

Chapter 10

Link Prediction

Summary

- Predicting Network Evolution
- Unsupervised approaches
- Supervised approaches
- Evaluation

Reading

- Liben-Nowel & Kleinberg paper



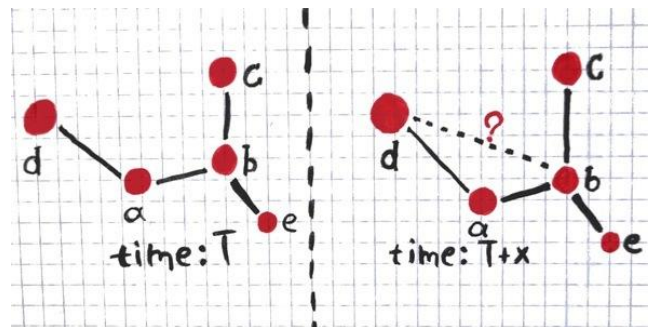
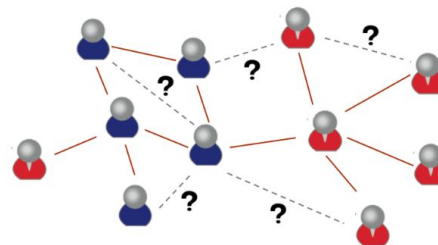
Link Prediction

Goal

Understanding how networks evolve

Problem definition

Given a snapshot of a network at time t ,
(accurately) predict the edges that will appear in
the network during the interval $(t, t+1)$



Examples of uses of

Link Prediction



Monitor terrorist networks – deducing possible interactions/missing links between terrorists (without direct evidence)



Suggest interactions or collaborations that haven't yet been exploited within an organization

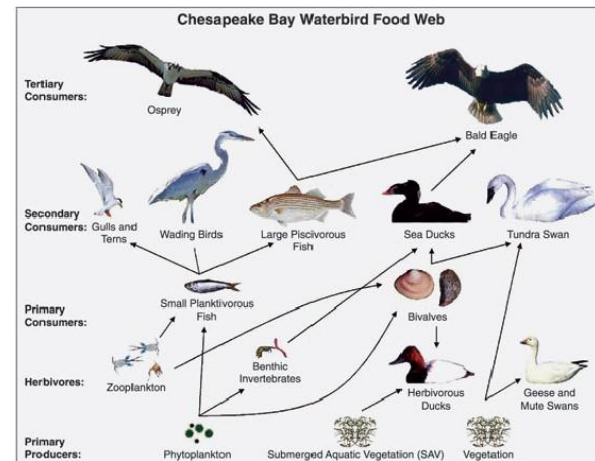


Friendship prediction (i.e., Facebook)

Link Prediction

Link prediction is used to predict **future possible links** in the network (e.g., Facebook).

Or, it can be used to predict **missing links** due to incomplete data (e.g., Food-webs)



facebook

tinder

amazon

PLOS ONE

RESEARCH ARTICLE

Link Prediction in Criminal Networks: A Tool for Criminal Intelligence Analysis

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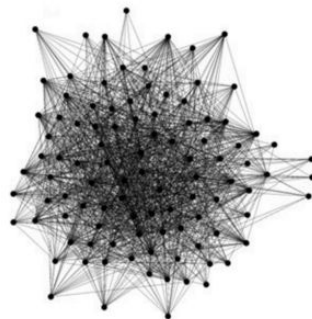
Link Prediction

Task Complexity

1. Given a graph $G = (V, E)$ the set of possible edges to be predicted is $O(|V|^2)$;
2. Real networks tend to be sparse



False Positive prediction issue!!
(i.e., we are likely to predict edges that will never appear)



Dense Graph



Sparse Graph

Concretizing an Intuition...

Scientists who are close in the network
(i.e., have common colleagues)

→ will likely collaborate in the future

Goal:

- make this intuitive notion precise and understand which measures of "proximity" leads to accurate predictions

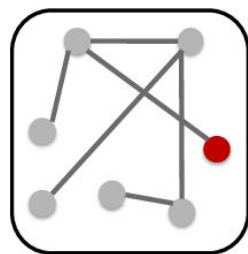


Link Prediction

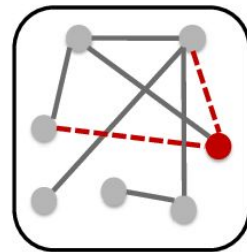
Workflow

1. Consider as input a graph G at time t
2. Consider all the possible pairs of nodes (u,v)
3. Compute a link formation scores:
 $\text{score}(u,v)$
4. Build a list of all possible edges ordered by scores (from highest to lowest)
5. Verify, following that ordering, the predictions on the graph at time $t+1$

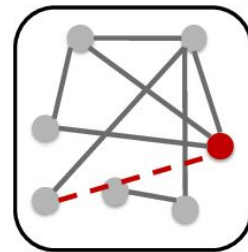
score is a measure of *proximity*



Time t



Time $t+a$



Time $t+2a$

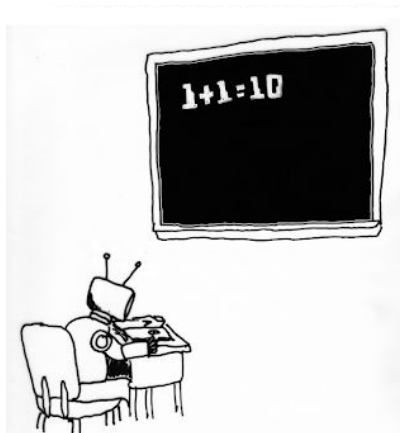
Link Prediction

Approaches

Unsupervised

defining a set of **proximity measures** unrelated to the particular network

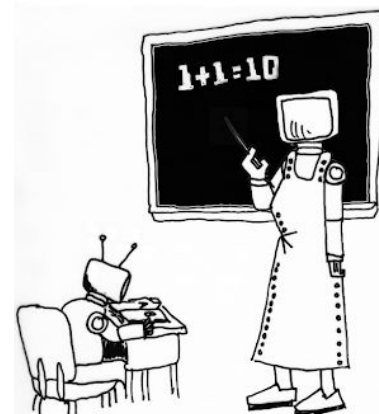
UNSUPERVISED MACHINE LEARNING



Supervised

extracting knowledge from the network in order to improve prediction accuracy

SUPERVISED MACHINE LEARNING



Unsupervised Link Prediction



Unsupervised Link Prediction

Unsupervised measurements rely on different structural properties of networks

Neighborhood measures

- Common Neighbors, Adamic Adar, Jaccard, Preferential Attachment

Path-based measures

- Graph distance, Katz

Ranking

- Sim Rank, Hitting time, Page Rank

Liben-Nowell, David, and Jon Kleinberg. "The link-prediction problem for social networks." *Journal of the American society for information science and technology* 58.7 (2007): 1019-1031.

Neighborhood measures

How many friends we have to share in order to become friends?

Common Neighbors:

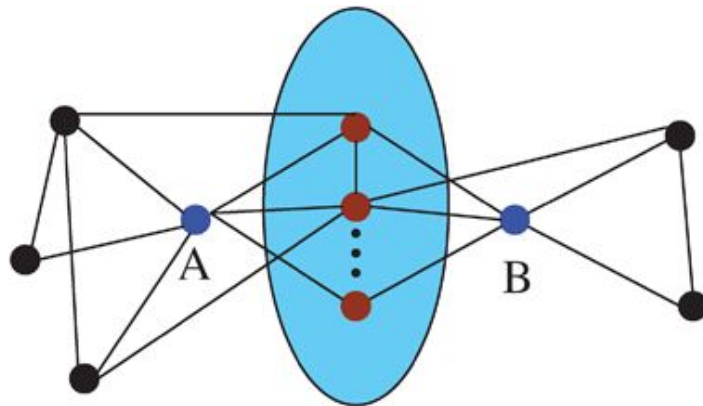
the more friends we share,
the more likely we will become friends

$$\text{score}(u, v) = |\Gamma(u) \cap \Gamma(v)|$$

Jaccard:

the more similar our friends circles are,
the more likely we will become friends

$$\text{score}(u, v) = \frac{|\Gamma(u) \cap \Gamma(v)|}{|\Gamma(u) \cup \Gamma(v)|}$$



Neighborhood measures

How many friends we have to share in order to become friends?

Adamic Adar:

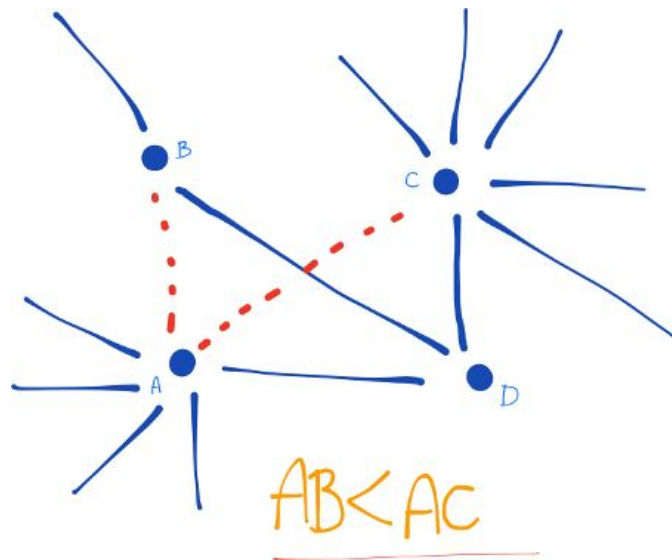
the more selective our mutual friends are,
the more likely we will become friends

$$\text{score}(u, v) = \sum_{z \in \Gamma(u) \cap \Gamma(v)} \frac{1}{\log(|\Gamma(z)|)}$$

Preferential Attachment:

the more friends we have,
the more likely we will become friends

$$\text{score}(u, v) = |\Gamma(u)| * |\Gamma(v)|$$



Path-based measures

How distant are we?

Graph Distance:

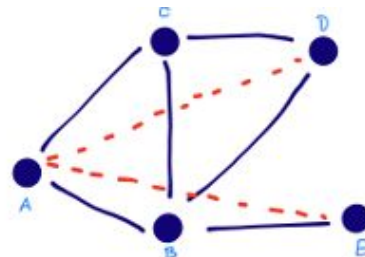
(negated) length of the shortest path between two nodes

Katz:

weighted sum over all the paths between two nodes

$$\text{score}(u, v) = \sum_{l=1}^{\infty} \beta^l |\text{paths}_{u,v}^{(l)}|$$

where: $\text{paths}_{u,v}^{(l)} = \{\text{paths of length exactly } l \text{ from } u \text{ to } v\}$



of Hops

$$\text{Path}_{A,D}^2 = 2 \quad \text{Path}_{A,D}^3 = 2$$
$$S = \frac{1}{2} \cdot 2 + \frac{1}{4} \cdot 2 + \dots$$

Damping Factor

$$\text{Path}_{A,E}^2 = 1 \quad \text{Path}_{A,E}^3 = 1$$
$$S = \frac{1}{2} \cdot 1 + \frac{1}{4} \cdot 1 + \dots$$

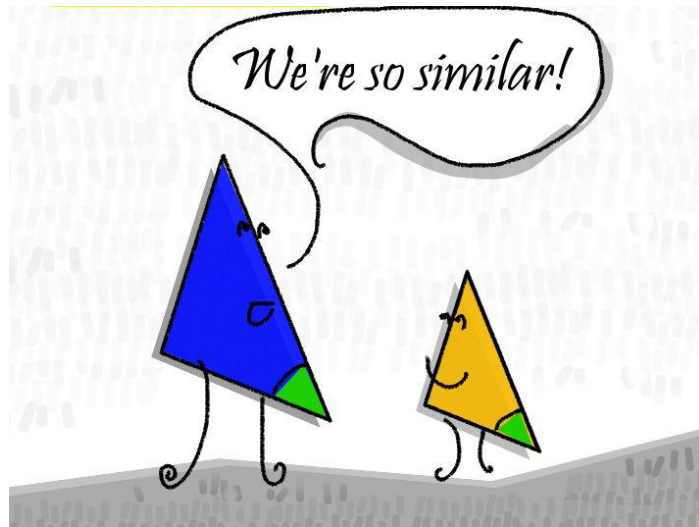
Ranking

How similar are we?

SimRank:

two nodes are *similar* to the extent that their neighborhoods are *similar*

$$\text{similarity}(u, v) = \gamma * \frac{\sum_{s \in r(u)} \sum_{n \in \Gamma(v)} \text{similarity}(a, b)}{|\Gamma(u)| * |\Gamma(v)|}$$
$$\text{score}(u, v) = \text{similarity}(u, v)$$



Unsupervised Link Prediction

Limits

- Different kinds of networks are described by the same general closed formula
- An average overall performance between 6% and 12%

Measure comparison

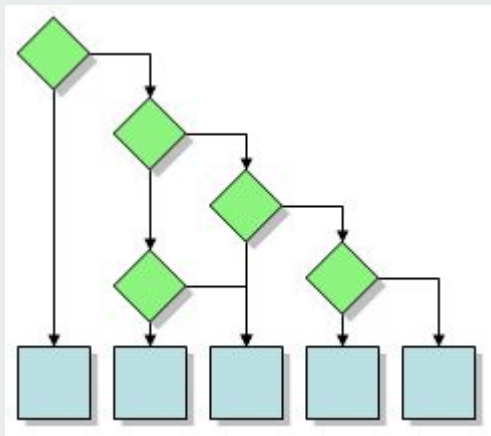
- No single winner
- Almost all predictors outperform the **random predictor**
⇒ there is useful information in network topology



Supervised Link Prediction



Supervised Link Prediction



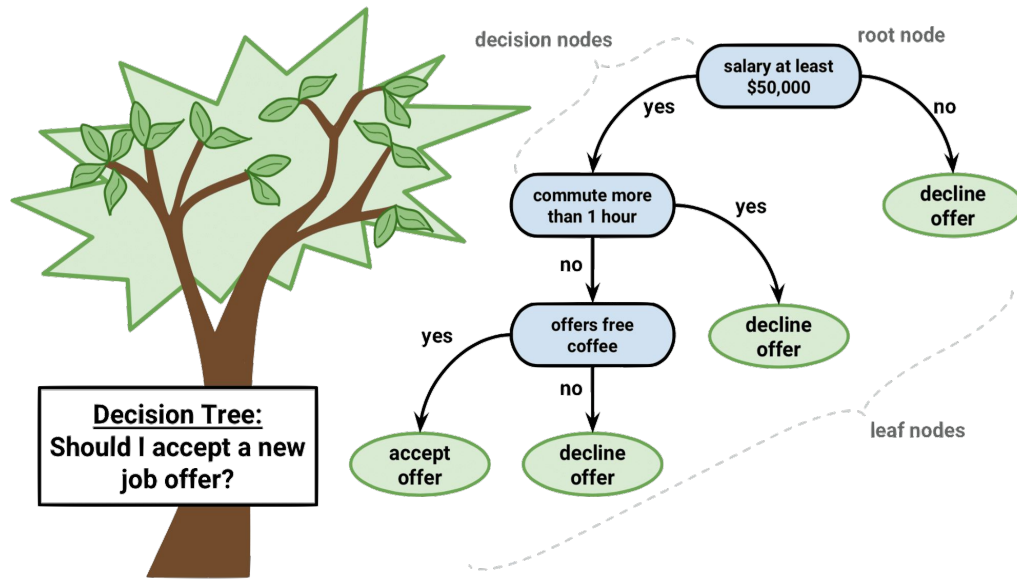
The process is now organized in 4 steps:

1. Split the data in train/test
2. Learning a model on the train
3. Use the model for prediction
4. Compare the prediction with the test

A natural way to do it:
build a “*classifier*” over a set of *network features*.

Staking Unsupervised Scores

Learn a Classifier (i.e., a Decision Tree) over unsupervised LP scores to generalize the assumption they made on the network growth model



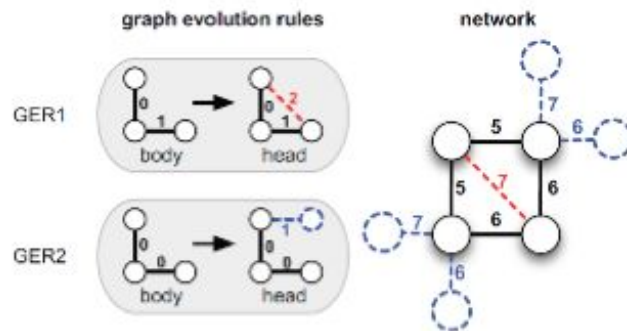
Supervised Link Prediction

Frequent Pattern Mining

GERM

Evolution rules can be extracted from the network history and used to identify/predict recurrent patterns

- e.g., generalization of triadic closure



Berlingiero, Michele, et al. "Mining graph evolution rules." *joint European conference on machine learning and knowledge discovery in databases* (2009).

Supervised Link Prediction

Network Embedding

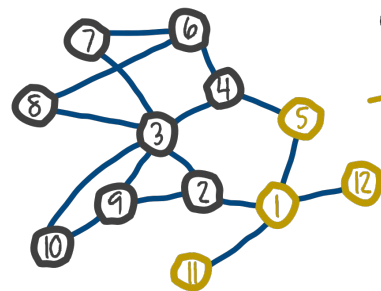
Idea

Graphs can be *mapped* into vector spaces

- Node/edge similarity scores can be used to define metric spaces
- Metric spaces enable a more natural application of approaches from DM/ML

NB: Different “mappings” facilitate the solution of different classes of problems

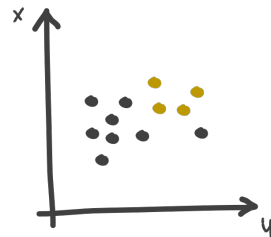
from a graph representation ...



embedding
algorithm



to real vector representation



Limits

- No Free Lunch
- Model construction is often complex and, usually, more time/resource *demanding* than directly applying unsupervised scores.

Results: Higher performances w.r.t. unsupervised approaches



Embedding is not The Answer,
only a different way to reason on graphs...



Evaluation

Given a predictor p is there a way to decide if it is a “good” one?

First Step:

verify that p outperforms the random predictor.

Random Predictor

each edge has the same probability to appear in the network

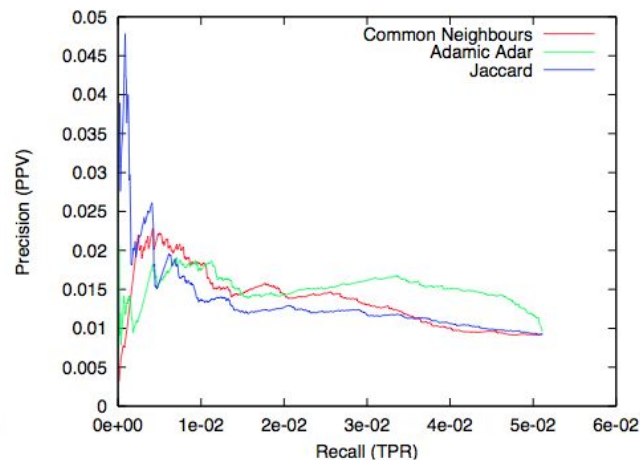
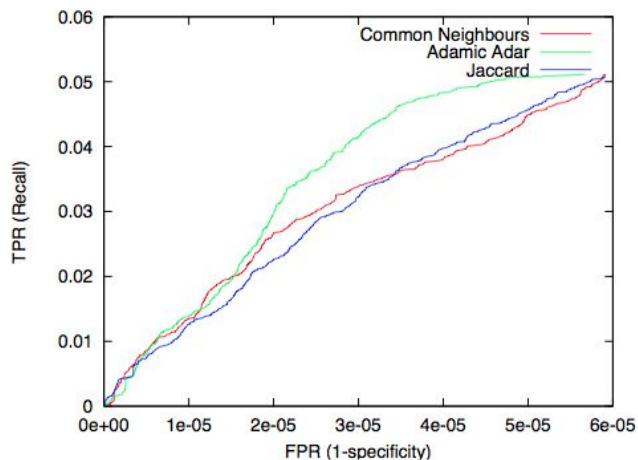
$$\text{performance}(p) = \frac{TP}{TP + FP}$$

$$\text{ratio} = \frac{\text{performance}(p)}{\text{performance}(\text{prandom})} = \frac{\text{performance}(p)}{\frac{\mathbb{V}(-(\sqrt{V}-1))}{2} - |E_{\text{old}}|}$$

if ratio > 1 then p is meaningful

Evaluation: Comparing Predictors

We need to analyze
either the
performances ratio,
ROC and/or Precision
Recall curve.



Evaluation: ROC and PR curve

Precision Vs. Recall

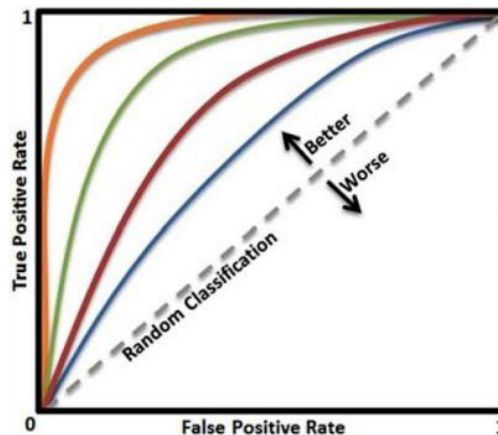
- Precision: $PPV = TP / (TP + FP)$
- Recall: $TPR = TP / (TP + FN)$

ROC (Receiver operating characteristic)

- 1-Specificity: $FPR = FP / (FP + TN)$
- Recall: $TPR = TP / (TP + FN)$

Note:

- ROC and PR spaces are isomorphic (the use of ROC is more widespread)
- Numerical comparison can be done using the AUROC (area under the ROC curve)



	p'	n'
p	TP	FN
n	FP	TN

Confusion Matrix

Link Prediction

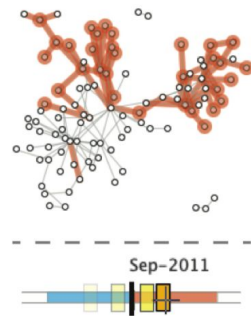
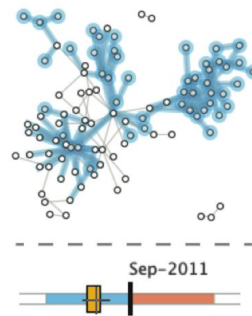
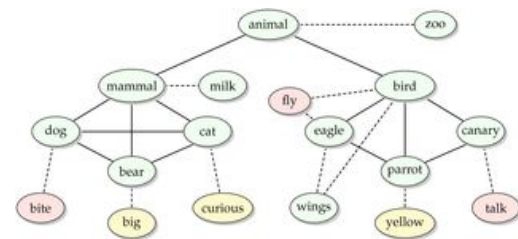
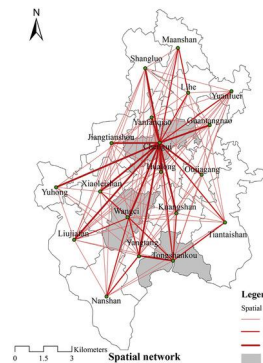
Something more...

Accuracy could be improved extending simple models with more complex (even semantic) informations:

- Link strength
- Geographical information
- ...

Link Prediction needs to be revised while in some scenarios:

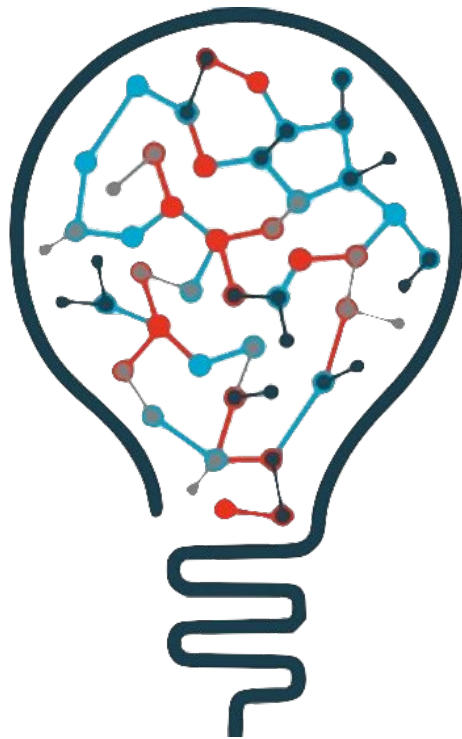
- **Dynamic Networks**
- Multiplex networks
- ...



Key Messages

Predict new links that will arise in a network is **not easy**:

1. Networks are, usually, **sparse**
2. **Cold Start Problem**
 - What if I don't have enough information?
 - *Can I predict bridges?*
3. Huge **False Positive** prediction
 - *Bridges !?!*
4. Simple approaches are “**too simple**”
5. Complex approaches are **costly**



Case Study:

Interaction Prediction in Dynamic Networks

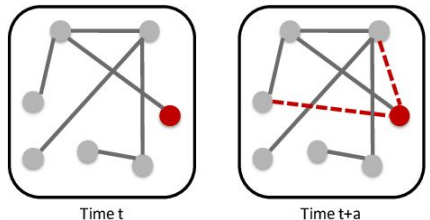


Case Study

Interaction Prediction

Link Prediction goal:

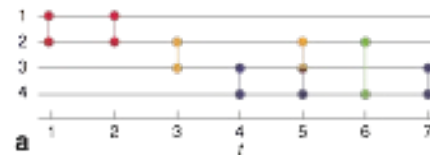
Predict ties that are not present in actual network configuration.



Ties are persistent structures that once appeared cannot disappear (i.e., friendship...)

Interaction Prediction goal:

Predict interactions that will occur (either for the first time or not) among nodes already observed in the network.



Interactions are volatile structures that can occur multiple times and whose value can vanish as time goes by (i.e., telephone calls...)

Rossetti et al. "Interaction prediction in dynamic networks exploiting community discovery." *IEEE/ACM ASONAM*, 2015.

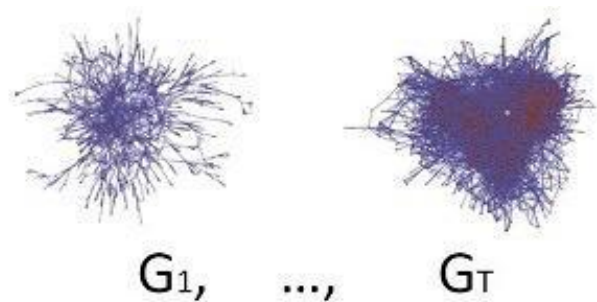
Case Study

Interaction Prediction

Idea:

- Model network evolution through temporal snapshots;
- False Positive reduction:
Community Discovery as a bound for strong ties;
- Time-Aware approach:
time series forecast of topological measures;
- Supervised Approach:
ensemble of classifiers learnt on the topological features, tested on the forecasted ones.

Given a set $G = \{G_0, \dots, G_t, \dots, G_T\}$ of ordered network observations, with $t \in T = \{0 \dots T\}$, the **interaction prediction** problem aims to predict new interactions that will take place at time $\tau + 1$ thus composing $G_{\tau+1}$.



Case Study

Interaction Prediction

Step 1:

For each temporal snapshot $t \in T$ compute a partition $C_t = \{C_{t,0}, \dots, C_{t,k}\}$ of G_t using a community discovery algorithm.

Step 2:

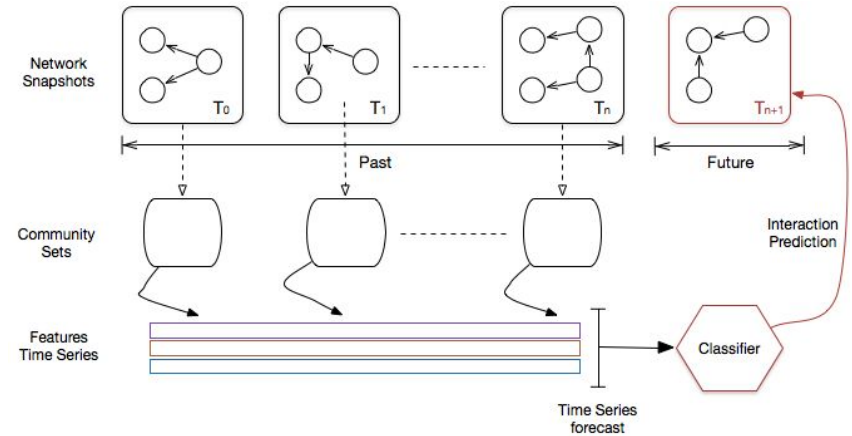
For each $t \in T$ compute a set of measures F for each nodes pair (u,v) belonging to at least a community in C_t

Step 3:

For each node pair (u,v) and feature $f \in F$ build a time series $S^{u,v}$ and apply a forecasting techniques in order to obtain its future expected value $f^{u,v}$

Step 4:

Use the set of expected values $f^{u,v}$ to predict future intra-community interactions.



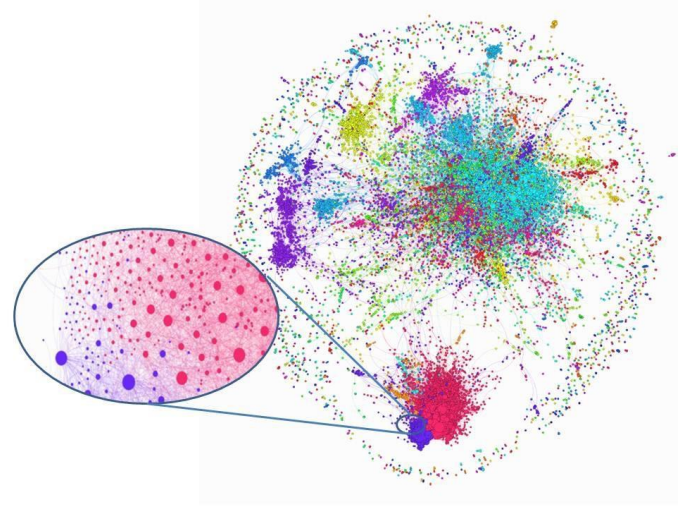
Case Study

Interaction Prediction

Step 1: Community Discovery (CD)

Each CD algorithm proposes its own Community Definition.

- **Demon**
(ego-network based, overlap)
- **Louvain**
(modularity, crisp partition)
- **Infohiermap**
(conductance, crisp partition)



Case Study

Interaction Prediction

Step 2: Features

On the identified communities we compute three set of features:

- **Pairwise Structural Features**
(i.e., Jaccard, CN, Adamic/Adar...)
- **Node Topology Features**
(PageRank, edge betweenness...)
- **Community Features**
(i.e., density, size, shared communities, avg. clustering...)



Case Study

Interaction Prediction

Step 3: Time Series Forecast

For each time series we apply several forecasting model in order to extract the expected future value.



Measure	Description
Last Value (Lv)	$\Theta_t = Z_{t-1}$
Average (Av)	$\Theta_t = \frac{\sum_{i=1}^T Z_i}{T}$
Moving Average (Ma)	$\Theta_t = \frac{\sum_{i=t-n}^t Z_i}{n}$
Linear Regression (LR)	$\Theta_{t+h} = \alpha_t + h\beta_t$

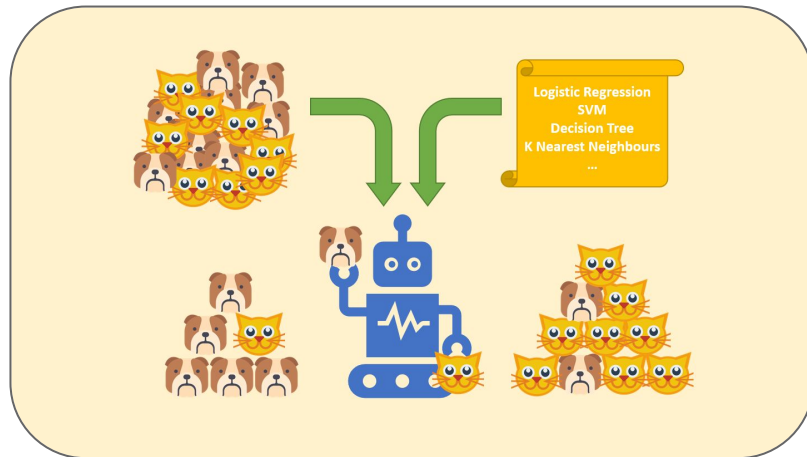
Case Study

Interaction Prediction

Step 4: Classification

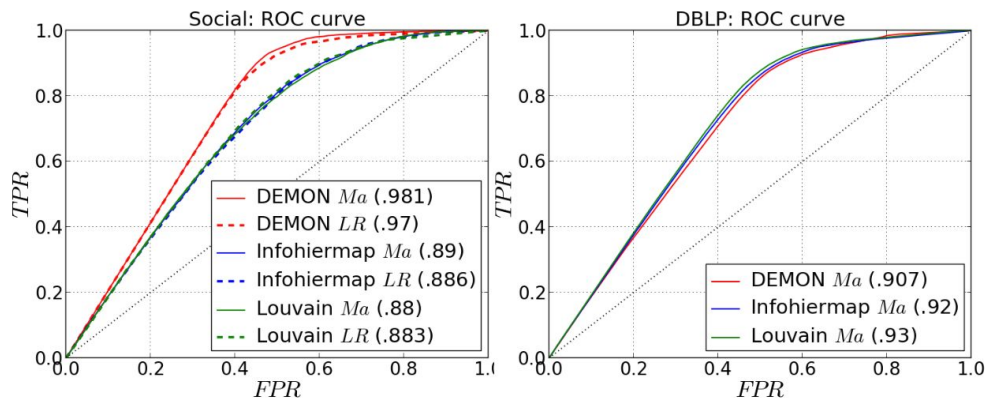
Once learned the features we design two different experiments:

- **Balanced Scenario**
The positive and negative class are balanced through downsampling in order to design a standard baseline
- **Unbalanced Scenario**
The data positive/negative class ratio is maintained. Due to network sparsity we observe a strong negative prevalence (~98%)



Case Study

Interaction Prediction: **Balanced Scenario**



Network	DBLP		Social	
Algorithm	AUC	ACC	AUC	ACC
DEMON <i>Ma</i>	0.907	85.68%	0.981	93.55%
DEMON <i>LR</i>	0.901	84.35%	0.970	91.87%
LOUVAIN <i>Ma</i>	0.930	87.72%	0.880	80.27%
LOUVAIN <i>LR</i>	0.926	87.48%	0.883	81.37%
INFOHIERMAP <i>Ma</i>	0.920	86.69%	0.890	81.34%
INFOHIERMAP <i>LR</i>	0.917	86.18%	0.886	80.89%

Very high accuracy and AUC

CD approaches contribution to IP
is topology sensitive

Case Study

Interaction Prediction: **Balanced Scenario** (cont'd)

Which feature set is the most predictive?

Algorithm	Structural		Topology		Community	
	AUC	ACC	AUC	ACC	AUC	ACC
DEMON	0.957	90.59%	0.962	91.44%	0.903	83.53%
LOUVAIN	0.850	78.63%	0.875	79.38%	0.724	66.64%
INFOHIERMAP	0.876	79.85%	0.887	80.81%	0.667	62.11%

False Positive Filtering (FSF)

vs.

No Filtering (SF)

Algorithm	<i>Ma</i>		<i>LR</i>	
	AUC	ACC	AUC	ACC
SF	0.901	82.88%	0.895	82.18%
FSF	0.956	90.10%	0.937	88.09%

All Forecast with Filtering

vs.

No Filtering

Algorithm	AUC	ACC
DEMON <i>All</i>	0.981	93.90%
LOUVAIN <i>All</i>	0.901	83.05%
INFOHIERMAP <i>All</i>	0.894	81.91%
FS <i>All</i>	0.959	90.44%

Case Study

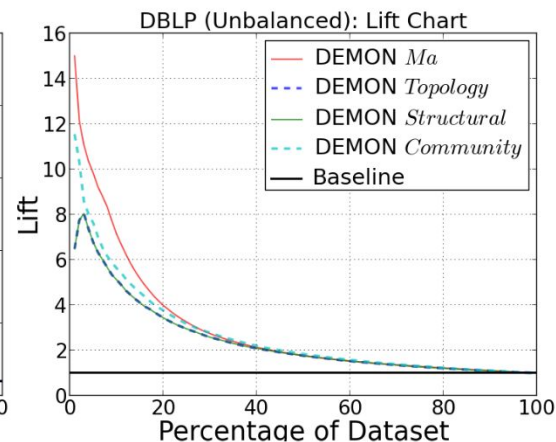
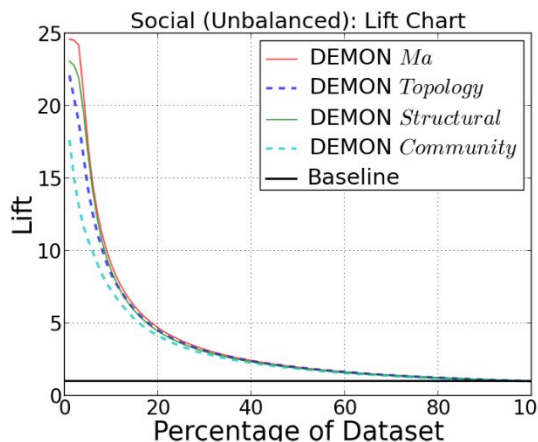
Interaction Prediction: Unbalanced Scenario

Negative class:

- Social 95.9%
- DBLP 98.9%

Very hard baselines

- majority classifiers scores $\sim .96$ and $\sim .99$ precision
(always predicting “no edges”)
- the proposed workflow is able to reach $\sim .96$ and $\sim .45$ precision
w.r.t. the positive class



Case Study

Interaction Prediction: What about weak links?

High accuracy is guaranteed by focusing the prediction on intra-community interactions.

Inter-Community Interaction Prediction

Focus on the predicting the presence/absence of *at least* a new interaction across two communities

- no identification of the “real” endpoints
- no identification of the multiplicity

Idea

1. Construct a new network where the **meta-nodes** are the communities
2. Apply the same workflow to such graph

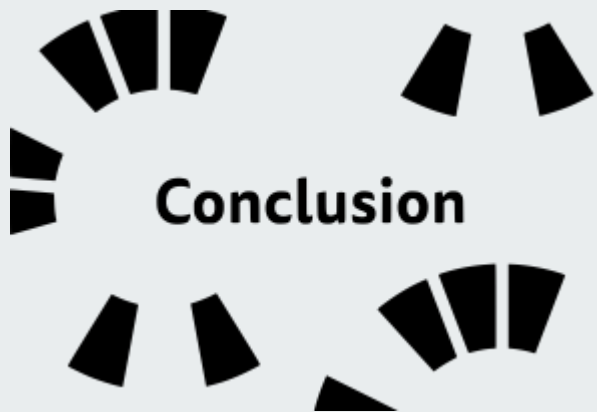
Algorithm	AUC	PPV (%)
Lv	0.594	33.33
Avg	0.632	07.02
Ma	0.647	50.00
LR	0.596	50.00
Flat Graph	0.316	57.20
Baseline	0.504	4.01

Infohiermap performances for the inter-community prediction. Like in the balanced scenario, the Moving Average Ma forecasted features allow for the best classification models

In bold the AUC of the best performing approach

Case Study

Interaction Prediction



Even though Interaction prediction is a complex problem it is possible to reach high accuracy through:

- **Target selection:**
False Positive reduction via Community Discovery
Weak interactions treated as "*special cases*"
- **Local topology history analysis:**
Feature forecast via Time Series analysis

Moreover, each type of datasets demands a specific CD algorithm:

- One-to-one interactions (i.e., social ones)
- Many-to-many interactions (i.e., co-authorship relations)

Chapter 10

Conclusion

Take Away Messages

1. Link wiring patterns define how network topology evolves
2. Predicting new links is a complex task due to network sparse topologies
3. Static and dynamic formulation of the problem account for different peculiarities

Suggested Readings

- Liben-Nowell & Kleinberg paper

What's Next

Chapter 11:
Dynamic Community Discovery

