

MrMine: Multi-resolution Multi-network embedding

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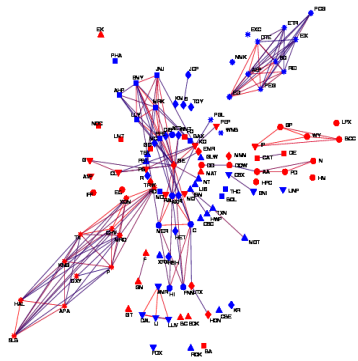


Hanghang Tong

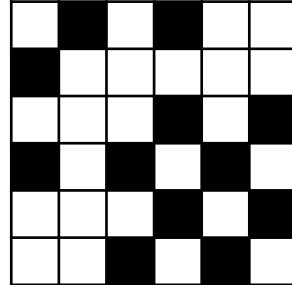
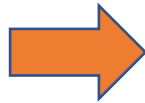
Network Embedding: Why?

Goal: Map network object (node/edge/subgraph/network) into a low-dimensional space.

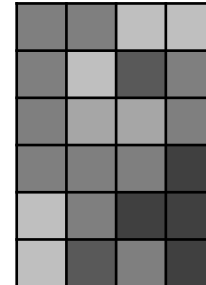
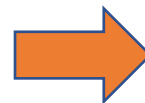
- Distributed representations for network objects
- Encode network characteristics into continuous vector space
- Automatic feature learning for downstream tasks on networks



Input Network

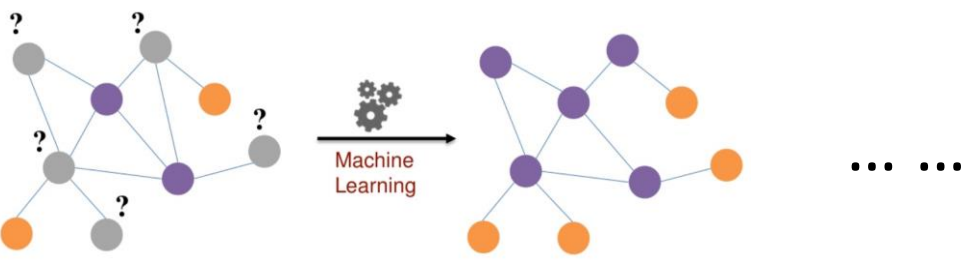


Adjacency matrix

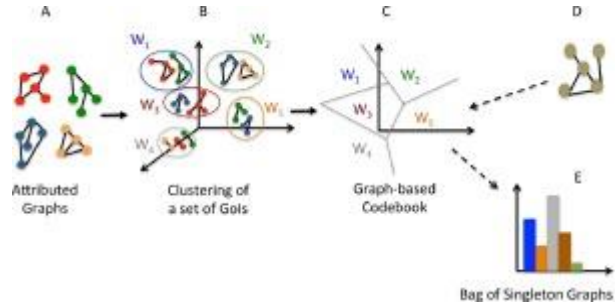


Low-dimensional representation

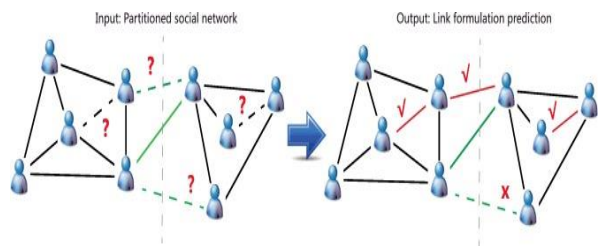
Network Embedding: Applications



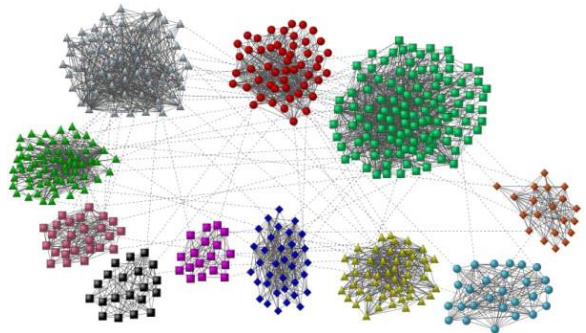
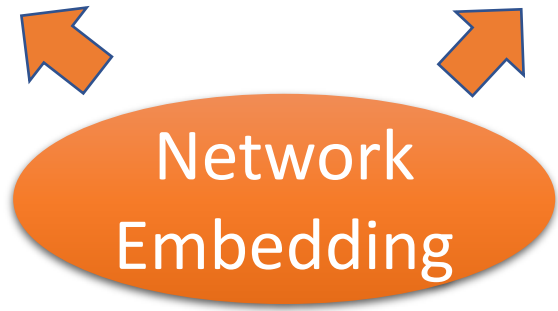
Node classification
[J Leskovec '17]



Network classification
[F Silva, et al '17]



Link prediction
[J Wu, et al '17]

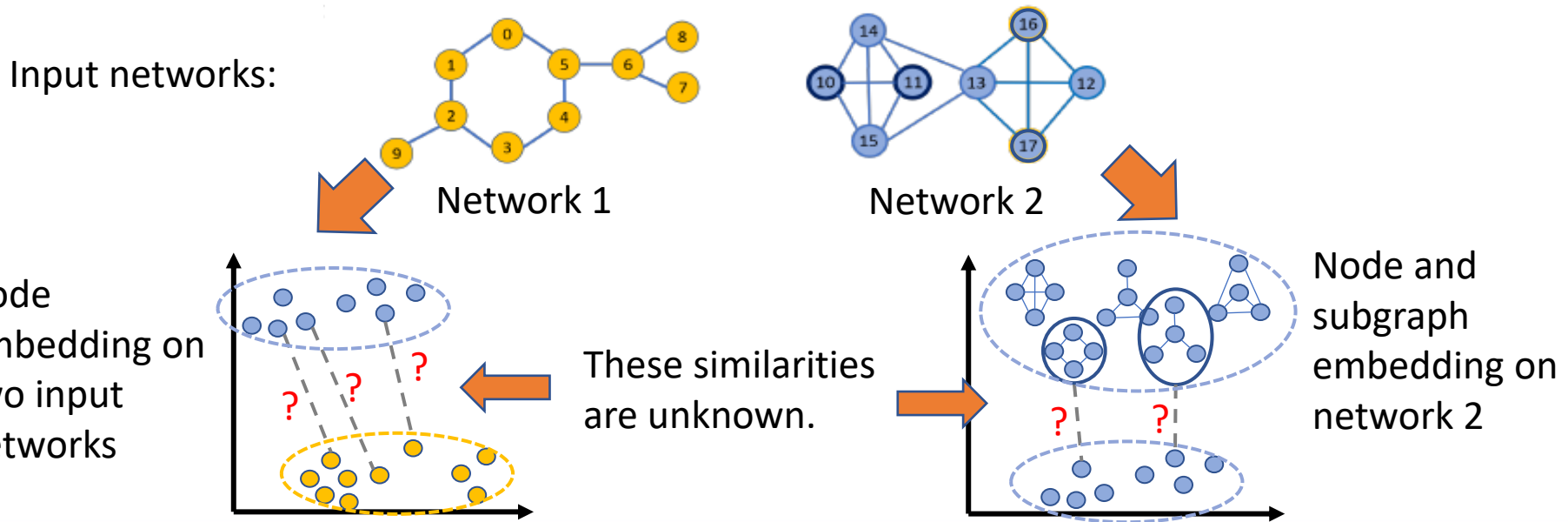


Community detection
[S Fortunato '09]

... ..

Limitations of Existing Methods

- Single resolution:
 - Embedding on nodes/subgraphs/networks
 - Problem: Embeddings across resolutions are separate
- Single network:
 - Embedding on one single network
 - Problem: Embeddings on multiple networks are separate



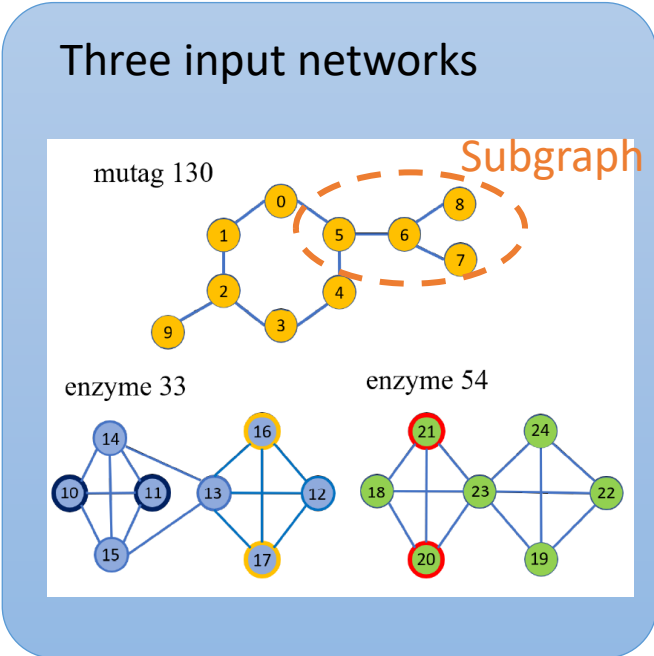
Limitations of Existing Methods (cont'd)

Name	Reference	Multi network	Multi resolution
<i>deepwalk</i>	B Perozzi, SIGKDD '14	✗	✗
<i>LINE</i>	J Tang, WWW '15	✗	✗
<i>Node2vec</i>	A Grover, SIGKDD '16	✗	✗
<i>subgraph2vec</i>	A Narayanan '16	✗	✓
<i>Deep graph kernel</i>	P Yanardag SIGKDD '15	✓	✗
<i>struc2vec</i>	LFR Ribeiro SIGKDD '17	✓	✗
...

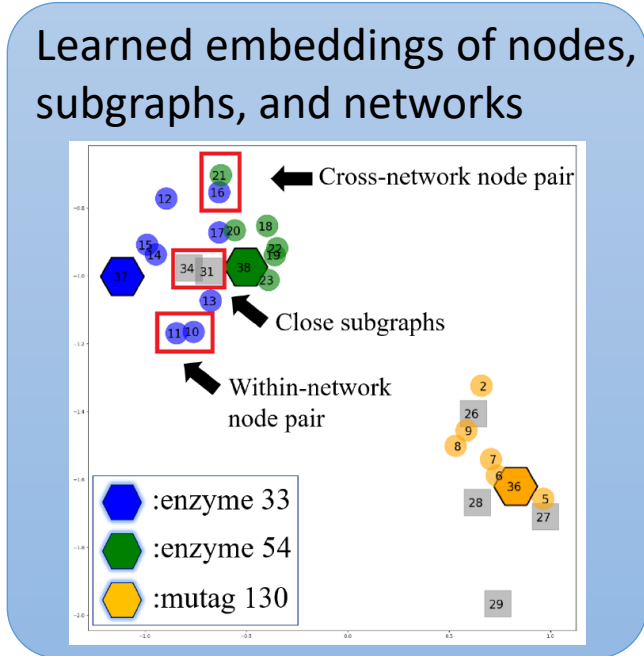
Name	Reference	Multi network	Multi resolution
MrMine	This paper	✓	✓
MrMine+	This paper	✓	✓

Our methods

Multi-resolution Multi-network embedding

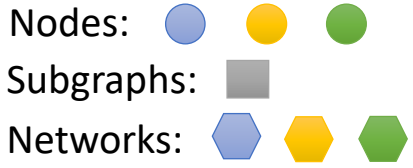


MrMine Learning model



Goals:

- Unsupervised embedding learning.
- Embeddings from different networks are comparable.
- Embeddings from different resolutions are comparable.
- Objects with close structural characteristics are close in embedding space.



Roadmap

- Motivation
- **Problem Definition**
- Proposed Solution: 'MrMine'
- Experiments
- Conclusions

Problem Definition

Given:

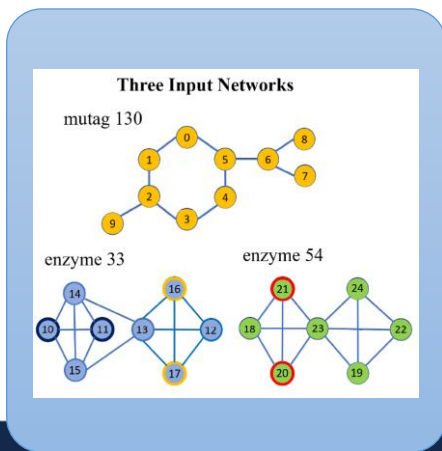
- The inputs for context building:
 - A set of networks \mathbf{G} ;
 - The dimension of embedding vectors \mathbf{p} ;
 - Subgraph constraints (e.g. the maximum height of WL subtrees H);
- Parameter set for language model for embedding learning.

Find: Embedding matrices \mathbf{F}_g , \mathbf{F}_s , and \mathbf{F}_n for:

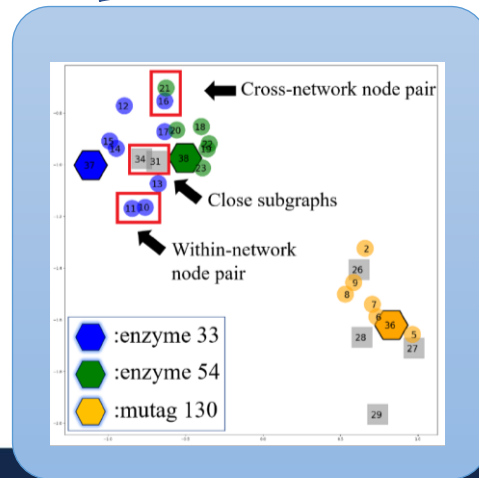
- All input networks in \mathbf{G} ;
- All extracted subgraphs in \mathbf{S} ;
- All nodes in \mathbf{G} .

We specify one subgraph type in proposed method.

With all embeddings in the same space.



MrMine
Learning
model

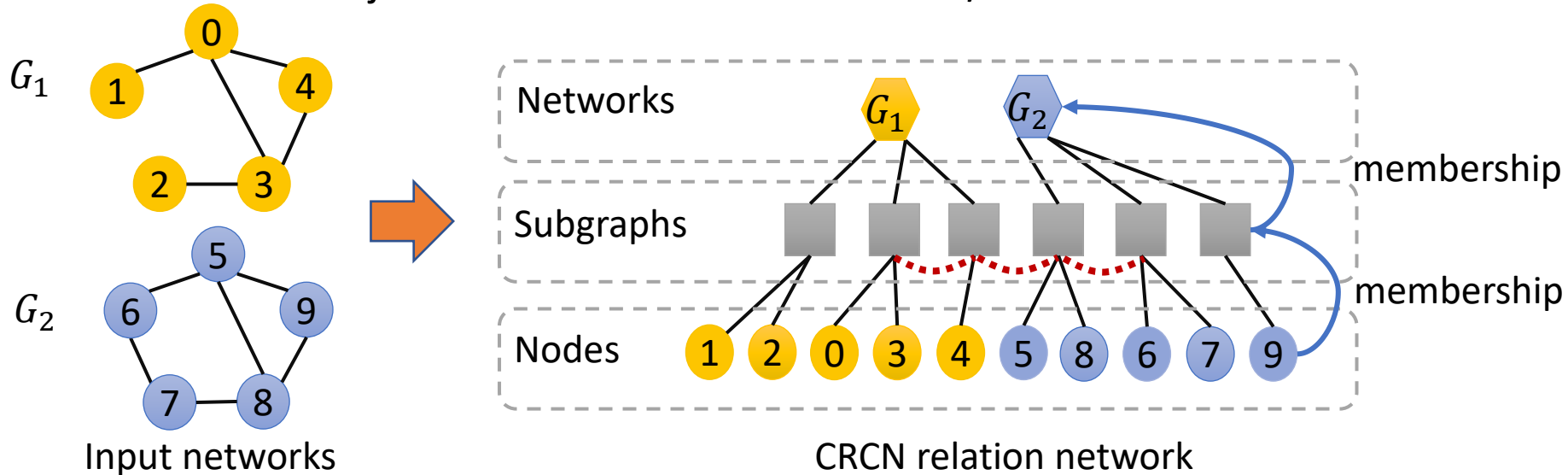


Roadmap

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- **Proposed Solution: 'MrMine'**
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MrMine: Challenges and Key Ideas

- C1: How to build context for cross-resolution cross-network objects?
 - Idea: Cross-Resolution Cross-Network (CRCN) relation network
 - Adv: Relate objects across different resolutions/networks

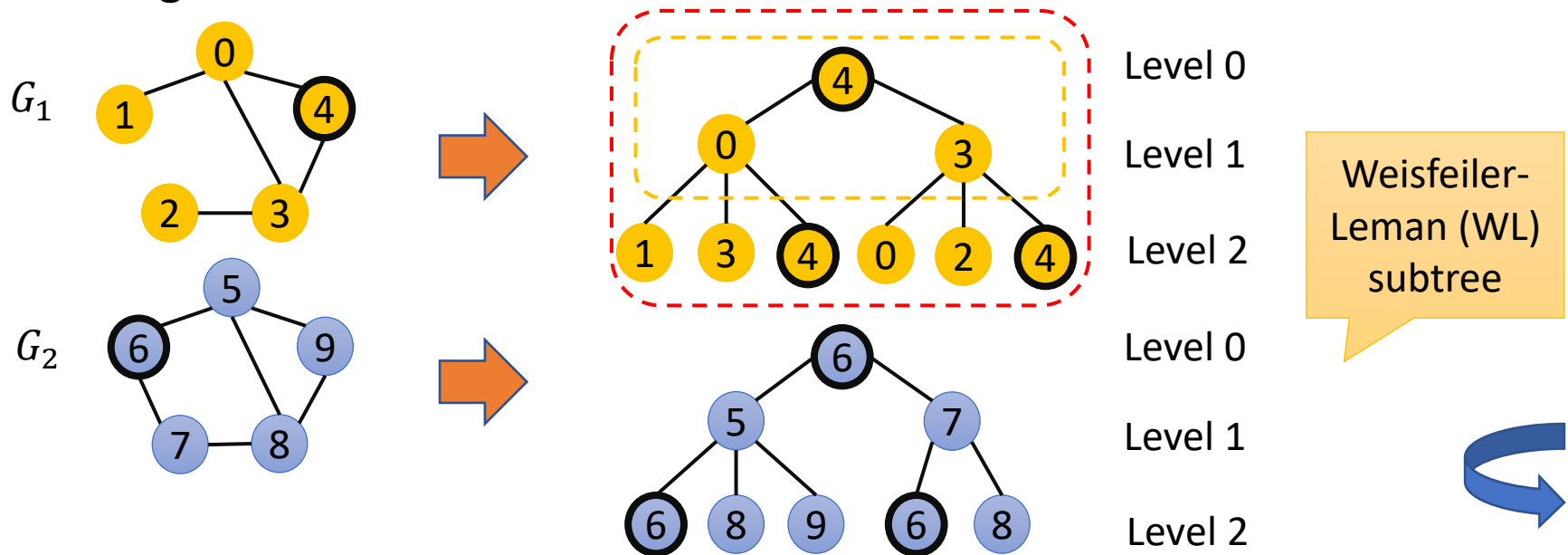


- Remarks:
 - Objects of multi-resolution and multi-network are vertices in the same network.
 - Links across different resolutions represent membership relation (e.g. blue arrow).
 - Links in the same layer on the subgraph resolution represents subgraph similarity.

MrMine: Challenges and Key Ideas (cont'd)

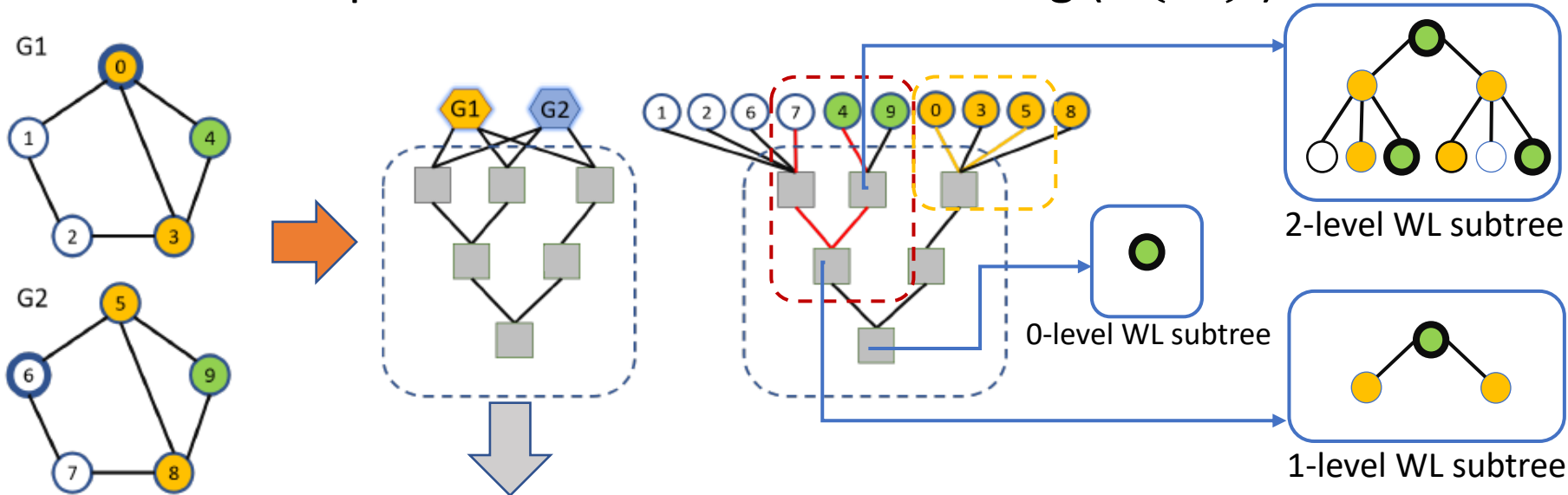
C2: How to construct the links of CRCN relation network?

- Idea: WL subtree as the subgraph resolution
- Adv:
 - WL label transformation is efficient ($O(hm)$, h : constant; m : # of edges)
 - 'Borderless' across different networks (allow links across networks)
 - Bridge between node resolution and network resolution



MrMine: Challenges and Key Ideas (cont'd)

- C3: How to reduce computation costs to build CRCN relation network?
 - Idea: Hierarchical structure of CRCN network (H-CRCN)
 - Adv:
 - Hierarchical structure of WL subtrees for finer resolutions
 - Avoid explicit cross-network link building ($O(n^2)$)

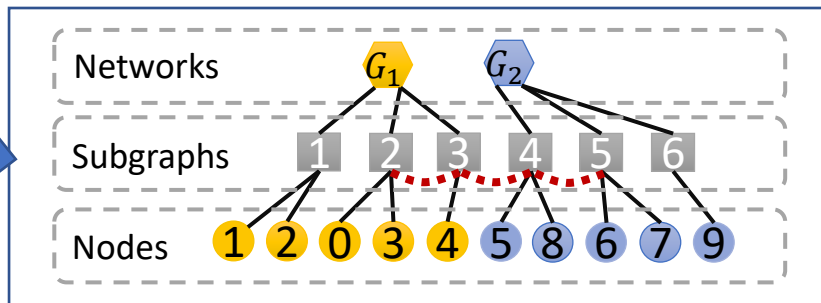


*Gray rectangle: hierarchical relation network of WL subtrees

MrMine: Method Details

- Step 1: H iterations of WL node label transformation:
 - Generate unique WL subtrees of height H as subgraphs
- Step 2: Function $f(S_i, S_j)$ to calculate the subgraph similarity:
 - Option 1: $f(S_i, S_j) = \sum_h DTW(Q_{S_i}^h, Q_{S_j}^h)$
 - Option 2: $f(S_i, S_j) = \sum_h \sum_t |\tilde{Q}_{S_i}^h(t) - \tilde{Q}_{S_j}^h(t)|$
- Step 3: Construct CRCN relation network:
 - Add nodes, subgraphs, networks as **new vertices** in CRCN network
 - Add **cross resolution links** based on membership relation
 - Add **cross network links** based on $f(S_i, S_j)$ and threshold σ .
- Step 4: Apply truncated random walk for corpus generation and Skipgram model
- Time complexity:
 - $O(Hn \log(n))$ (n: # of nodes, H: small constant)

Graph kernel can also be used, but suffers from computational cost ($O(n^3)$) [1].



Notation:

$Q_{S_i}^h$: sorted degree sequence of S_i on level h
 DTW : Dynamic Time Wrapping
 $\tilde{Q}_{S_i}^h$: sorted degree sequence on level h with zero filling

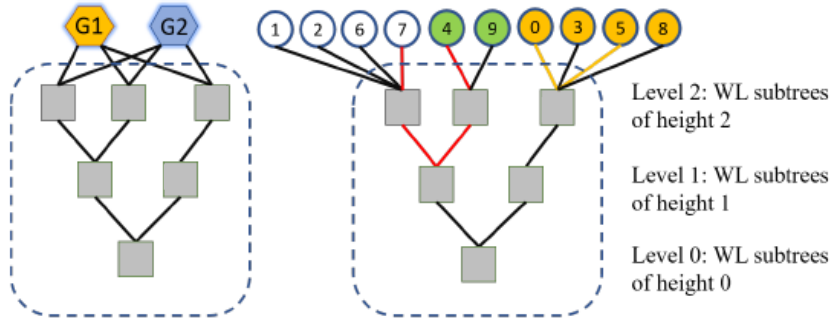
[1] Vishwanathan, S. V. N., Karsten M. Borgwardt, and Nicol N. Schraudolph. "Fast computation of graph kernels." *NIPS*. Vol. 19. 2006.

[2] Ribeiro, Leonardo FR, Pedro HP Saverese, and Daniel R. Figueiredo. "struc2vec: Learning node representations from structural identity." *Proceedings of the 23rd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining*. ACM, 2017.

MrMine+: Method Details

- Step 1: H iterations of WL node label transformation:
 - Generate unique WL subtrees of up to height H as subgraphs
- Step 2: Construct H-CRCN relation network:
 - Add **subtree vertices** to hierarchical relation network of WL subtrees (HRN)
 - Add **links** to HRN based on WL subtree generation relation
 - Attach **network vertices/node vertices** to the last level of HRN
- Step 3: Apply truncated random walk on two H-CRCN networks for corpus generation and Skipgram model
- Time complexity:
 - $O(Hm + cn)$ (n: # of nodes; m: # of edges; H, c: small constants)

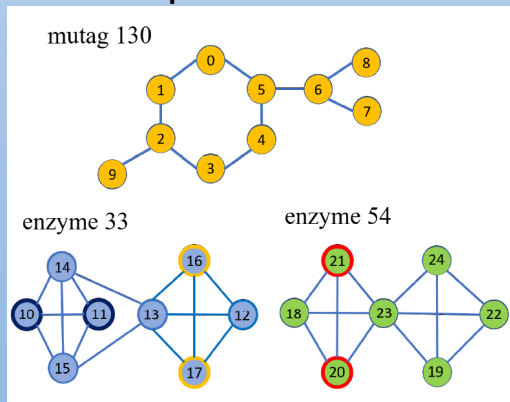
More in the paper



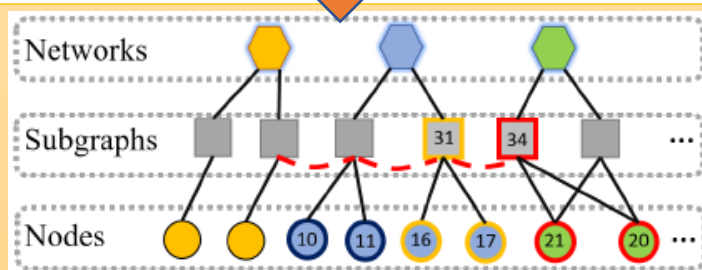
Nodes in the (H-)CRCN relation networks are called vertices

Summary of major steps

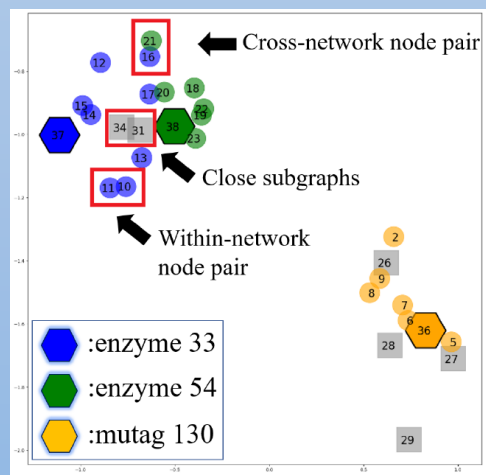
Three input networks



WL label transformation for subgraph generation

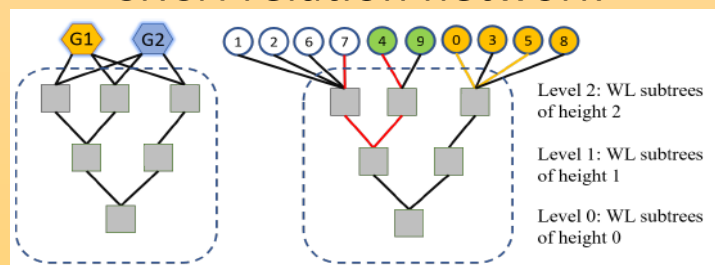


Learned embeddings of nodes, subgraphs, and networks



Or

CRCN relation network



H-CRCN relation network

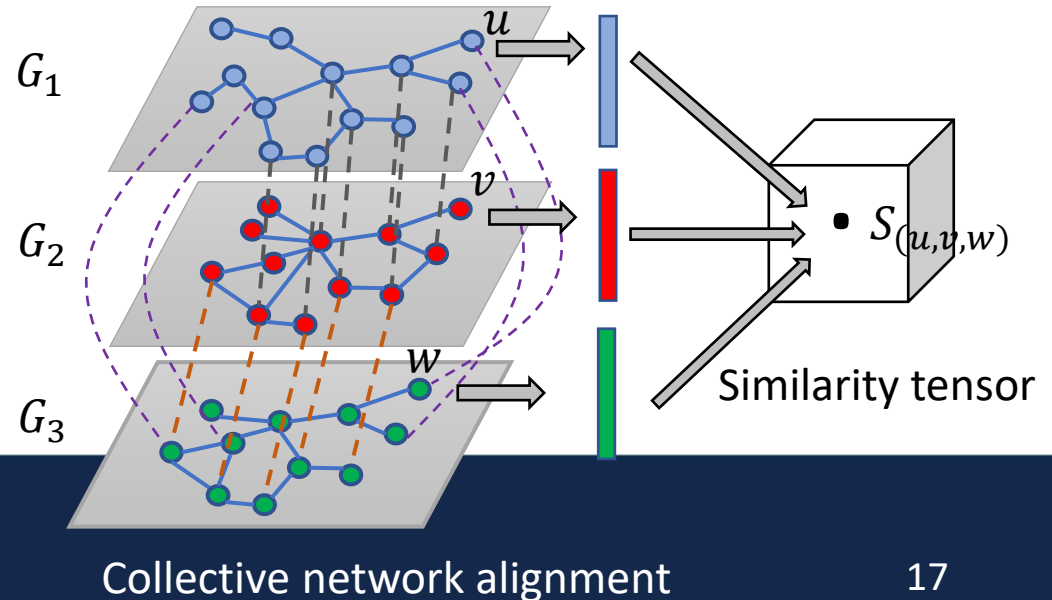
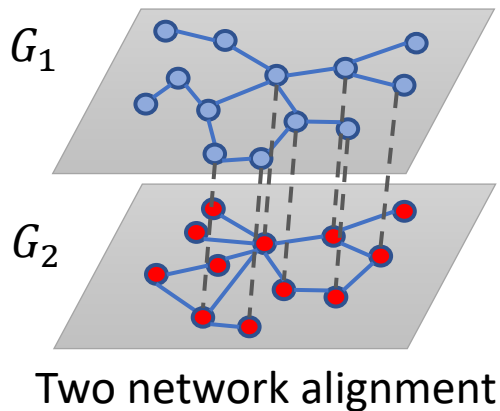
Truncated random walk for corpus building and Skipgram model

Roadmap

- Motivation
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- Proposed Solution: 'MrMine'
- **Experiments**
- Conclusions

Experimental Setup (downstream tasks)

- Query node retrieval:
 - Given a set of nodes from G_1 , retrieve similar nodes from G_2 .
- Network classification:
 - Classify networks into different categories.
- Two network alignment:
 - Align nodes from two input network G_1, G_2 .
- Collective Network alignment:
 - Collectively align nodes in input networks G_1, G_2, G_3 .



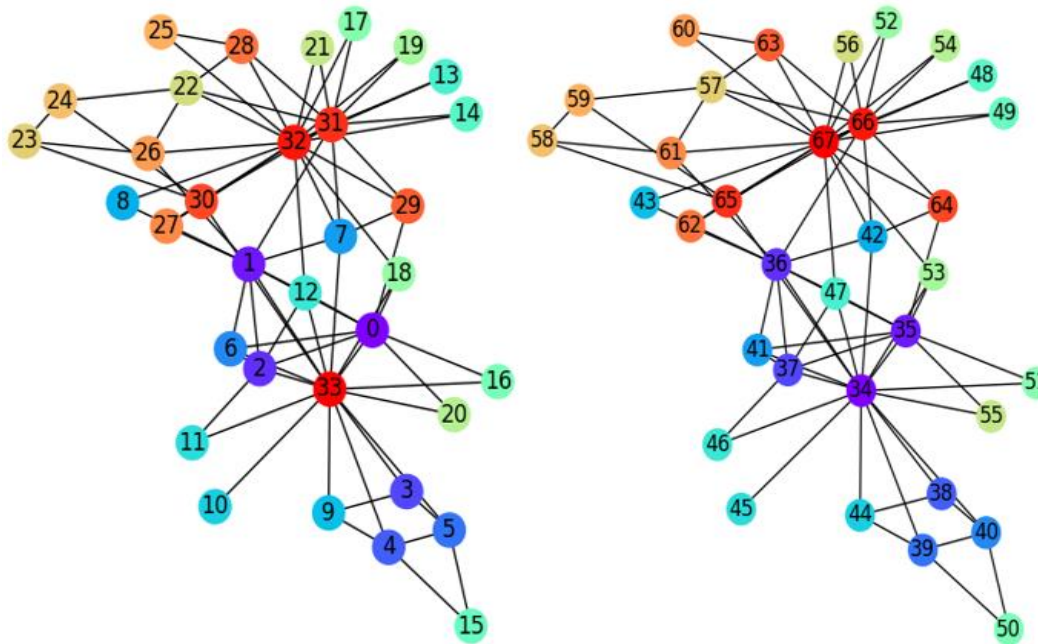
Real-world Datasets

Name	Category	# of Nodes	# of Edges
DBLP	Co-authorship	1,013	3,022
Flickr	User relationship	3,911	4,152
LastFm	User relationship	4,068	4,347
Douban	User relationship	1,118	3,022
MySpace	Social network	6,362	6,514
Aminer	Academic network	1,274,360	4,756,194

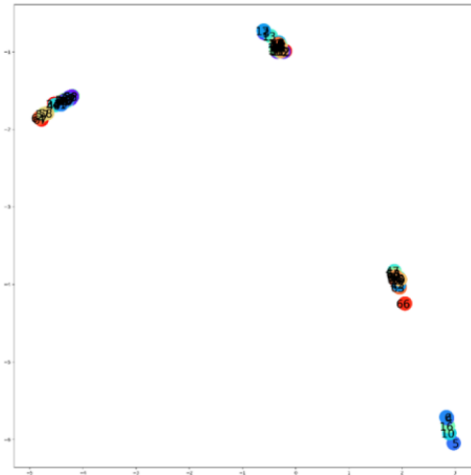
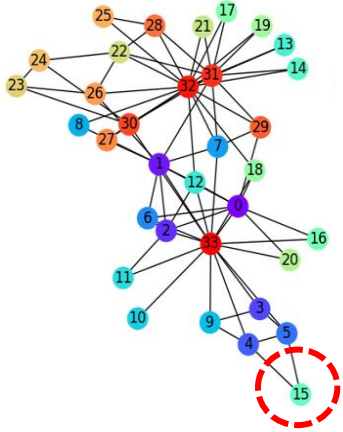
Bioinformatics	# of graphs	Classes	Avg. nodes
MUTAG	188	2	17.9
PTC	344	2	25.5
PROTEINS	1113	2	39.1
NCI1	4110	2	29.8
NCI109	4127	2	29.6

Visualization

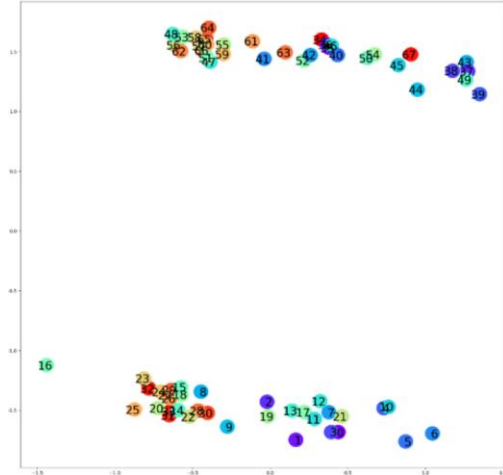
- Dataset: Mirrored Zachary's Karate Club data
- Corresponding nodes are colored the same.
- Learn the node embeddings first and project onto 2-D space.



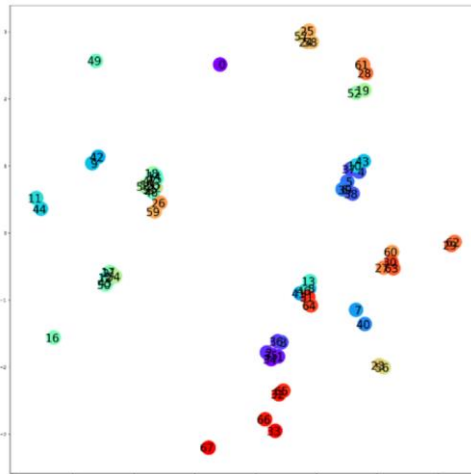
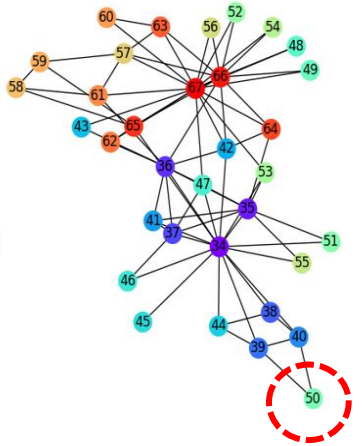
Visualization results



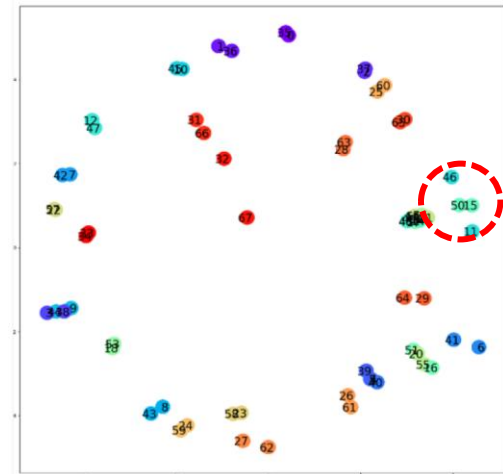
deepwalk



node2vec



struc2vec



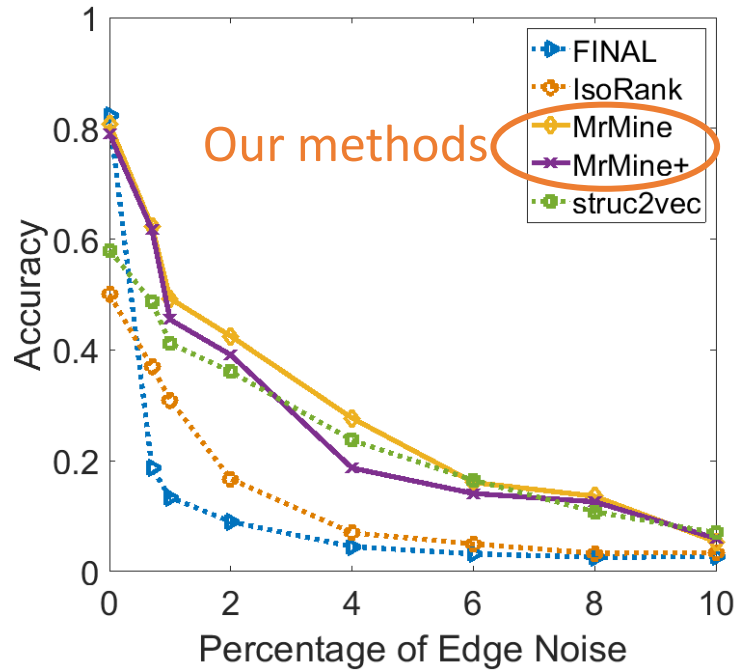
MrMine

Paired

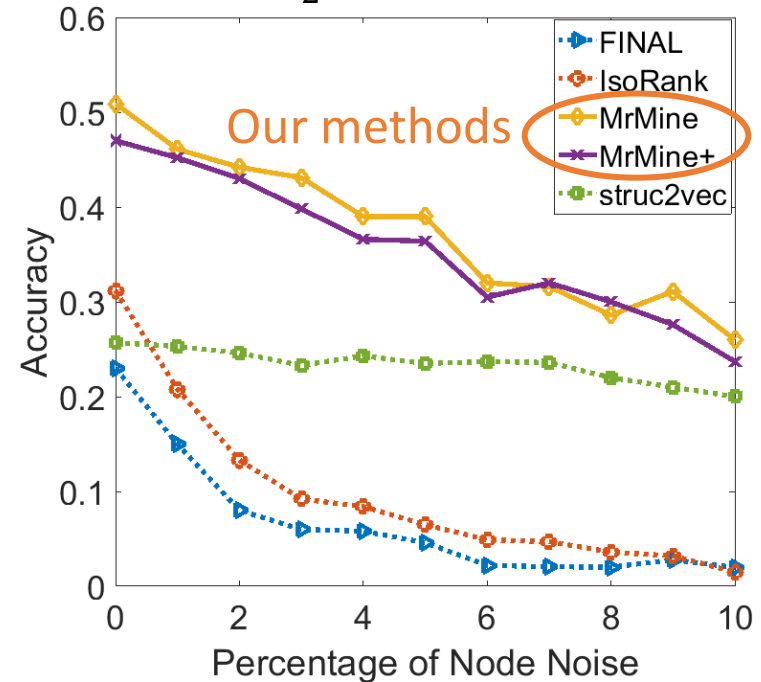
Identical nodes (same color) are supposed to be paired (close)

Network alignment

G_1 : DBLP & G_2 : DBLP
with random noises



G_1 : Douban online
& G_2 : Douban offline



- Observation:

- Our methods are competitive in DBLP data; outperform baselines in douban data.
- Our methods are more robust against noises.

Network classification

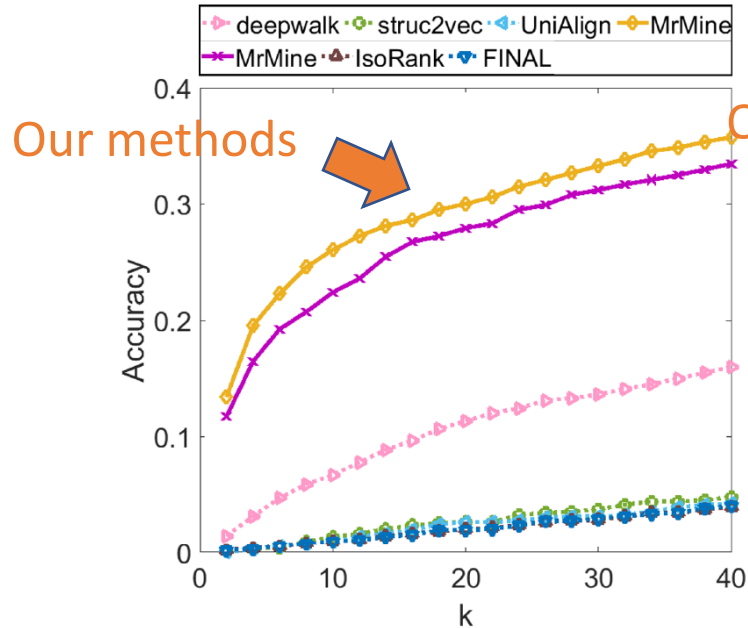
	MUTAG	PTC	PROTEINS	NCI1	NCI109
MrMine+	83.47±2.01	62.00±0.07	71.22±0.62	68.50±0.03	65.57±0.02
MrMine	82.19±1.58	55.41±2.52	70.88±0.38	66.90±0.05	64.53±0.01
WL Kernel	80.66±3.07	59.94±2.79	64.45±1.14	63.42±0.22	62.94±0.42
Deep WL Kernel	82.95±1.96	53.29±1.53	69.49±0.26	62.83±0.25	62.47±0.15
subgraph2vec	79.33±0.07	42.29±0.09	73.04±0.04	63.01±0.01	49.20±0.02

(± standard deviation)

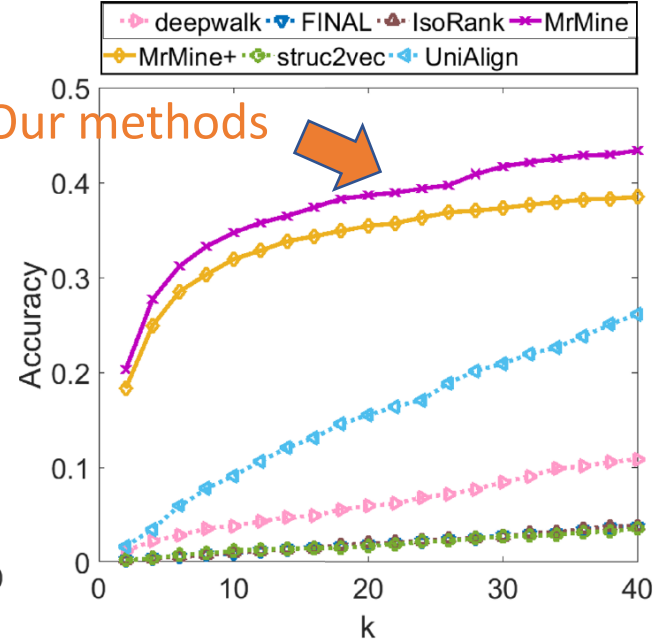
- Observation:
 - Our methods are competitive against baseline methods.
 - MrMine+ consistently outperform MrMine
 - (Cross-network relation captured by H-CRCN relation network is more effective than the basic CRCN relation network in network classification task)

Node retrieval

G_1 : DBLP & G_2 : DBLP
with random noises

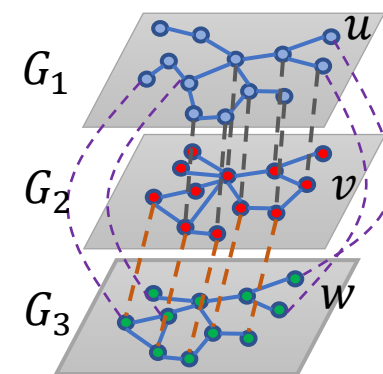
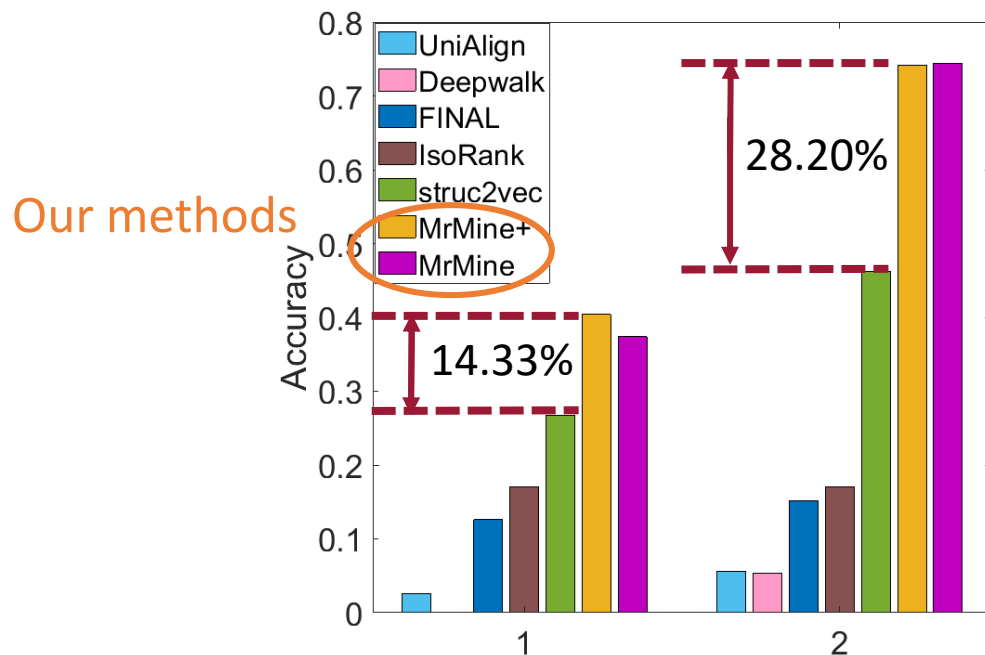


G_1 : Douban online &
 G_2 : Douban offline



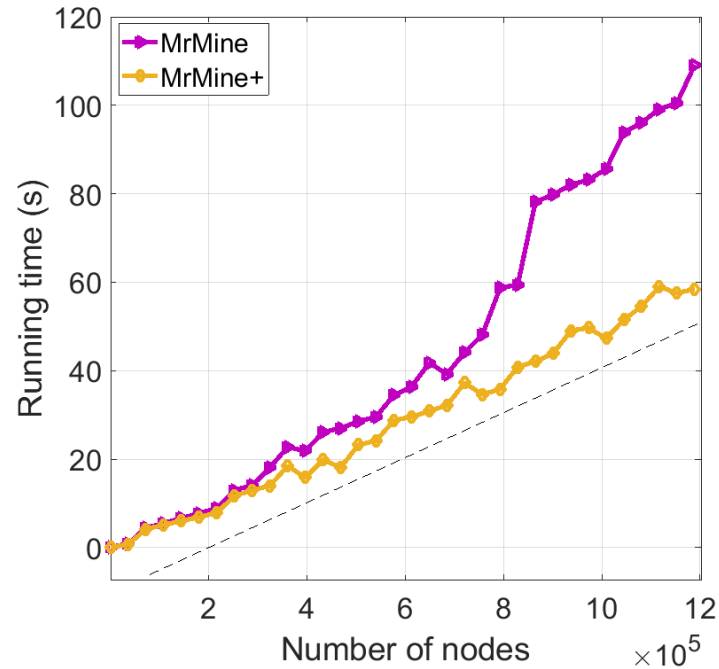
- K: For one query node of G_1 , top-k node list are retrieved from G_2 .
- Accuracy = # of hits/# of query nodes (hit: correct node appears in top-k list)
- Observation: Our methods outperform baselines.

Collective network alignment



- Dataset: Douban-online, douban-offline, douban-online with random noises
- Metrics: For each pair of three-node alignment:
 - Metric 1: successfully alignment when all nodes are aligned correctly.
 - Metric 2: successfully alignment when two of the three nodes are aligned correctly.
- Obs.: our methods outperform all baselines (embedding-based and non embedding-based)

Scalability



- Observation:

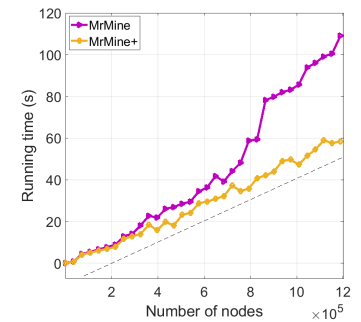
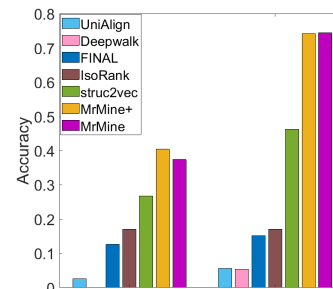
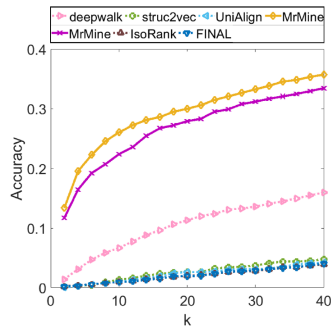
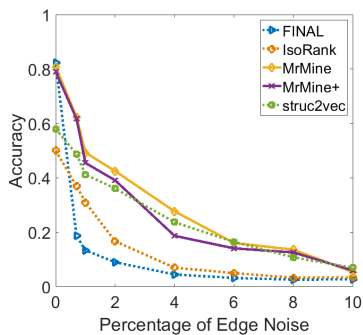
- MrMine scales super-linearly w.r.t. # of nodes of input networks
- MrMine+ scales linearly w.r.t. # of nodes of input networks

Roadmap

- Motivation
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- Proposed Solution: 'MrMine'
- Experiments
- **Conclusions**

Conclusion

- Goal: *Unsupervised* **multi-resolution multi-network** embedding.
- Solution: MrMine, MrMine+
 - Key idea 1: Cross-Resolution Cross-Network (CRCN) relation network
 - Key idea 2: WL subtree as the subgraph resolution
 - Key idea 3: Hierarchical structure of CRCN network
- Results:
 - Boost traditional network mining tasks (e.g. network classification)
 - Enable novel network mining tasks (e.g. collective network alignment)
 - Accelerated method has *linear* time complexity



Thank you!

- Q & A Session