## On Embedding Uncertain Graphs

Jiafeng Hu, Reynold Cheng, Zhipeng Huang, Yixiang Fang and Siqiang Luo

# Department of Computer Science <br> The University of Hong Kong 

jhu@cs.hku.hk

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## Graphs are everywhere



Social Network


Collaboration Network


Protein-Protein interaction Network

## Uncertainty in Graph data

- Wireless sensor networks (WSNs)
- Nodes: sensors
- Edges: wireless connectivity between sensors
- Uncertainties: probabilities of wireless connectivity



## Uncertainty in Graph data

- Protein-Protein Interaction Networks (PPI)
- Nodes: Proteins
- Edges: an interaction between proteins
- Uncertainties: probabilities of interactions between proteins derived from experimental evidence


Gabriele Cavallaro [Genome-wide analysis of eukaryotic twin $\mathrm{CX}_{9} \mathrm{C}$ proteins]

## Uncertainty in Graph data

Uncertain Graphs: each edge has an existence probability.


- Social Networks
- Traffic Networks
- Wireless Sensor Networks
- Protein-interaction Networks
- ...


## Possible World Semantics (PWS)

- Given an uncertain graph $\mathcal{G}$, a possible world of $\mathcal{G}$ is a deterministic graph $G=\left(V, E_{G} \subseteq E\right)$. Assume the existence probabilities of edges are mutually independent [VLDB'10, KDD'10].

$$
\operatorname{Pr}[G]=\prod_{e \in E_{G}} \mathbf{P}_{e} \prod_{e \in E \backslash E_{G}}\left(1-\mathbf{P}_{e}\right)
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$$


(a) An uncertain graph $\mathcal{G}$
(b) A possible world of $\mathcal{G}$

$$
\begin{aligned}
\operatorname{Pr}[G] & =\mathbf{P}_{a d} \mathbf{P}_{b d} \mathbf{P}_{b c}\left(1-\mathbf{P}_{a b}\right)\left(1-\mathbf{P}_{c d}\right) \\
& =0.1 \times 0.8 \times 0.6 \times 0.1 \times 0.8=0.00384
\end{aligned}
$$

## Uncertain Graph Mining

Tasks:

- Clustering [TKDE'13, ICDM'12]
- Classification [ICDM'09, SSDBM'14]
- k-NN queries [VLDB'10]


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- k-NN queries [VLDB'10]
- ...


## Shortcomings:

- High computational cost: expensive to compute similarities between nodes under PWS.
- Low adaptability: solutions are tailored for a particular mining task.


## Graph Embedding



Graph G

Embedding
|V|

$K=4$ dimensions

## Graph Embedding



Graph G

## Embedding

## |V|


$K=4$ dimensions

- Existing embedding solutions (for deterministic graphs): DeepWalk [KDD'14], LINE [WWW'15], node2vec [KDD'16], etc


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- Not designed for uncertain graphs


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$K=4$ dimensions

- Existing embedding solutions (for deterministic graphs): DeepWalk [KDD'14], LINE [WWW'15], node2vec [KDD'16], etc
- Not designed for uncertain graphs
- Simply remove the probabilities $\rightarrow$ poor performance


## Uncertain Graph Embedding



## URGE: UnceRtain Graph Embedding



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## Step 1: Proximity matrix S

- (second order) Expected Jaccard Similarity (EJS)
- (high order) Probabilistic Random Walk with Restart (PRWR)


## URGE: UnceRtain Graph Embedding



Step 2: Objective function

$$
\min _{\mathbf{U}, \tilde{\mathbf{U}}}\left\|\mathbf{S}-\mathbf{U} \tilde{\mathbf{U}}^{T}\right\|^{2}+\frac{\lambda}{2}\left(\|\mathbf{U}\|^{2}+\|\tilde{\mathbf{U}}\|^{2}\right)
$$

- (Input) $\mathbf{S} \in \Re^{n \times n}$ : proximity matrix;
- (Input) $\lambda$ controls the weight of the regularization term.
- (Output) matrices $\mathbf{U} \in \Re^{n \times K}$ and $\tilde{\mathbf{U}} \in \Re^{n \times K}$


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$$

Negative sampling + asynchronous stochastic gradient algorithm (ASGD)

## How to compute the proximity matrix efficiently?

## Second-order Proximity: Expected Jaccard Similarity

Jaccard similarity between node $u$ and $v$ on a deterministic graph $G$ :

$$
S_{u v}^{J}=\frac{\left|N_{G}(u) \cap N_{G}(v)\right|}{\left|N_{G}(u) \cup N_{G}(v)\right|}
$$

$N_{G}(x)$ : the neighbor set of node $x$ on $G$.

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## Expected Jaccard Similarity (EJS):

$$
S_{u v}^{\mathrm{EJS}}=\sum_{G \in \Omega(\mathcal{G})}\left(S_{u v}^{\mathrm{J}}\right)_{G} \operatorname{Pr}[G]
$$

$\Omega(\mathcal{G})$ : the set of all possible worlds of $\mathcal{G}$.

## Computation of EJS

## Lemma (A. Stuart, 1998; Z. Zhou, ICDM'13)

Given two nodes $u$ and $v$ of $\mathcal{G}$, let $X_{u v}=\left|N_{G}(u) \cap N_{G}(v)\right|$ and $Y_{u v}=\left|N_{G}(u) \cup N_{G}(v)\right|$, where $G$ is a possible world of $\mathcal{G}$. Then,

$$
\mathbf{S}_{u v}^{\mathrm{EJS}}=E\left[\frac{X_{u v}}{Y_{u v}}\right] \approx \frac{E\left[X_{u v}\right]}{E\left[Y_{u v}\right]}-\frac{\operatorname{Cov}\left(X_{u v}, Y_{u v}\right)}{E\left[Y_{u v}\right]^{2}}+\frac{E\left[X_{u v}\right] \operatorname{Var}\left(Y_{u v}\right)}{E\left[Y_{u v}\right]^{3}}
$$

- Z. Zou et al. study the problem of computing EJS between a pair of nodes $\rightarrow \mathcal{O}(d)$
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- (Basic method) for each node $u$ on $\mathcal{G}$, enumerate all 2-hop neighbors, $v$, and compute $\mathbf{S}_{u v}^{\mathrm{EJS}} \rightarrow \mathcal{O}\left(n d^{2} * d\right)$
- Z. Zou et al. study the problem of computing EJS between a pair of nodes $\rightarrow \mathcal{O}(d)$
- (Basic method) for each node $u$ on $\mathcal{G}$, enumerate all 2-hop neighbors, $v$, and compute $\mathbf{S}_{u v}^{\mathrm{EJS}} \rightarrow \mathcal{O}\left(n d^{2} * d\right)$
- (Our solution) compute the EJS for all pair of nodes incrementally (i.e., the whole EJS matrix $\left.\mathbf{S}^{\text {EJS }}\right) \rightarrow \mathcal{O}\left(n d^{2}\right)$


## High-order Proximity: Probabilistic Random Walk with Restart

- Random walk (transition procedure) on uncertain graphs (for node $u$ ) [VLDB'10]:
(1) generate a possible world $G$ for $u$;
(2) walk to a neighbor uniformly at random if there exists any neighbors of $u$ in $G$, otherwise stay at $u$.


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- Probabilistic Transition Matrix, PTM [VLDB'10, IS'15]:

$$
\mathbf{W}_{u v}= \begin{cases}\prod_{(u, q) \in E}\left(1-\mathbf{P}_{u q}\right), & u=v \\ \sum_{G \in \Omega(\mathcal{G}) \wedge(u, v) \in E_{G}} \frac{1}{d_{u}^{G}} \operatorname{Pr}[G], & u \neq v\end{cases}
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- Probabilistic Random Walk with Restart, PRWR

$$
\mathbf{S}^{\mathrm{PRWR}}=(1-\alpha) \mathbf{S}^{\mathrm{PRWR}} \mathbf{W}+\alpha \mathbf{I}, \quad(\mathbf{I} \text { is an identity matrix })
$$

## Computation of PTM and PRWR

Computation of PTM:

- (Basic method) an existing algorithm $\rightarrow \mathcal{O}\left(n d^{3}\right)$ [IS'15]
- (Our method) further improvement, a hash-based method $\rightarrow \mathcal{O}\left(h n d^{2}\right)$, $h \ll d$.

Computation of PRWR:

- Monte Carlo method $\rightarrow \mathcal{O}\left(n R \frac{1}{\alpha}\right)$
- $R$ : number of walkers;
- $1 / \alpha$ : expected length of random paths


## Experiments

Tasks: node clustering, node classification and $k$-NN search

Algorithms:

- Our algorithms:
- URGE EJs : URGE algorithm based on EJS
- URGE ${ }_{\text {prur }}$ : URGE algorithm based on PRWR
- Existing embedding algorithms:
- DeepWalk
- LINE
- node2vec ${ }_{p}^{q}$ ( node2vec $_{0.25}^{0.25}$ and node2vec ${ }_{4}^{1}$ )
- Existing non-embedding algorithms:
- MCL (for deterministic graph clustering)
- pKwikCluster (for uncertain graph clustering)
- uBayes ${ }^{+}$(for uncertain graph classification)
- MostProbPath (for uncertain graph $k$-NN)


## Task 1: Clustering

Dataset: 4 real uncertain Protein-Protein Interaction (PPI) networks ${ }^{1}$ Ground truth: the complex-memberships lists from the MIPS database

| Name | \#Nodes | \#Edges | Avg. Prob. |
| :--- | :---: | :---: | :---: |
| Krogan_core | 2,708 | 7,123 | 0.68 |
| Krogan_extend | 3,672 | 14,317 | 0.42 |
| Collins | 1,622 | 9,074 | 0.78 |
| Gavin | 1,855 | 7,669 | 0.36 |

Table: Statistics of the PPI networks.

[^0]
## Task 1: Clustering (Cont'd)

- Metric: F1 score based on the confusion matrix (true positive, false positive, true negative and false negative)
- Hierarchical clustering in vector space (embedding-based algorithms)


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| Algorithm | Krogan_core | Krogan_extend | Collins | Gavin |
| :--- | :---: | :---: | :---: | :---: |
| DeepWalk | 39.21 | 33.43 | 55.15 | 47.33 |
| LINE | 38.73 | 33.07 | 48.28 | 44.14 |
| node2vec $_{4}^{1}$ | 39.30 | 33.06 | 52.42 | 46.17 |
| node2vec 0.25 $_{0.25}$ | 38.96 | 33.75 | 53.23 | 46.13 |
| MCL | 36.01 | 30.83 | 57.55 | 47.84 |
| pKwikCluster | 16.94 | 12.88 | 24.59 | 5.65 |
| URGE EJS $^{\text {URGE }}$ PRWR | 38.39 | 30.08 | 55.61 | 54.54 |
| Table : F1 scores (\%) for clustering tasks. |  |  |  |  |

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| URGE ${ }_{\text {PRWR }}$ | 44.86 | 35.58 | 58.16 | 52.59 |

## Task 2: Classification

## Dataset:

- DBLP (co-authorship network): 45,583 edges, 14,376 papers, 4 classes
- Cora (citation network): 8,365 edges and 2,708 papers, 7 classes *Use the method proposed by P. Boldi et al. [VLDB'12] to do obfuscation.

Other setting:

- $k$ nearest neighbor classifiers (embedding-based algorithms).
- Varying training ratio ( $T_{R}$ ) from $20 \%$ to $80 \%$
- Metric: Micro-F1, Macro-F1.


## Task 2: Classification (Cont'd)

| Metric | Algorithm | 20\% | 40\% | 60\% | 80\% |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Micro-F1(\%) | DeepWalk | 40.81 | 49.71 | 54.33 | 57.27 |
|  | LINE | 40.87 | 47.96 | 53.26 | 56.52 |
|  | node2vec ${ }_{4}^{1}$ | 41.25 | 51.29 | 55.67 | 59.27 |
|  | node2vec ${ }_{0}^{0.25}$ | 40.16 | 49.33 | 53.53 | 56.98 |
|  | uBayes ${ }^{+}$ | 32.43 | 45.13 | 44.57 | 57.44 |
|  | URGE ${ }_{\text {EJS }}$ | 58.00 | 63.31 | 66.36 | 69.45 |
|  | URGE ${ }_{\text {PRWR }}$ | 52.16 | 58.08 | 61.12 | 63.97 |
| Macro-F1(\%) | DeepWalk | 38.12 | 48.22 | 52.95 | 55.98 |
|  | LINE | 39.02 | 46.37 | 51.70 | 55.08 |
|  | node2vec ${ }_{4}^{1}$ | 39.42 | 49.21 | 53.93 | 57.69 |
|  | node2vec ${ }_{0.25}^{0.25}$ | 38.11 | 47.64 | 51.93 | 55.37 |
|  | uBayes ${ }^{+}$ | 31.07 | 42.02 | 45.03 | 55.33 |
|  | URGE ${ }_{\text {EJS }}$ | 55.48 | 61.45 | 64.49 | 67.50 |
|  | URGE ${ }_{\text {PRWR }}$ | 49.86 | 56.41 | 59.64 | 62.42 |

Table: Results of classification on DBLP under different training ratio(\%).

## Task 2: Classification (Cont'd)

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## Efficiency

Datasets: DBLP, Cora, and BlogCatalog (relationships of the bloggers, 10K nodes and 665 K edges)
Algorithms:

- EJS: BasicEJS vs FastEJS (6+ times faster)
- PTM: BasicPTM vs DP hash (20+ times faster)

(c) EJS

(d) PTM


## Conclusion

- Formulate the problem of uncertain graph embedding.
- Propose URGE, a proximity preserved embedding method for uncertain graphs.
- Develop efficient algorithms for two kinds of proximities (EJS and PRWR).
- Detailed evaluation on various tasks demonstrates the efficiency and effectiveness of the URGE solution.


## Our team

THE UNIVERSITY OF HONG KONG



Dr. Reynold Cheng (ckcheng@cs.hku.hk)


Jiafeng


Zhipeng


Yixiang


Siqiang

# Thanks! 

## Q\&A


[^0]:    ${ }^{1}$ http://www.nature.com/nmeth/journal/v9/n5/full/nmeth. 1938.html

