Link Prediction

Online Social Networks Analysis and Mining

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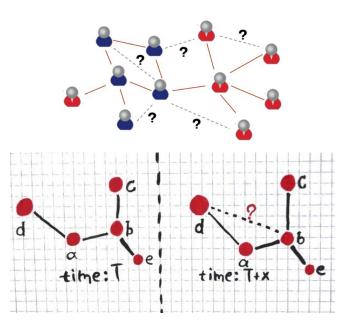
Link Prediction

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**)



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. 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

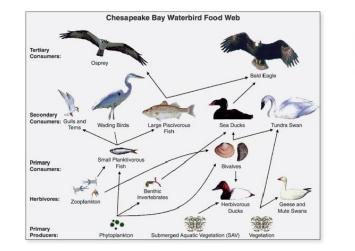


Friendship prediction (i.e., Facebook)

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)





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

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facebook

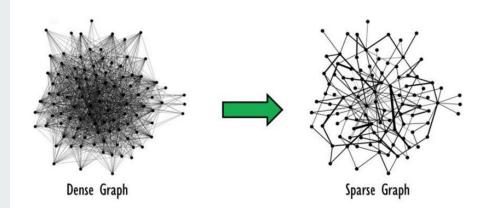
amazon

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Link Prediction Task Complexity

- 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)



Concretizing an Intuition...

Scientists who are close in the network

(i.e., have common colleagues)

 \rightarrow will likely collaborate in the future

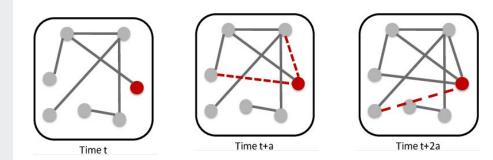
Goal:

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



Link Prediction

- 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 score: 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

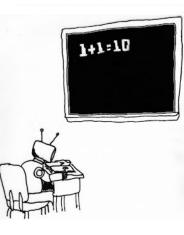
Link Prediction



Unsupervised

Define a set of proximity measures unrelated to the particular network

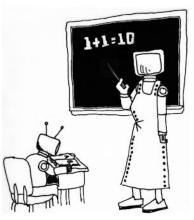
UNSUPERVISED MACHINE LEARNING



Supervised

Extract knowledge from the network in order to improve prediction accuracy

SUPERVISED MACHINE LEARNING



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.

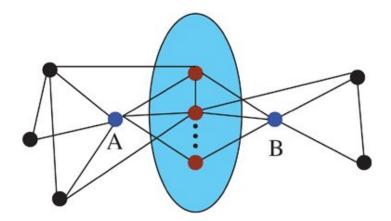
Neighborhood measures

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

score $(u, v) = |\Gamma(u) \cap \Gamma(v)|$ Jaccard: the more similar our friends circles are, the more likely we will become friends

$$ext{ score } (u,v) = rac{|\Gamma(u)n\Gamma(v)|}{\Gamma(u)\cup r(v)|}$$



Neighborhood measures

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

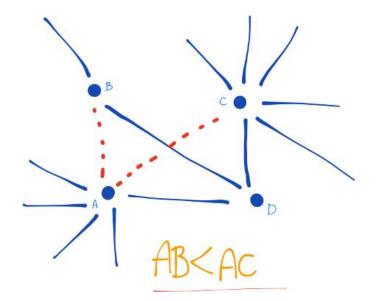
Adamic Adar:

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

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

Preferential Attachment:

the more friends we have, the more likely we will become friends score $(u, v) = |\Gamma(u)| * |\Gamma(v)|$



Path-based measures

How distant are we?

Graph Distance:

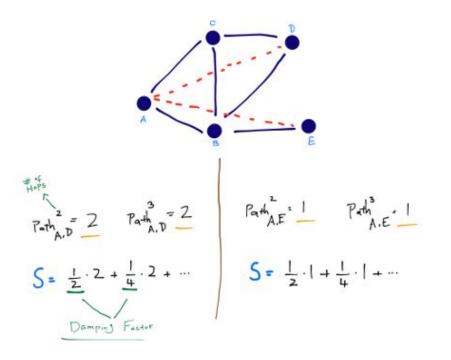
(negated) length of the shortest path between two nodes

Katz:

weighted sum over all the paths between two nodes

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

where: $paths_{u,v}^{(I)} = \{ paths of length exactly I from u to v \}$



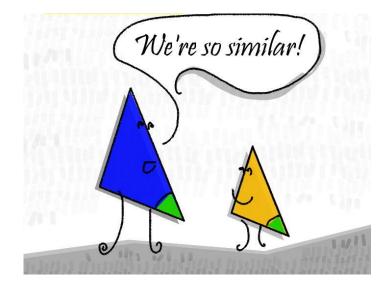
Ranking

How similar are we?

SimRank:

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

 $egin{aligned} ext{similarity} & (u,v) = \gamma st rac{\sum_{s \in r(u)} \sum_{n \in \Gamma(v)} ext{similarity} (a,b)}{|\Gamma(u)| st \Gamma(v)|} \ ext{score} & (u,v) = ext{similarity} & (u,v) \end{aligned}$



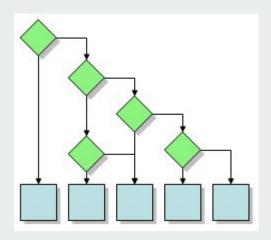
Limitations

- 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
 - \Rightarrow there is useful information in network topology





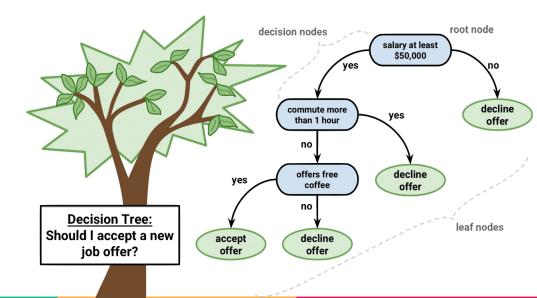
The process is now organized in 4 steps:

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

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

Stacking 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

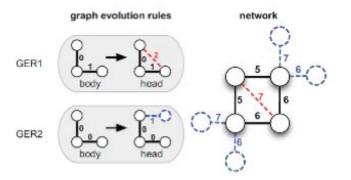


Frequent Pattern Mining

GERM

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

- e.g., generalization of triadic closure



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

Network Embedding

Idea

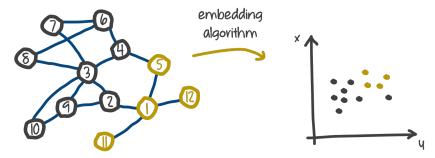
Graphs can be *mapped* into vector spaces

- 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 ...

to real vector representation



- 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

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.

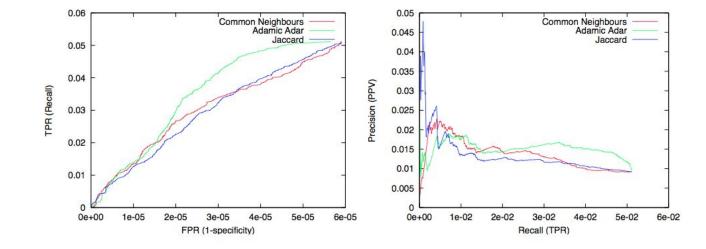
Random Predictor

each edge has the same probability to appear in the network

 $ext{performance } (p) = rac{TP}{TP + FP}$ $ext{ratio} = rac{ ext{performance } (p)}{ ext{performance } (ext{prandom})} = rac{ ext{performance } (p)}{rac{ ext{V}(-(\sqrt{V}-1)|}{2} - |E_{ ext{old }}|}$ $ext{if ratio > 1 then p is meaningful}$

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

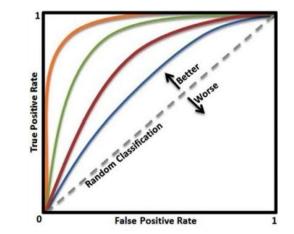
- Precision: PPV = TP/(TP+FP)
- Recall: TPR = TP/(TP+FN)

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'
р	TP	FN
n	FP	TN

Confusion Matrix

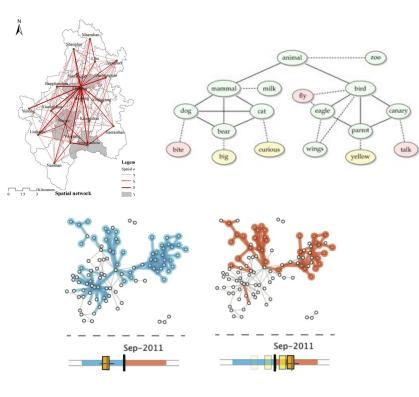
Link Prediction Something more...

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:

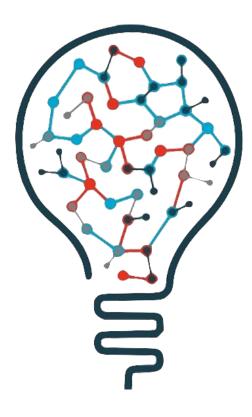
- Dynamic Networks
- Multiplex networks
- ...



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. False Positive prediction
 - Bridges !?!
- 4. Simple approaches are "too simple"
- 5. Complex approaches are **costly**

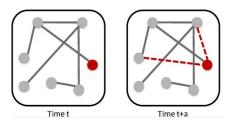


Case Study: Interaction Prediction in Dynamic Networks



Link Prediction goal:

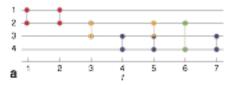
Predict ties that are not present in actual network configuration.



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.

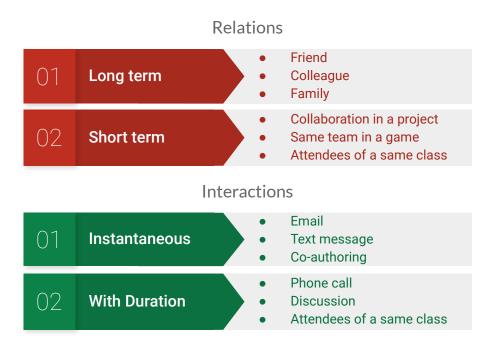
Semantic

Relations Vs. Interactions

Topological perturbations may have different temporal scales depending on their intrinsic semantic value.

Two families:

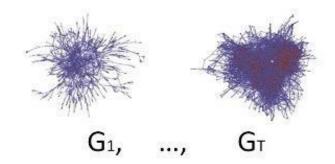
- Relations (stable ties)
- Interactions (unstable ties)



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



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:

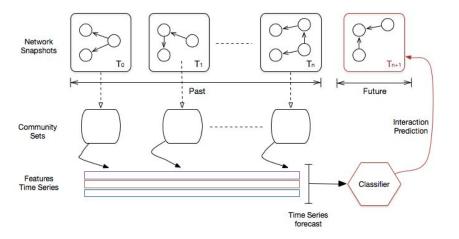
For each t \in T compute a set of measures F for each nodes pair (u,v) belonging to at least a community in C_t

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:

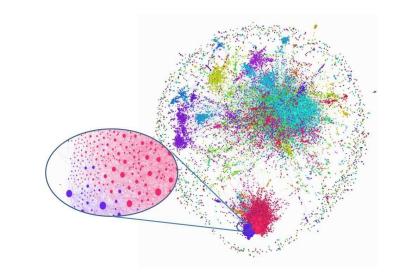
Use the set of expected values $f^{u,v}$ to predict future intra-community interactions.



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)



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...)



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)	$\Theta_t = \frac{\sum_{i=1}^{T} Z_i}{\sum_{i=1}^{T} T}$
Moving Average (Ma)	$\Theta_t = \frac{\sum_{i=\tau-n} Z_i}{n}$
Linear Regression (LR)	$\Theta_{t+h} = \alpha_t + h\beta_t$

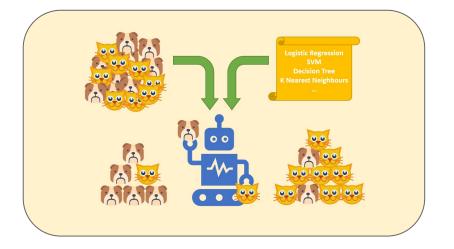
Step 4: Classification

-

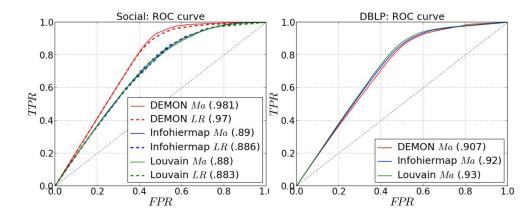
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Once learned the features we design two different experiments:

- Balanced Scenario The positive and negative class are balanced through downsampling 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 Ma	0.907	85.68%	0.981	93.55%	>
Demon LR	0.901	84 35%	0.970	91.87%	
Louvain Ma $<$	0.930	87.72%	0.880	80.27%	-
Louvain LR	0.926	87.48%	0.883	81.37%	
Infohiermap Ma	0.920	86.69%	0.890	81.34%	-
Infohiermap LR	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)

Algorithm		Structural		Topology		Community	
					200 - 200 Berlin - 2008		
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%	
	DEMON Louvain	AlgorithmAUCDEMON0.957LOUVAIN0.850	Algorithm AUC ACC DEMON 0.957 90.59% LOUVAIN 0.850 78.63%	Algorithm AUC ACC AUC DEMON 0.957 90.59% 0.962 LOUVAIN 0.850 78.63% 0.875	Algorithm AUC ACC AUC ACC DEMON 0.957 90.59% 0.962 91.44% LOUVAIN 0.850 78.63% 0.875 79.38%	Algorithm AUC ACC AUC ACC AUC DEMON 0.957 90.59% 0.962 91.44% 0.903	

Folco Docitivo Filtoring (FSF)	Algorithm		Ia	LR	
False Positive Filtering (FSF)	Algorithm			AUC	ACC
VS.	SF		82.88%		
No Filtering (SF)	FSF	0.956	90.10%	0.937	88.09%

	Algorithm	AUC	ACC
All Forecast with Filtering	DEMON All	0.981	93.90%
VS.	LOUVAIN All	0.901	83.05%
No Filtering	Infohiermap All	0.894	81.91%
NOTILETING	FS All	0.959	90.44%

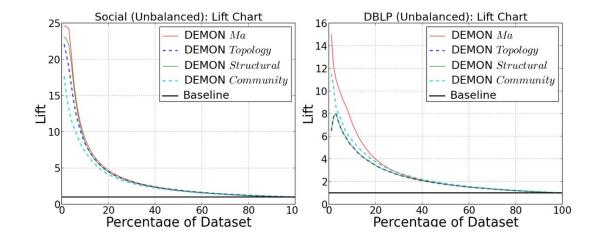
Case Study Interaction Prediction: Unbalanced Scenario

Negative class:

- Social 95.9%
- DBLP 98.9%

Very hard baselines

- majority classifiers scores ~.96 and
 ~.99 precision
 (always predicting "no edges")
- the proposed workflow is able to reach ~.96 and ~.45 precision w.r.t. the <u>positive class</u>



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

Lv0.59433.33Avg0.63207.02Ma 0.647 50.00LR0.59650.00	Algorithm	AUC	PPV (%)
Ma 0.647 50.00	Lv	0.594	33.33
	Avg	0.632	07.02
LR 0.596 50.00	Ma	0.647	50.00
	LR	0.596	50.00
Flat Graph 0.316 57.20	Flat Graph	0.316	57.20
Baseline 0.504 4.01	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

Conclusion

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)