

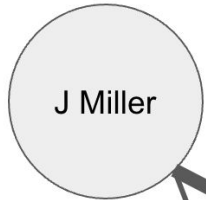
Project Status

Sammi Abida Salma

- Problem definition
- Previous work
- Proposed approach
- Evaluation Methodology

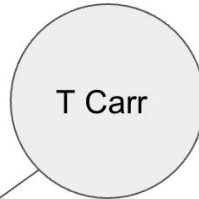
- **Problem definition**
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[publications, grants, patents,
research interest ...]



J Miller

[publications, grants, patents,
research interest ...]



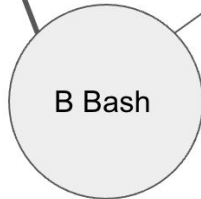
T Carr

Node ->

Researchers

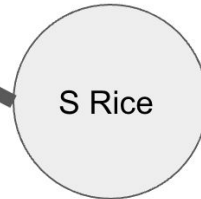
Link ->

collaborative research



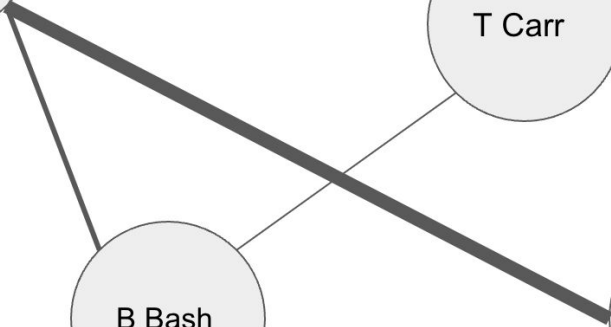
B Bash

[publications, grants, patents,
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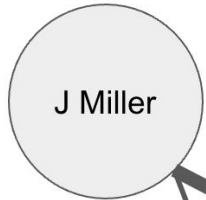


S Rice

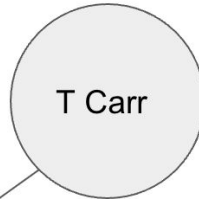
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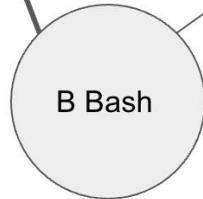


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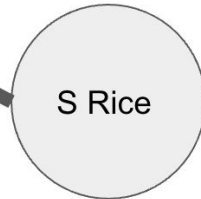


Node ->
Researchers

Link ->
collaborative research



[publications, grants, patents,
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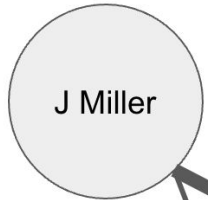


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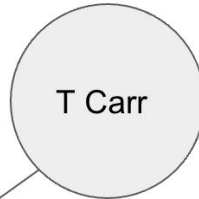
nodes : **4,884**
edges/links : **19,241**
Average degree : **7.8792**

Problem : *Infer Future Collaboration*

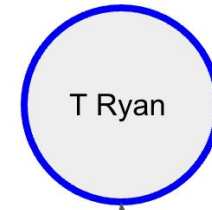
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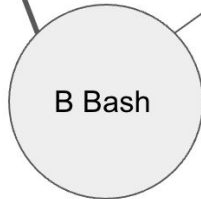
[publications, grants, patents,
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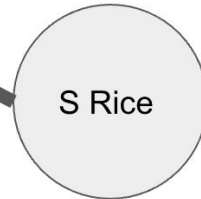
[publications, grants, patents,
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New faculty
/ Candidate

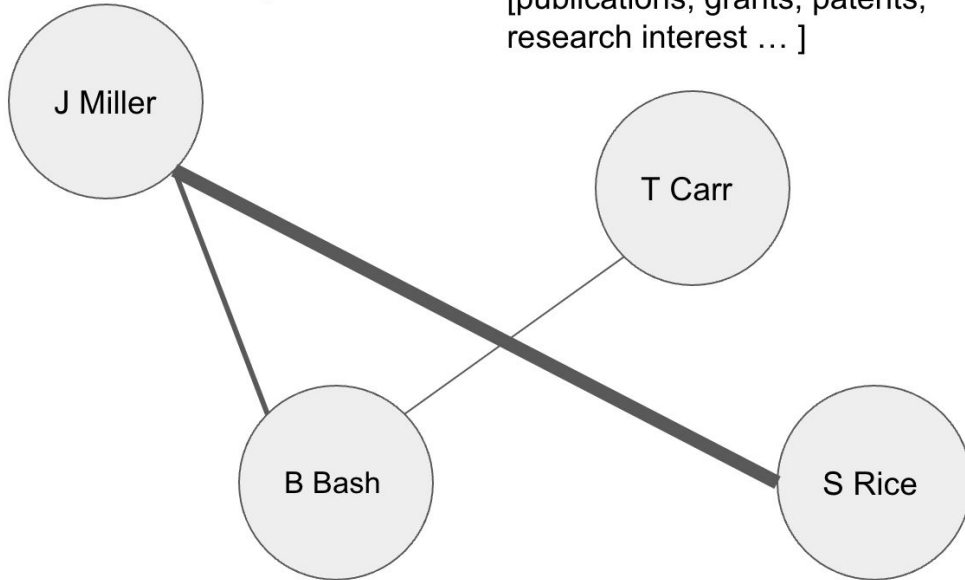


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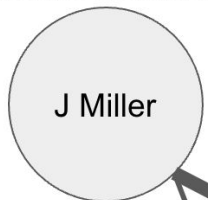
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Problem:
Predict links for "T Ryan"

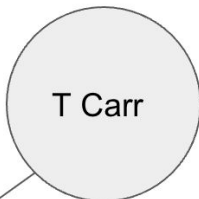


Link Prediction

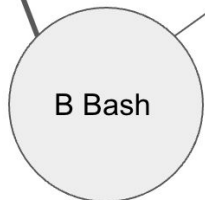
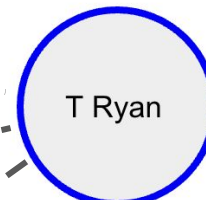
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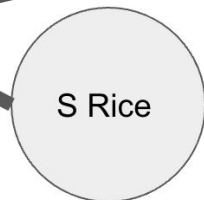
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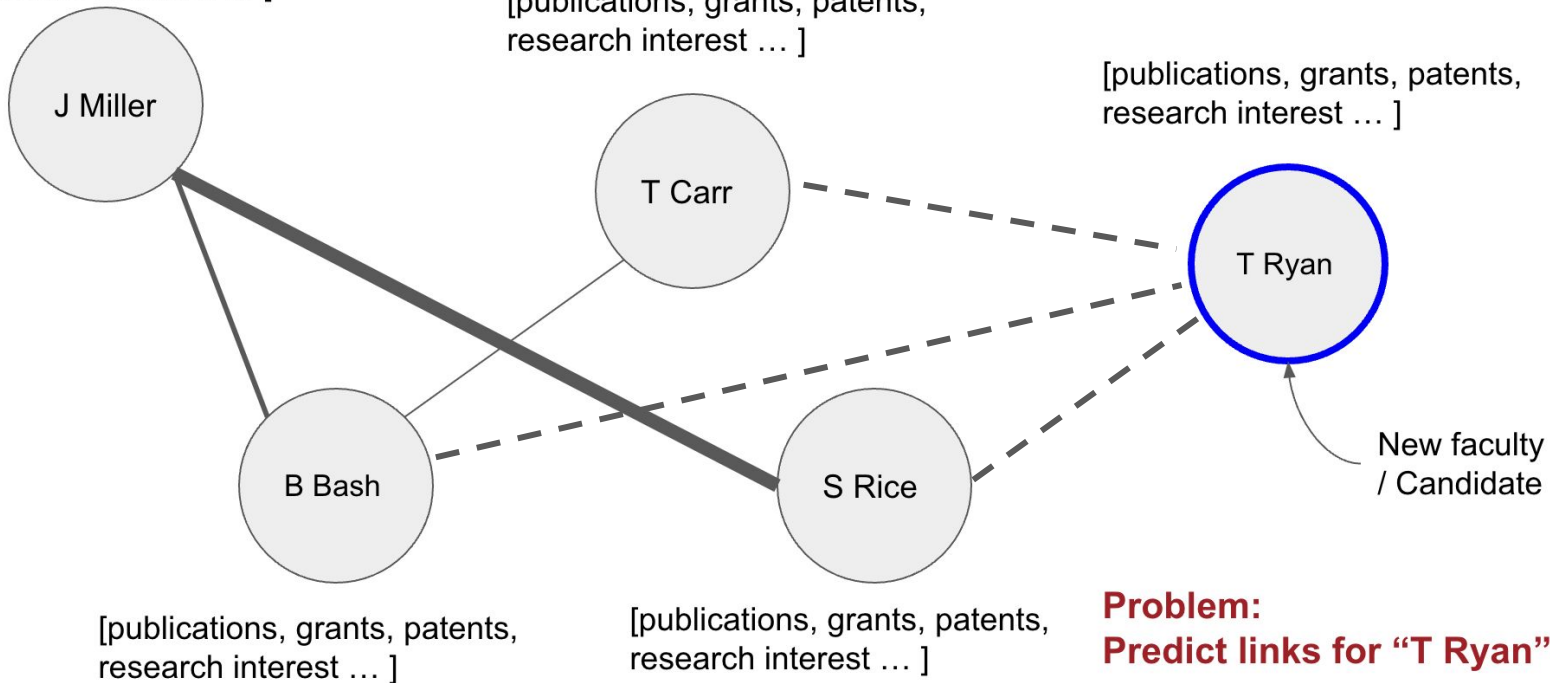
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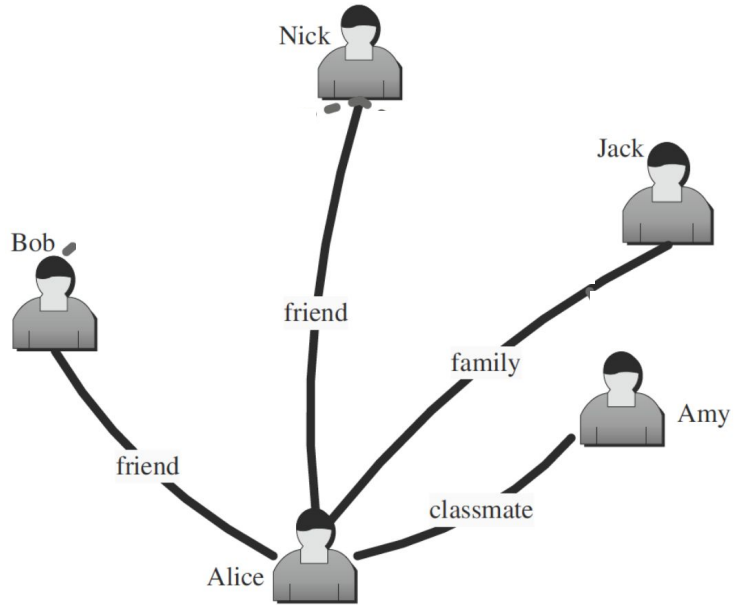
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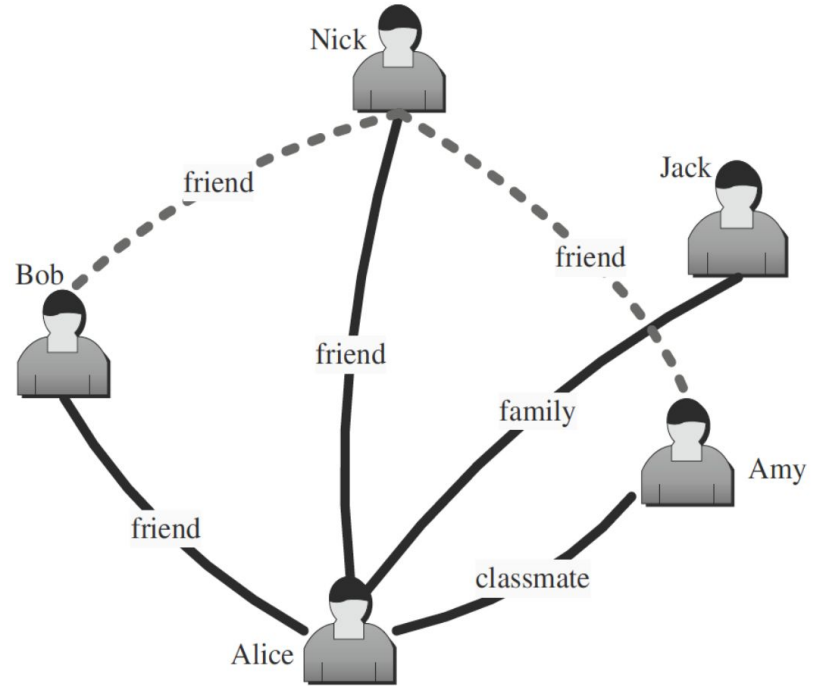
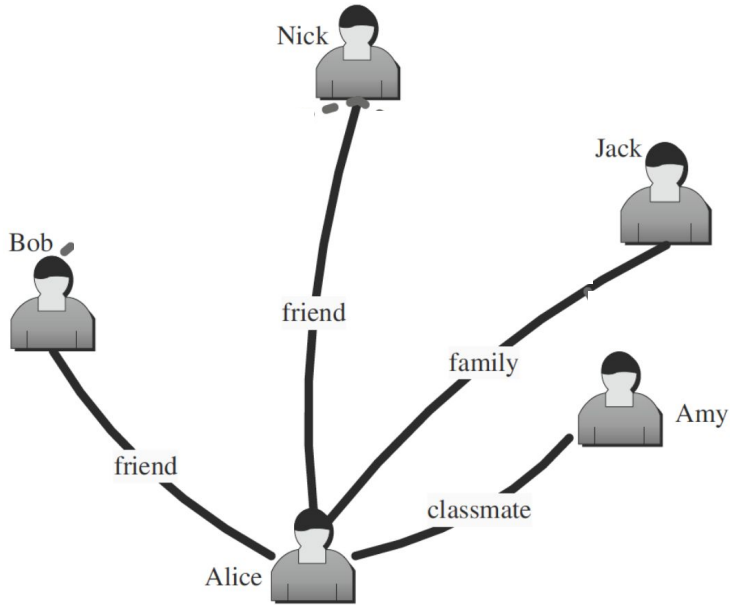
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Predict links for "T Ryan"



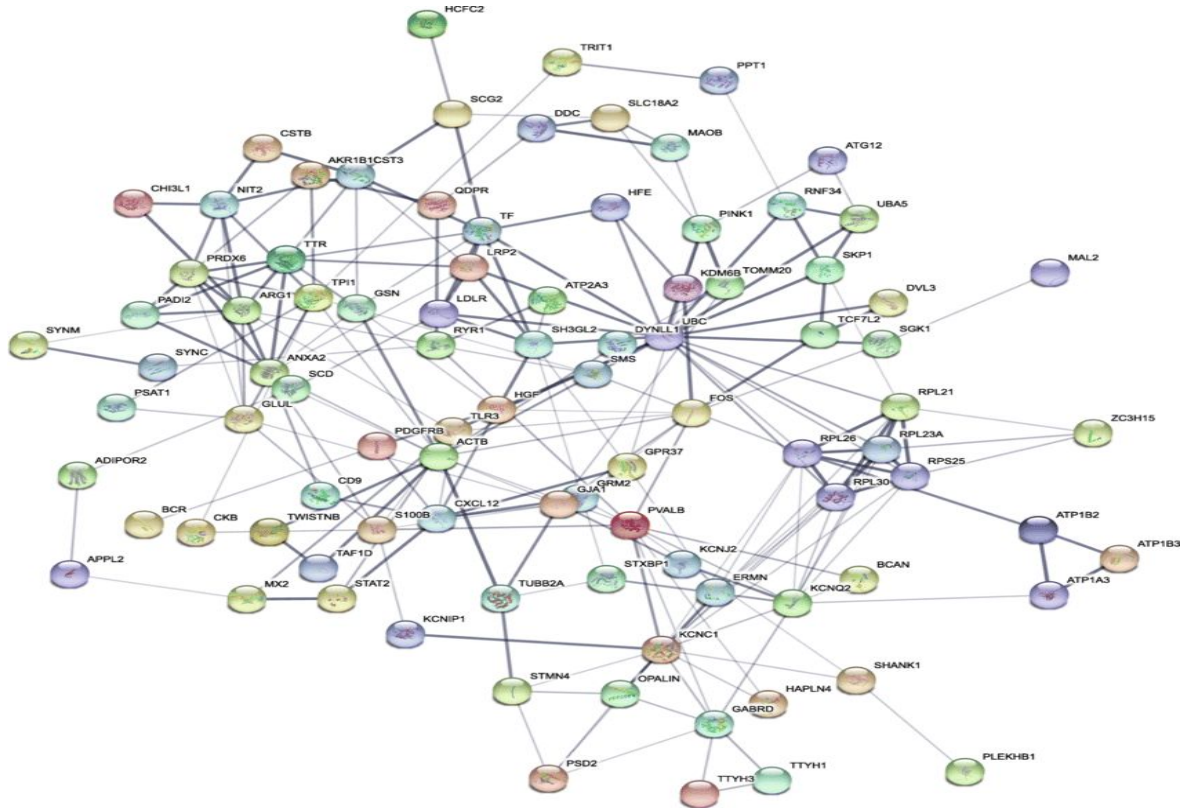
Link Prediction : *In Social Network*



Link Prediction : *Recommend Friend*



Link Prediction : *Infer Protein-protein interaction*



- Problem definition
- **Previous work**
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The generic link prediction framework

- Similarity-based approach (topological feature)
- Learning-based approach (topological feature + latent feature)

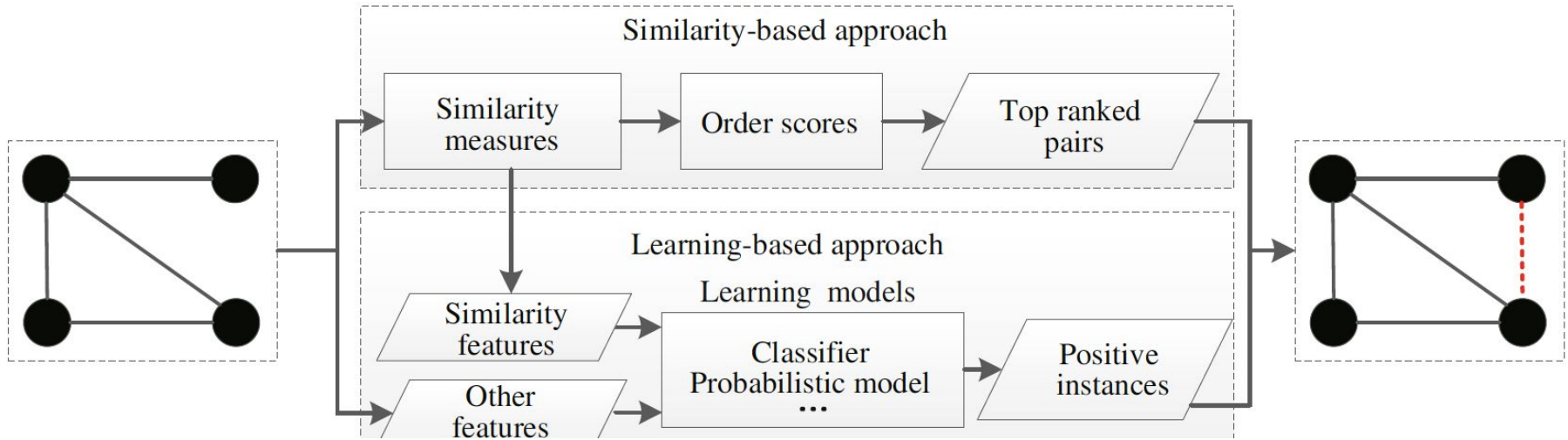


Table 1: Popular Heuristics for Link PredictionSimilarity-
based
approach

Name	Formula	Order
common neighbors	$ \Gamma(x) \cap \Gamma(y) $	first
Jaccard	$\frac{ \Gamma(x) \cap \Gamma(y) }{ \Gamma(x) \cup \Gamma(y) }$	first
preferential attachment	$ \Gamma(x) \cdot \Gamma(y) $	first
Adamic-Adar	$\sum_{z \in \Gamma(x) \cap \Gamma(y)} \frac{1}{\log \Gamma(z) }$	second
resource allocation	$\sum_{z \in \Gamma(x) \cap \Gamma(y)} \frac{1}{ \Gamma(z) }$	second
Katz	$\sum_{l=1}^{\infty} \beta^l \text{path}(x, y) = l $	high
PageRank	$q_{xy} + q_{yx}$	high
SimRank	$\gamma \frac{\sum_{a \in \Gamma(x)} \sum_{b \in \Gamma(y)} \text{score}(a, b)}{ \Gamma(x) \cdot \Gamma(y) }$	high
resistance distance	$\frac{1}{l_{xx}^+ + l_{yy}^+ - 2l_{xy}^+}$	high

Similarity-based approach

→ account (only) topological feature

- ◆ degree of nodes
- ◆ path information

→ works well (*metric common neighbor*)

- ◆ social network

→ works poorly (*metric common neighbor*)

- ◆ protein-protein interaction network

→ Limitation

- ◆ Non-universal
 - Different domains need different metrics
- ◆ Fails to predict link where
 - **similarity scores do not capture the network's latent formation mechanisms.**

Learning-based approach

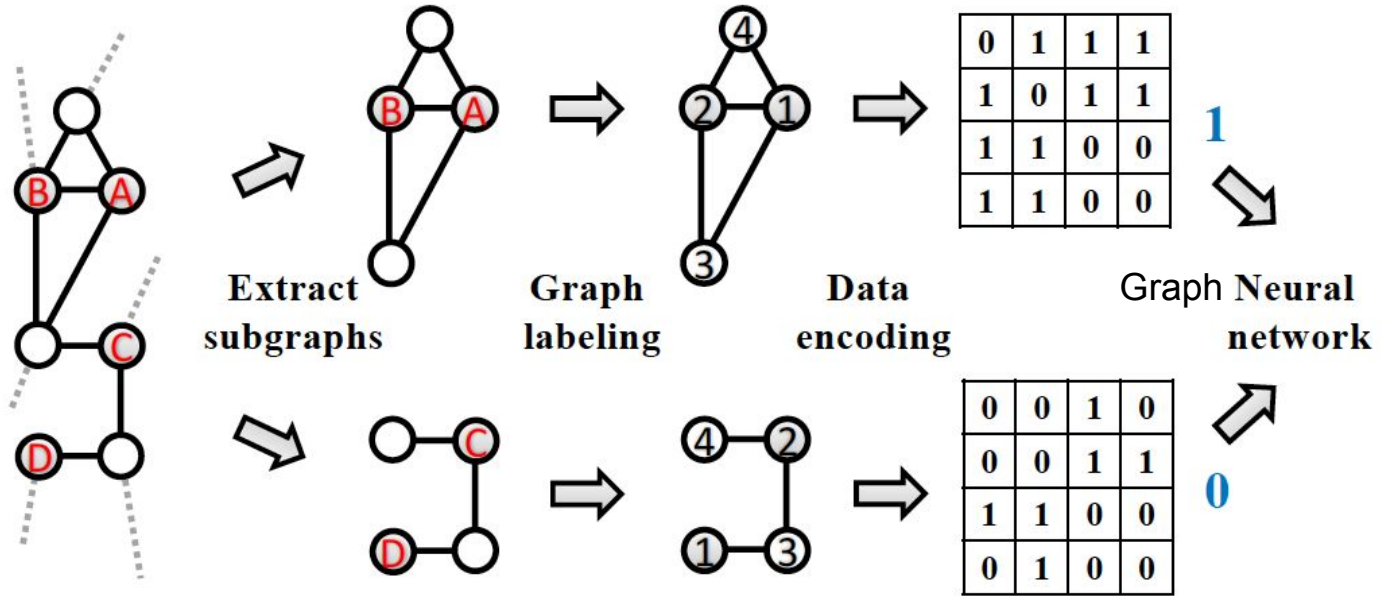
- Probabilistic Graphical Model
 - ◆ Ranking Method
 - ◆ Stochastic Block Model
- Matrix factorization
- Deep Learning
 - ◆ CNN (Convolution Neural Network)
 - ◆ GNN (Graph Neural Network)

- Problem definition
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Proposed approach

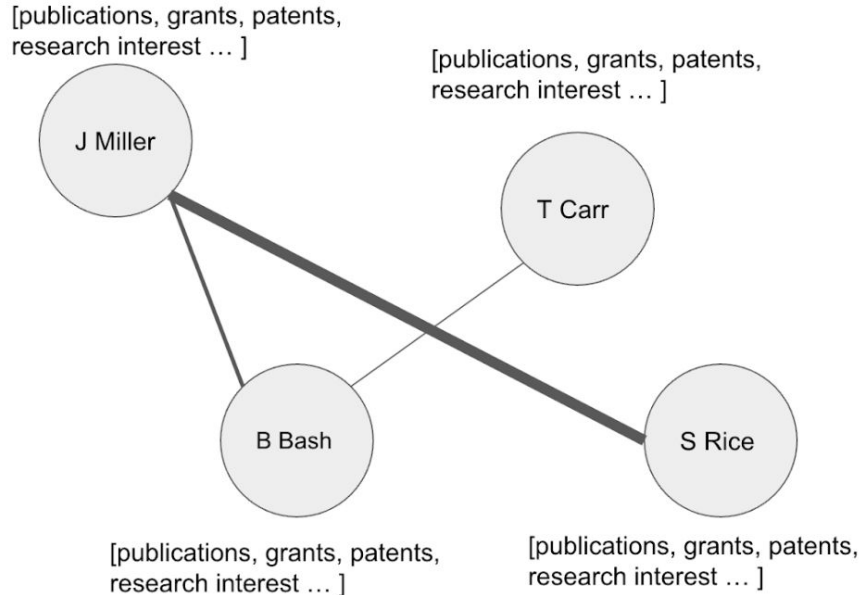
- Extract an enclosing subgraph of each target link
- Label the subgraphs using WL algorithm
- Encode the subgraph as an adjacency matrix
- Encode the features to vector
 - Publication title to a vector using word2vec transformation
- Split edges into train, validation, test set
- Feed the adjacency matrices along with feature vector to the graph neural network (GNN)
- Test and evaluate accuracy
- **Python -> pytorch-geometric**

Proposed approach



Data Encoding

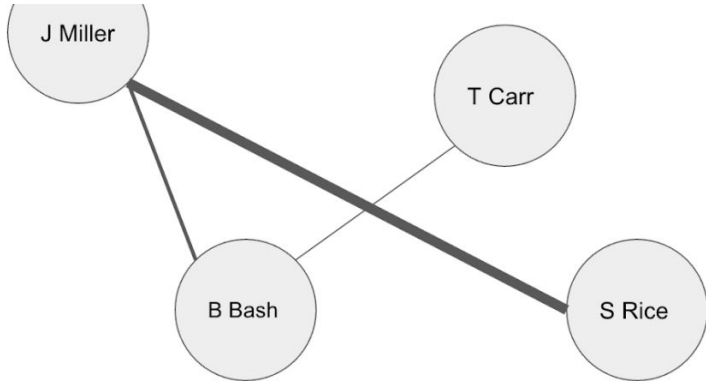
- Encodes the subgraph as an adjacency matrix
- Encode **publication titles** (text data) to feature vector using word2vec



Each node has the following attributes

- publications
- patents
- grants
- research interests
- weight
- kmapId

Fetching Data



KMAP Data

People

GET `/api/v0/people/{kmap_id}` Returns information about a person.

GET `/api/v0/people/{kmap_id}/publications` Returns list of publications of a person.

GET `/api/v0/people/{kmap_id}/grants` Returns list of Grants of a person

GET `/api/v0/people/{kmap_id}/technologies` Returns list of technologies of a person. by the kmapId

GET `/api/v0/people/{kmap_id}/patents` Returns list of patents of a person. by the kmapId

Data Cleaning

0it [00:00, ?it/s][{'kmapId': 'palmerio', 'titles': ''}]

1it [00:00, 8.39it/s][{'kmapId': 'cwesterl', 'titles': "The Judicial Common Space 1 # All Along the Watchtower: Acculturation, Fear, Anti-Latino Affect, and Immigration # Strategic Defiance in the US Courts of Appeals # Strategic Defiance in the US Courts of Appeals # Legislators in Robes?....."}]

{'kmapId': 'lumbee', 'titles': ''}

3it [00:00, 10.78it/s][{'kmapId': 'skaib', 'titles': 'Exploring Perceived Medical Student Mistreatment from Interdisciplinary Perspectives # Survey Information to Improve Competitiveness'}]

{'kmapId': 'witte', 'titles': 'Congenital chylothorax: Current evidence-based prenatal and post-natal diagnosis and management #'}]

5it [00:00, 11.21it/s][{'kmapId': 'shonad', 'titles': 'Comprehensive Lifestyle Improvement Program for Prostate Cancer (CLIPP): Protocol for a Feasibility and Exploratory Efficacy Study in Men on Androgen Deprivation Therapy (Preprint) #'}]

{'kmapId': 'dcorso', 'titles': ''}]

7it [00:00, 11.66it/s][{'kmapId': 'ghuck', 'titles': ''}]

{'kmapId': 'macmccallum', 'titles': ''}]

9it [00:00, 11.84it/s][{'kmapId': 'adriannah', 'titles': ''}]

10it [00:00, 11.58it/s]

Data Cleaning

nodes : **4,884**

edges/links : **19,241**

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Remove nodes with empty publication data

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Calculate correctness

given network $G = (V, E)$

- ❖ Take positive samples by selecting all edges $(x,y) \in E$.
- ❖ Take negative samples by randomly selecting $\alpha |E|$ pairs of $x,y \in V$ such that $(x,y) \notin E$.
- ❖ Split both positive and negative samples to
 - 90% training set
 - 10% testing set
- ❖ Train GNN with training set
- ❖ Evaluate correctness using test set

Compare Performance

- Calculate accuracy using Stochastic Block Model
- Compare correctness

Thank You!