{-\frac{1}{2}} M_{l} W_{att}^{l}}_{idx = top(s, [\gamma n]),} \right)$$
$$s_{mask} = s_{idx}, M_{sel} = M \odot s_{mask}$$

Graph attention pooling

attention poolinç

$$\begin{split} s_{l} &= \sigma \left(\widetilde{D^{-\frac{1}{2}} \widetilde{A} D^{-\frac{1}{2}} M_{l} W_{att}^{l}} \right) \\ & i dx = top(s, \lceil \gamma n \rceil), \\ s_{mask} &= s_{idx}, M_{sel} = M \odot s_{mask} \end{split}$$

 $s_{l} = \sigma \begin{pmatrix} l \\ idx = \\ s_{mask} = s_{id} \end{pmatrix}$

- *M_{sel}* is the representation matrix for the selected atoms in the molecule graph
- After selection, the combination of mean pooling and sum pooling is used to produce the representation for the molecule graph, which is also the input for interaction graph neural network.

is the representation nolecule graph

r selection, the com ing is used to produ cule graph, which i h neural network.

Graph attention network:

 With the learned molecule graph representations, the embedding is updated based on the interaction network. The multihead attention mechanism is utilized:

$$\begin{aligned} x_{G_{i}}^{l+1} &= \left\| \begin{matrix} K \\ \kappa = 1 \end{matrix} \sigma \left(\sum_{j \in \eta_{G_{i}}} \alpha_{ij}^{\kappa} W_{\kappa}^{l} x_{G_{j}}^{l} \right) \\ exp\left(\text{LeakeyRelu}\left(a \left[W x_{G_{i}} \right] || \left[W x_{G_{j}} \right] \right) \right) \\ \overline{\sum_{n \in \eta_{G_{i}}} exp\left(\text{LeakeyRelu}\left(a \left[W x_{G_{i}} \right] || \left[W x_{G_{n}} \right] \right) \right)} \end{aligned} \right. \end{aligned}$$

Training objectives

The predicted interaction probability: $p_{ij} = \sigma \left(x_{G_i}^T \cdot x_{G_j} \right)$

Cross-entropy loss function:

$$\mathcal{L}_{CCJ} = \sum_{\left(G_{i}, G_{j}\right) \in G_{CCI}} - \log(p_{ij}) - E_{m \sim P_{j}} \log(1 - p_{im})$$

Datasets:

- CCI tasks: The CCI dataset assigns a score from 0 to 999 to describe the interaction probability where a higher score indicates higher interaction probability. According to threshold score, we get two datasets with chemical interaction probability score over 900 and 950: CCI900 and CCI950.
- DDI tasks. For the drug-drug interaction prediction problem, DDI dataset and the side effect dataset SE are used. A vector representation (attribute) is assigned to each side effect type produced by pre-trained BERT model

Experiment results:

| | CCI900 | | CCI950 | |
|--------------|--------|-------|--------|-------|
| | AUC | AP | AUC | AP |
| DeepCCI | 0.925 | 0.918 | 0.957 | 0.957 |
| DeepDDI | 0.891 | 0.886 | 0.916 | 0.915 |
| MR-GNN | 0.927 | 0.921 | 0.934 | 0.924 |
| MLRDA | 0.922 | 0.907 | 0.959 | 0.948 |
| SEAL | 0.894 | 0.886 | 0.941 | 0.937 |
| GoGNN | 0.937 | 0.932 | 0.963 | 0.962 |
| GoGNN-M | 0.914 | 0.909 | 0.938 | 0.931 |
| GoGNN-I | 0.921 | 0.898 | 0.929 | 0.912 |
| GoGNN-noPool | 0.931 | 0.930 | 0.958 | 0.954 |
| GoGNN-noAttn | 0.909 | 0.905 | 0.956 | 0.948 |

Table 1: Result of chemical-chemical interaction prediction task.

| | DDI | | SE | |
|--------------|-------|-------|-------------|-------------|
| | AUC | AP | AUC | AP |
| DeepCCI | 0.862 | 0.856 | 0.819 | 0.806 |
| DeepDDI | 0.915 | 0.912 | 0.827 | 0.809 |
| MR-GNN | 0.932 | 0.922 | 0.769^{*} | 0.752^{*} |
| MLRDA | 0.931 | 0.926 | 0.847^{*} | 0.825^{*} |
| Decagon | - | - | 0.872 | 0.832 |
| SEAL | 0.925 | 0.921 | N/A | N/A |
| GoGNN | 0.943 | 0.933 | 0.930 | 0.927 |
| GoGNN-M | 0.905 | 0.902 | 0.862 | 0.817 |
| GoGNN-I | 0.922 | 0.917 | 0.860 | 0.834 |
| GoGNN-noPool | 0.900 | 0.891 | 0.912 | 0.909 |
| GoGNN-noAttn | 0.925 | 0.921 | 0.897 | 0.883 |

* indicates that the result is the output of the baselines after two weeks' training.

 DDI dataset has no protein data which is required by Decagon

Table 2: Result of drug-drug interaction prediction task.

Case study:

Red: the substructures that are responsible of the interactions.

Our recent work

- "Denoising Variational Graph of Graphs Auto-Encoder for Predicting Structured Entity Interactions" -- TKDE 2023

Subgraph Isomorphism

Subgraph Counting

Subgraph Isomorphism

- Query graph $q = (V, E, f_l)$
- Data graph $G = (V', E', f_l)$
- Subgraph Isomorphism: injective function $f_{iso}: V \rightarrow V'$:
 - $\forall u \in V, f_l(u) = f_l(f_{iso}(u))$
 - $\forall e(u, u') \in E, e(f_{iso}(u), f_{iso}(u')) \in E'$
- Determining the existence of subgraph isomorphism is NP-complete.

<u>Graph isomorphism (</u> a more complicated example)

| Graph G | Graph H | An isomorphism between G and H |
|-----------------------|---------|--|
| a b b i j | | f(a) = 1 f(b) = 6 f(c) = 8 f(d) = 3 f(g) = 5 f(h) = 2 f(i) = 4 f(j) = 7 |

https://en.wikipedia.org/wiki/Graph_isomorphism

Subgraph Counting

Subgraph Counting: Given a query graph q and a data graph G, the problem is to count the number of subgraphs in the data graph that match the query graph by subgraph <u>isomorphism</u>.

Subgraph isomorphisms

1.
$$(u_1, u_2, u_3, u_4) \rightarrow (v_1, v_4, v_5, v_{10})$$

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2. $(u_1, u_2, u_3, u_4) \rightarrow (v_1, v_4, v_5, v_{11})$



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3.
$$(u_1, u_2, u_3, u_4) \to (v_1, v_4, v_6, v_{11})$$



Why Subgraph Counting?

Applications

Analysis on Social Networks:

• Find co-authorships, community detection.

Analysis on Biological Networks:

- On brain, regulation, protein and molecule graphs.
- Summarize the structural patterns for the biological graphs.

Query Optimization for Subgraph Matching Queries:

• Cardinality estimation for multi-way join.







Algorithmic Methods:

Enumeration-based methods:

• Computational complexity.

Sampling-based methods:

Sampling failure.

Summary-based methods:

• Independent assumption.

Learning-based Methods:

Neural Subgraph Isomorphism Counting



A Learned Sketch for Subgraph Counting





Neural Subgraph Isomorphism Counting:

- The query graph and data graph are input into the graph neural networks for representation learning.
- The learned representations are input into the RNN-based network named DIAMNet to predict the subgraph counts.





Neural Subgraph Isomorphism Counting:

- The data graph is usually large-scale.
- The model will face the efficiency and scalability issue.
- Since the data graph contains more information, it is hard to distinguish the counting results of different query graphs when the data graph is large. The representation of data graph will dominate the computation in this case.





A Learned Sketch for Subgraph Counting:

- The initial features of query nodes are computed based on the labels and structure of data graph by the graph embedding methods like DeepWalk.
- The query graph is decomposed into small substructures and fed into the graph neural network with active learner to predict the subgraph counts.





A Learned Sketch for Subgraph Counting:

- Cannot fully utilize the data graph information. The topological information of data graph is somehow ignored.
- Consequently, the model has limited robustness, i.e., the result can be easily affected by minor modification on the query graph.





Goal

- Utilize both query and data graph information.
- Avoid efficiency and scalability issue.
- Improve the subgraph counting accuracy.



Neural Subgraph Counting method: NeurSC





Our recent work

- "Neural Subgraph Counting with Wasserstein Estimator" -- SIGMOD 2022

Substructure Extraction

- Complete Candidate Vertex Set (CS):
 - CS(u) for query vertex $u \in V$ is a set of data vertices $v \in V'$
 - If (u, v) exists in a match from q to G, then $v \in CS(u)$
- First, we determine the complete candidate vertex set for all query vertices using *local pruning* and *global refinement*.
- Based on neighboring and label information
- Candidate set of query $q: CS(q) = \bigcup_{u \in V} CS(u)$
- Induced subgraph of G with vertices CS(q) is used as the candidate substructures, denoted as G_{sub}





Feature Initialization

 $\boldsymbol{x}_{v} = f_{b}(deg_{v})||f_{b}(f_{l}(v))||_{i=1}^{k} MP_{\forall v' \in N^{(i)}(v)} f_{b}(deg_{v'})||f_{b}(f_{l}(v'))$

Degree information Label information Neighbor information

- || denotes the concatenation.
- f_b denotes the binary encoding that converts the decimal digits into binary numbers.
- MP denotes the mean pooling.
- $N^{(i)}(v)$ denotes the *i*-hop neighbors of *v*.



Wasserstein Estimator

- Intra-Graph Neural Network
 - For both query graph and substructure.
 - Capture structural and attribute information.

•
$$h_u^{(k)} = MLP^{(k)}((1+\epsilon^{(k)})h_u^{(k-1)}, \sum_{u'\in N_q(u)}h_{u'}^{(k)})$$

- Inter-Graph Neural Network
 - Construct a bipartite graph for inter-relationship.
 - Capture the mapping relationship between query vertices and corresponding candidate vertices
 - $h_u^{(k)} = \sigma(a_{uu}^{(k)}\theta^{(k)}h_u^{(k-1)}, \sum_{v \in N_{G_B}(u)} a_{uv}^{(k)}\theta^{(k)}h_v^{(k)})$







Wasserstein Estimator

- Readout
 - Sum Pooling
 - Concatenation of intra- and inter-graph representations.
- Prediction
 - Multi-layer perceptron.
- Wasserstein Discriminator
 - Minimize Wasserstein distance between q and G
 - Further utilize the vertex correspondence information between *q* and *G*

•
$$L_w(q, G_{sub}) = \sum_{u \in V'(q)} f_{\omega(h_u)} - \sum_{v \in V'(G_{sub})} f_{\omega(h_v)}$$

- Expressive Power
 - WEst is as powerful as 1-Weisfeiler-Lehman test.





Learning objective and training procedure

- q-error loss
 - $L_c(q) = \max\left(\frac{c(q)}{\hat{c}(q)+\varepsilon}, \frac{\hat{c}(q)}{c(q)}\right)$
 - Optimize the model by reducing the prediction error.
- Wasserstein loss
 - $L_w(q, G_{sub}) = \sum_{u \in V'(q)} f_{\omega(h_u)} \sum_{v \in V'(G_{sub})} f_{\omega(h_v)}$
 - Minimize the Wasserstein distance.
- Overall loss

•
$$L(q) = (1-\beta)L_c(q) - \frac{\beta}{|G_{sub}|} \sum_{g \in G_{sub}} L_w(q,g)$$

Algorithm 3: Training Procedure of WEst **Input:** training query graph set Q_t , data graph G, estimation network f_{θ} , discriminator f_{ω} , learning rates $\alpha_{\theta}, \alpha_{\omega}$, batch size n_{batch} , number of training iterations *iter*_{ω}. 1 Initialize optimizers opt_{θ} , opt_{ω} with learning rates α_{θ} , α_{ω} . ² Separate Q_t into batches $\{Q_b = \{q^{(i)}\}\}$ with n_{batch} query graphs. 3 for $Q_b \in Q_t$ do for $q^{(i)} \in Q_b$ do Generate $\mathcal{G}_{sub}^{(i)}$ for $q^{(i)}$ 5 $X_a^{(i)} \leftarrow$ initial features of vertices in query graph $q^{(i)}$ 6 for $j = 1, ..., |\mathcal{G}_{sub}^{(i)}|$ do 7 $X_{sub}^{(j)} \leftarrow \text{initial features of vertices in } G_{sub}^{(j)}$ $H_q^{(i)}, H_{sub}^{(j)}, \hat{c}_j(q^{(i)}) \leftarrow f_\theta(q^{(i)}, G_{sub}^{(j)}, X_q^{(i)}, X_{sub}^{(j)})$ 8 9 for *iter* $_{\omega}$ do 10 Sample $V'(q^{(i)})$ and $V'(G^{(j)}_{sub})$ 11 Update ω by opt_{ω} minimizing $-\mathcal{L}_{w}$ in Eq. 9 12 Compute $\mathcal{L}_{w}(q^{(i)}, G_{cub}^{(j)})$ 13 $\hat{c}(q^{(i)}) = \sum_{j=1}^{|\mathcal{G}_{sub}|} \hat{c}_j(q^{(i)})$ 14 Compute $\mathcal{L}_c(q^{(i)})$ using Eq. 10 15 Update θ by opt_{θ} with $\sum_{q^{(i)} \in Q_h} \mathcal{L}(q^{(i)})$ defined in Eq. 11 16

Experiment

Experimental Setup

- Dataset
 - 7 data graphs + 5 query sets.
- Compared methods
 - 5 non-learning methods.
 - 2 learning-based methods with 2 variants.
 - 2 variants of *NeurSC* for ablation study.

Parameter settings

- Initial dimension: 64
- Hidden and output dimension 128
- Intra- and Inter-GNN have 2 layers.
- Prediction network is a 4-layer MLP.
- Implementation
 - Substructure extraction: C++
 - WEst: Python + Pytorch Geometric

Table 2: Statistics of Data Graphs

| Dataset | V | E | L | d |
|---------|-----------|------------|-----|------|
| Yeast | 3,112 | 12,519 | 71 | 8.0 |
| Human | 4.674 | 86,282 | 44 | 36.9 |
| HPRD | 9,460 | 34,998 | 307 | 7.4 |
| Wordnet | 76,853 | 120,399 | 5 | 3.1 |
| DBLP | 317,080 | 1,049,866 | 15 | 6.6 |
| EU2005 | 862,664 | 16,138,468 | 40 | 37.4 |
| Youtube | 1,134,890 | 2,987,624 | 25 | 5.3 |

Table 3: Details of Query Graphs

| Dataset | # Queries | Query Sizes | Counts Range |
|---------|-----------|--------------------|-------------------|
| Yeast | 1,632 | {4, 8, 16, 24, 32} | $[10^0, 10^{11}]$ |
| Human | 339 | {4, 8, 16} | $[10^0, 10^{10}]$ |
| HPRD | 1,000 | {4, 8, 16} | $[10^0, 10^4]$ |
| Wordnet | 600 | {4, 8} | $[10^1, 10^9]$ |
| DBLP | 600 | {4, 8} | $[10^3, 10^8]$ |
| EU2005 | 372 | {4, 8} | $[10^4, 10^9]$ |
| Youtube | 811 | $\{4, 8, 16\}$ | $[10^0, 10^{11}]$ |



Experiment Results

Accuracy Evaluation



Evaluation metric







Subgraph Matching

The objective of the *subgraph matching* is searching for all *subgraph isomorphisms* from query graph q to data graph G

Definition II.1 (Subgraph Isomorphism). Given a query graph q = (V, E) and a data graph G = (V', E'), a subgraph isomorphism is an injective function f_{iso} from V to V' such that (1) $\forall v \in V, f_l(v) = f_l(f_{iso}(v))$; and (2) $\forall e_{(u,v)} \in E, e_{(f_{iso}(u), f_{iso}(v))} \in E'$.



Subgraph Isomorphism



(a) Query Graph q

(b) Data Graph G



Subgraph Isomorphism



Subgraph isomorphisms

1.
$$(u_1, u_2, u_3, u_4) \rightarrow (v_1, v_4, v_5, v_{10})$$

2. $(u_1, u_2, u_3, u_4) \rightarrow (v_1, v_4, v_5, v_{11})$
2. $(u_1, u_2, u_3, u_4) \rightarrow (v_1, v_4, v_5, v_{11})$

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$$(u_1, u_2, u_3, u_4) \to (v_1, v_4, v_6, v_{11})$$



Existing Subgraph Matching Methods

There are two major categories of subgraph matching methods:

- Backtracking-based methods
- Join-based methods

In this work, we focus on the backtracking-based methods.



Existing Subgraph Matching Methods

The backtracking-based methods can be partitioned in three main phases:

- 1. The complete candidate vertex set generation.
- 2. Matching order generation.
- 3. Matching enumeration.



Backtracking-based Methods

Complete candidate vertex set generation is to filter out the unpromising vertices, and hence reduce the search space before the enumeration process begins.

Definition II.2 (Complete Candidate Vertex Set C). Given q and G, a complete candidate vertex set C(u) of $u \in V(q)$ is a set of data vertices such that for each $v \in V(G)$, if (u, v) exists in a match from q to G, then $v \in C(u)$.



Backtracking-based Methods

Matching order generation phase generates the matching order ϕ to guide the enumeration of matched subgraphs.

Definition II.3 (Matching Order). A matching order ϕ is a permutation (i.e., sequence) of query graph's vertex set V(q).



Backtracking-based Methods

The enumeration procedure finds all matches of the query subgraph q in the data graph G with given matching order φ .

Definition II.5 (Enumeration Procedure). An enumeration procedure is performed recursively to find subgraph matches f_{iso} with given matching order ϕ and candidate vertex set C.



Subgraph Matching

Subgraph matching has wide applications such as query in graph database and biological graph analytics.

However, it has been proven that the subgraph matching is *NP*-*complete*. We cannot optimize the worst-case time cost.

In this work, we aim to reduce the enumeration time *on the average case* by proposing a novel query vertex ordering method.



- Background
- Motivation
- Framework
- Feature Representations
- Query Vertex Ordering as Markov Decision Process
- Policy Training
- Experiments



Limitations of Existing Order Generation Methods

The existing subgraph matching methods usually generate the matching order based on the heuristic values, here are some examples:

- Infrequent edge first ordering
- Infrequent label first ordering
- Path-based ordering.



Limitations of Existing Order Generation Methods

Two major limitations:

- Cannot fully use the graph information.
- Greedy heuristics can lead to local optimum.



If ordering based on degree (RI)





If ordering based on label frequency





Greedy heuristics can lead to local optimum

The heuristic-based greedy methods can reduce the most redundant intermediate results.

However, these methods cannot consider the long-term query time cost.

The exact optimal order can only be found after all possible order permutations are evaluated.



Our solutions

Motivated by the aforementioned limitations, we proposed the following approaches:

- Capture the graph information with graph neural network.
- Try to approach the global optimal with reinforcement learning.



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Framework



Our recent work

- "Reinforcement learning based query vertex ordering model for subgraph matching" -- ICDE 2022



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We use the important statistical heuristics of query vertices to initialize the input query representations.

Based on the input features, our model can fully exploit the information within the features while preserving the relations between the query and data graphs.



Degree of node:

$$\boldsymbol{h}_{u}^{(0)}(1) = degree(u)/\alpha_{degree},$$

Label of node:

$$\boldsymbol{h}_{u}^{(0)}(2) = label(u)$$

Query node ID:

$$\boldsymbol{h}_u^{(0)}(3) = id(u)$$



Frequency of data vertices with greater degree than the query node
 h⁽⁰⁾_u(4) = |{v ∈ G|d(u) < d(v)}| / (|V(G)| × α_d);

Frequency of data vertex with same label as query vertex:

$$\boldsymbol{h}_{u}^{(0)}(5) = \left| \{ v \in G | L(u) = L(v) \} \right| / (|V(G)| \times \alpha_{l});$$



Lastly, we put two indicator variables in the initial feature:

Number of unordered vertices:

$$h_u^t(6) = |V(q)| - t + 1$$

Trailing indicator that shows whether the node has been ordered:

$$\boldsymbol{h}_{u}^{t}(7) = \mathbb{1}(u \in \phi_{t-1})$$



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- Policy Training
- Experiments



Query Vertex Ordering as Markov Decision Process

To exploit the reinforcement learning, we need to model our query vertex ordering problem as a Markov decision process (MDP).

Specifically, we need to define:

State space

Action probability

Action space

Reward





In this work, the state space is defined as the all (partial) vertex order sequence.

Specifically, we use ϕ_t (the order at time step t) to denote the state at time step t.



Action Space

With the order ϕ_t (state of the MDP), we define the action space as $N(\phi_t) = \{N(u) | \forall u \in \phi_t, N(u) \notin \phi_t\}$ which is the neighbor vertices set of the ordered vertices in ϕ_t .

Consequently, the action at each time step t is to select a node from the action space (neighbor set $N(\phi_t)$).



Action Probability

With the state and corresponding action space.

The key problem is to compute the probabilities for each action in the action space.

To this end, we design a Graph Neural Network (GNN)-based policy network.





Policy Network

Graph convolutional network:

$$\boldsymbol{H}^{(l+1)} = \sigma(\tilde{\boldsymbol{D}}^{-\frac{1}{2}} \tilde{\boldsymbol{A}} \tilde{\boldsymbol{D}}^{-\frac{1}{2}} \boldsymbol{H}^{(l)} \boldsymbol{W}^{(l)})$$

Action probability with the state at time step t:

$$\mathbb{P}_{u'}^{(t)} = \pi(\cdot|S^{(t)}) = Softmax(mask_{u'\in AS(t)}(\boldsymbol{W}_2 \cdot \sigma(\boldsymbol{W}_1 \boldsymbol{h}_{u'}^{(t)})))$$



Reward Design

Three main rewards:

- Enumeration reward
- Step-wise validate reward
- Entropy reward



Enumeration Reward

Definition II.6 (Enumeration Number). An enumeration number $\#_{enum}$ is the number of *recursive* calls of the enumeration procedure to find all matches with given q, G, ϕ and C.

The reduced enumeration number:

$$\Delta \#_{enum} = \#_{enum}(\phi) - \#_{enum}(\phi_{base})$$

Enumeration reward:

 $r_{enum} = f_{enum}(\Delta \#_{enum}).$



Other rewards

Step-wise validation: $r_{val,t}$



Entropy reward:

$$r_{h,t} = H(\dot{P}_{\pi_{\theta}}(\phi_t, N(\phi_t)))$$

Ensure the model can perform more actions.



Reward Design

With the aforementioned rewards, our overall rewards at time step t is as follows:

$$R_t = r_{enum} + \beta_{val} \cdot r_{val,t} + \beta_h \cdot r_{h,t}$$

The overall reward is as follows:

$$R_{q,\theta} = \sum_{t=1}^{|V(q)|} \gamma^t R_t,$$



- Background
- Motivation
- Framework
- Feature Representations
- Query Vertex Ordering as Markov Decision Process
- Policy Training
- Experiments



Policy Training

In this work, we use the proximal policy optimization (PPO) to train our policy network with the following loss function:

$$J_r^{(t)}(\theta) = \sum_{(a_t, s_t)} \min(\frac{\pi_{\theta}(a_t|s_t)}{\pi_{\theta'}(a_t|s_t)} r_t(\theta), clip(\frac{\pi_{\theta}(a_t|s_t)}{\pi_{\theta'}(a_t|s_t)}, 1 - \epsilon, 1 + \epsilon) r_t(\theta))$$
$$J(\theta) = \sum_{t=1}^{|V(q)|} J_r^{(t)}(\theta)$$



- Background
- Motivation
- Framework
- Feature Representations
- Query Vertex Ordering as Markov Decision Process
- Policy Training
- Experiments



Experiment Setup

Dataset Statistics

| Dataset | <i>V</i> | <i>E</i> | L | d |
|----------|-----------|------------|----|------|
| Citeseer | 3,327 | 4,732 | 6 | 1.4 |
| Yeast | 3,112 | 12,519 | 71 | 8.0 |
| DBLP | 317,080 | 1,049,866 | 15 | 6.6 |
| Youtube | 1,134,890 | 2,987,624 | 25 | 5.3 |
| Wordnet | 76,853 | 120,399 | 5 | 3.1 |
| EU2005 | 862,664 | 16,138,468 | 40 | 37.4 |

| Compared methods: | |
|----------------------|--|
| QuickSI | |
| RI | |
| VF2++ | |
| VEQ | |
| Hybrid | |
| RL-QVO | |



Evaluation Metrics

- Query processing time
- Enumeration time



Average Query Processing Time





Query Processing Time Percentile Comparison





Enumeration Time Comparison





Fraud Detection



Attributed Bipartite Graph

An attributed bipartite graph is a type of graph which consists of two sets of vertices that are linked by edges. The vertices have additional attributes, making this graph particularly useful for **representing information in the field of e-commerce**.





Group-based Frauds on Attributed Bipartite Graphs

Kate

*u*₂

 u_3

 (u_4)

Group-based fraud is becoming increasingly common: "Ride Item's Coattails" attack (edge classification) Sockpuppet-based Targeted Attack on Reviewing Systems

(STARS attack) (vertex classification) Lisa Sam Bob Andy Kane Eric (u_1) (u_2) (u_3) (u_4) (u_5) (u_6)







SOTA method for "Ride Item's Coattails" attack

RICD ((α , k1, k2)-biclique): **fraud detection method** for "Ride Item's Coattails" attack. Can only utilize structural information.

Tianchi competition winner's algorithm: classification method. Can only use attribute information.





SOTA method for STARS attack

RTV: fraud detection method for Sockpuppet-based Targeted Attack on Reviewing Systems (STARS). Unable to make good use of label information.

Algorithm RTV **Input:** Rating graph $G = (\mathcal{U} \cup \mathcal{P}, \mathcal{R}, sc)$, weights $\alpha_1, \alpha_2, \beta_1, \beta_2, \gamma_1, \gamma_2, \gamma_3, \gamma_4$, threshold ϵ **Output:** fair(u) $\forall u \in \mathcal{U}$, good(p) $\forall p \in \mathcal{P}$, rel(u, p) $\forall (u, p) \in \mathcal{R}$ for each $u \in \mathcal{U}$, fair₀(u) \leftarrow norm(u) for each $p \in \mathcal{P}$, $good_0(p) \leftarrow norm(p)$ for each $(u, p) \in \mathcal{R}$, rel₀ $(u, p) \leftarrow \text{norm}(u, p)$ $\mu_f \leftarrow \frac{\sum_{u \in \mathcal{U}} \operatorname{fair}_0(u)}{|\mathcal{U}|}, \mu_g \leftarrow \frac{\sum_{p \in \mathcal{P}} \operatorname{good}_0(p)}{|\mathcal{P}|}$ $t \leftarrow 1$ 5 for each $u \in \mathcal{U}$, fair_t(u) \leftarrow value computed as specified in Section 4.1, with rel(u, p) = rel_{t-1}(u, p) 6 for each $p \in \mathcal{P}$, good_t(p) \leftarrow value computed as specified in Section 4.1, with rel(u, p) = rel_{t-1}(u, p) for each $(u, p) \in \mathcal{R}$, rel_t $(u, p) \leftarrow$ value computed as specified in Section 4.1, with fair $(u) = \text{fair}_t(u)$ $\Delta \leftarrow \max\left(\sum_{u \in \mathcal{U}} |\operatorname{fair}_t(u) - \operatorname{fair}_{t-1}(u)|, \sum_{p \in \mathcal{P}} |\operatorname{good}_t(p) - \operatorname{good}_{t-1}(p)|, \sum_{(u,p) \in \mathcal{R}} |\operatorname{rel}_t(u,p) - \operatorname{rel}_{t-1}(u,p)|\right)$ 9 if $\Delta > \epsilon$ or t = 1 then $t \leftarrow t + 1$ and go to Line 6 10 **return** fair_t(u) $\forall u \in \mathcal{U}$, good_t(p) $\forall p \in \mathcal{P}$, rel_t(u, p) $\forall (u, p) \in \mathcal{R}$



Existing methods

Classification Methods:

• Imbalanced labeled vertices, community information.

Cohesive Subgraph Mining Methods:

• Attribute and label information, suffer from NP-completeness.

Fraud Detection Methods:

• Global topological and attribute information, label information, manual parameter setting.



Group-based Fraud Detection method: GFDN





Our recent work

- "Group-based Fraud Detection Network on e-Commerce Platforms" -- KDD 2023

Group-based Fraud Detection method: GFDN





Group-based Fraud Detection method: GFDN





Group-based Fraud Detection method: GFDN





Structural Feature Initialization

(*α*, *β*)-core:

Given a bipartite graph G and integers α , $\beta \in Z^+$, (α, β) -core of G is denoted as G ' which consists of two vertex sets U' \subseteq U and V' \subseteq V.

The (α, β) -core G ' is a maximal bipartite subgraph induced by U' \cup V' from G in which all the vertices in U' have degrees at least α and all the vertices in V' have degrees at least β .





Structural Feature Initialization

GFDN will generate structural features for vertices based on their existence in different (α , β)-core.

$$\hat{X}_{(\mathcal{U},s)} = X_{(\mathcal{U},s)} \odot (I_{\mathcal{U}}W_{(\mathcal{U},s)}), \ \hat{X}_{(\mathcal{V},s)} = X_{(\mathcal{V},s)} \odot (I_{\mathcal{V}}W_{(\mathcal{V},s)})$$
Structural Features Element-wise Product All-ones Vector Weight Matrix

 $W_{(\mathcal{V},s)}$ Attributed Bipartite Graph $(\alpha,\beta) - \operatorname{core}$

 $W_{(\mathcal{U},s)}$

Fraudster Community Detection

BDCN - Autoencoder:

Autoencoder in Bipartite Deep Clustering Network (BDCN) can:

1. preserving both structural and attribute information from the input features.

2. Generate high-quality community representation for customer vertices.

It can achieve self-supervised fraud **community detection** using a loss function measures with Student's t-distribution kernel.





Fraudster Community Detection

BDCN - GNN:

GNN in BDCN can aggregate on attribute bipartite graph and preserve the attribute information and structural information of the graph. The output of encoding layer will be used.




GFDN

Training Objective

"Ride Item's Coattails" Attack:

In "Ride Item's Coattails" attack, not all edges related to fraudsters necessarily have attack implications. GFDN will perform **multi-task training** on this issue, predicting both **fraudsters** and **fraudulent attack**.

STARS Attack:

STARS attack detection aims to **detect fraudsters**, in which case GFDN only needs to perform the vertex classification task.





GFDN

Training Objective

The final loss function will be composed of the loss functions of the aforementioned modules, including **autoencoder**, **community prediction**, **fraudster prediction**, and **fraudulent prediction**. The sum of the weights of all parts of them is 1.

$$\mathcal{L} = \omega_{ae} \mathcal{L}_{ae} + \omega_c \mathcal{L}_c + \omega_l \mathcal{L}_l + \omega_e \mathcal{L}_e$$
Autoencoder Community Fraudster Fraudulent



Experimental Setup

Dataset

- 4 real-life datasets.
- Compared methods
 - 5 learning-based methods.
 - 2 pattern-based methods.
 - 4 fraud detection methods.
 - A naïve model and four ablated GFDNs

Parameter settings

- The number of GNN layer: 4.
- The number of community: 32.
- Hidden dimension: 128.
- The selected GNN is GraphSAGE.
- Implementation
 - Structure information extraction: C++
 - Other Parts of the Model : Python + Pytorch Geometric.

Table 1: Datasets for "Ride Item's Coattails" Attack Detection

| Dataset | 3 | $ \mathcal{U} $ | $ \mathcal{V} $ | % Fraudulent | % Legitimate |
|---------|-----------|-----------------|-----------------|--------------|--------------|
| TB | 3,085,653 | 996,090 | 381,611 | 0.62% | 3.53% |
| TC | 1,050,000 | 532,345 | 239,840 | 2.86% | 11.43% |

Table 2: Datasets for STARS Attack Detection

| Dataset | 3 | $ \mathcal{U} $ | $ \mathcal{V} $ | % Fraudulent | % Legitimate |
|---------|--------|-----------------|-----------------|--------------|--------------|
| Alpha | 24,186 | 3,286 | 3,754 | 3.10% | 4.20% |
| OTC | 35,592 | 4,814 | 5,858 | 3.70% | 2.80% |



Effectiveness Evaluation Results for "Ride Item's Coattails" Detection

| | TB Data | | | | TC Data | | | | | |
|-------------------------|---------|--------|--------|--------|---------|--------|--------|--------|--------|--------|
| | F1 | Acc | AUC | Pre | Recall | F1 | Acc | AUC | Pre | Recall |
| LPA | 0.2737 | 0.4627 | 0.5517 | 0.1715 | 0.6785 | 0.2056 | 0.4284 | 0.5276 | 0.1219 | 0.6557 |
| SBGNN | 0.4789 | 0.8228 | 0.7947 | 0.4279 | 0.5438 | 0.3676 | 0.8074 | 0.7666 | 0.2900 | 0.5018 |
| BiGI | 0.5359 | 0.8540 | 0.8491 | 0.5097 | 0.5649 | 0.4039 | 0.8292 | 0.8044 | 0.3331 | 0.5129 |
| SIHG | 0.6449 | 0.8709 | 0.8692 | 0.5470 | 0.7853 | 0.5947 | 0.8771 | 0.8985 | 0.4735 | 0.7992 |
| Tianchi | 0.6446 | 0.8752 | 0.9342 | 0.5606 | 0.7581 | 0.5364 | 0.8717 | 0.9107 | 0.4527 | 0.6583 |
| RICD | 0.6518 | 0.8405 | 0.9063 | 0.4834 | 1.0000 | 0.4784 | 0.8482 | 0.7474 | 0.3906 | 0.6171 |
| (α, β) -core | 0.8081 | 0.9449 | 0.8757 | 0.8417 | 0.7770 | 0.6348 | 0.8907 | 0.8696 | 0.5093 | 0.8423 |
| FRAUDAR | 0.2580 | 0.1481 | 0.4963 | 0.1483 | 0.9927 | 0.2020 | 0.1124 | 0.4981 | 0.1124 | 0.9961 |
| CF1 | 0.2407 | 0.7698 | 0.5532 | 0.2371 | 0.2445 | 0.1620 | 0.7981 | 0.5253 | 0.1523 | 0.1731 |
| CF2 | 0.4675 | 0.7603 | 0.7376 | 0.3497 | 0.7052 | 0.3588 | 0.6837 | 0.7277 | 0.2326 | 0.7844 |
| Naive | 0.8109 | 0.9473 | 0.9844 | 0.8736 | 0.7565 | 0.6397 | 0.9090 | 0.9516 | 0.7816 | 0.5414 |
| GFDN-S | 0.6867 | 0.9202 | 0.9653 | 0.8284 | 0.5864 | 0.6122 | 0.8783 | 0.9342 | 0.4780 | 0.8514 |
| GFDN-F | 0.9212 | 0.9754 | 0.9886 | 0.8821 | 0.9639 | 0.6401 | 0.8976 | 0.9287 | 0.5302 | 0.8076 |
| GFDN-L | 0.9398 | 0.9813 | 0.9964 | 0.9050 | 0.9775 | 0.7015 | 0.9192 | 0.9654 | 0.6014 | 0.8417 |
| GFDN-C | 0.9423 | 0.9821 | 0.9967 | 0.9086 | 0.9785 | 0.7048 | 0.9226 | 0.9646 | 0.6181 | 0.8198 |
| GFDN | 0.9522 | 0.9853 | 0.9974 | 0.9254 | 0.9806 | 0.7226 | 0.9242 | 0.9713 | 0.6154 | 0.8752 |



Comparison with Pattern-based Algorithms





Query Time Evaluation of "Ride Item's Coattails" Detection





Effectiveness Evaluation Results for STARS Detection

| | | | Alpha | | | | | OTC | | |
|----------------------------------|------------------|-------------------------|------------------|-------------------------|-------------------------|------------------|------------------|------------------|------------------|-------------------------|
| | F1 | Acc | AUC | Pre | Recall | F1 | Acc | AUC | Pre | Recall |
| FRAUDAR RTV-SUP | 0.3800 0.8652 | 0.2626 0.9452 | 0.5236 0.8859 | 0.2346 0.9747 | 1.0000 0.7778 | 0.3780 0.7010 | 0.2547 0.8082 | 0.5183 0.8736 | 0.2330 0.5417 | 1.0000 0.9931 |
| (α, β) -core Naive | 0.7857 0.8089 | 0.8767 0.9018 | 0.9204 0.9789 | 0.6471 0.7222 | 1.0000 0.9192 | 0.7784 0.7937 | 0.8711 0.8978 | 0.9167 0.9508 | 0.6372 0.7310 | 1.0000 0.8681 |
| GFDN | 0.8919 | 0.9452 | 0.9913 | 0.8049 | 1.0000 | 0.9231 | 0.9623 | 0.9746 | 0.8571 | 1.0000 |



Efficiency Evaluation Results for STARS Detection





Thank you!

Q&A

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