Jan Ramon
ECML 2013

## LEARNING AND MINING WITH NETWORK-STRUCTURED DATA



## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Learning


## Introduction

Network = Objects + Relations

| Application | Objects | Relations |
| :--- | :--- | :--- |
| Social network | Person | Friendship, colleague |
| Traffic network | Crossroad | Road |
| Chemical interaction net | Chemicals | Interaction |
| Telecommunication net | Person | Phone call |
| Citation network | Paper, researcher | Citation, authorship |
| Shareholder network | Company, Person | Shareholdership |
| Computer network | Computer, router | Cable, wifi registration |

## Introduction

- In this tutorial:
- Who studies networks?
- Network patterns \& mining them
- Learning in networks
- Focus on
- Local patterns
- Not so much on large scale properties


## Related ECML/PKDD 2013 tutorials

- [Fri-PM] Algorithmic techniques for modeling and mining large graphs (Alan Frieze et al.)
- Focus is more on global properties
- [Mon-PM] Discovering Roles and Anomalies in Graphs: Theory \& Applications (T. Eliassi-Rad et al.)
- Anomaly detection is not covered here
- [Fri-AM] Statistically sound pattern discovery (G. Webb \& W. Hamalainen)
- Different statistical aspects


## Introduction <br> Prerequisites

Supervised learning: Given i.i.d. training examples, learn a function from example to target value.

- You could use SVM, DT, NB, IBL, GP, ... or any of your favorite supervised techniques


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Learning


## Contents

- Introduction
- Networks from different points of view
- Basic concepts
- Data mining tasks
- Relevant fields of research
- Patterns \& pattern mining
- Learning


## Basic concepts - Graphs

- An undirected (labeled) graph is a tuple $G=(V, E, \lambda)$ where
- $V$ is a set of vertices (nodes) [punten (knopen)]
- $E \subseteq\{\{v, w\} \mid v, w \in V\}$ is a set of edges [takken]
- $\lambda: V \cup E \rightarrow \Sigma$ is a labeling function
- Unlabeled graph:
- If $\lambda$ is constant (all vertices/edges have the same label), $\lambda$ may be omitted


## Basic concepts - Graphs

- A directed graph is a tuple $G=(V, E, \lambda)$ where
- $V$ is a set of vertices (nodes) [punten (knopen)]
- $E \subseteq\{(v, w) \mid v, w \in V\}$ is a set of arcs [bogen]
- $\lambda: V \cup E \rightarrow \Sigma$ is a labeling function
- Further notations:
- $V(G)$ is the set of vertices of the graph $G$
- $E(G)$ is the set of edges / arcs of the graph $G$
- $\lambda_{G}$ is the labeling function of the graph $G$
- $N_{H}(v)=\{w \in V(H) \mid\{v, w\} \in E(H)\}$ is the neighborhood of $v$
- $\Delta v=N_{H}(v)$ is the degree of $v$


## Basic concepts <br> adjacency matrix

- Adjacency matrix of graph $G$ is a square matrix $A$ of dimension $V(G) \times V(G)$ such that
$A_{u, v}=0$ if $u$ and $v$ are not connected
- $A_{u, v}=1$ if there is an edge between $u$ and $v$


$$
\begin{array}{lllll}
1 & 2 & 3 & 4 \\
1 & 0 & 1 & 1 & 1 \\
2 & 1 & 0 & 0 & 0 \\
3 & 1 & 0 & 0 & 1 \\
4 & 1 & 0 & 1 & 0
\end{array}
$$

## Basic concepts - Walk/Path

- A walk $P$ between vertices $v$ and $w$ in a graph $G$ is a sequence of vertices $u_{1}, u_{2}, \ldots, u_{n} \in$ $V(G)$ such that
- $u_{1}=v$,
- $u_{n}=w$ and
- $\left(u_{i}, u_{i+1}\right) \in E(G)$ for all $1 \leq i \leq n-1$.
- The length of such walk $P$ is $n-1$.
- A path is a walk where all vertices are distinct
- Slightly abusing terminology, a path $P$ can also be seen as a subgraph of $G$


## Basic concepts - Shortest path

- A shortest path is a path of minimal length.
- Distance $d(u, v)$ between $u$ and $v$ is length of shortest path between $u$ and $v$
- The diameter of $G$ is

$$
\operatorname{diam}(G)=\max _{u, v \in V(G)} d(u, v)
$$

## Basic concepts - Diameter

The diameter of $G$ is

$$
\operatorname{diam}(G)=\max _{u, v \in V(G)} d(u, v)
$$

Many real-world graphs have small diameter.

- V : all persons
- E : an edge connects persons who have ever met each other
- Many people have met a local politician who met the national prime minister


## Basic concepts - connected, tree

- A graph $G$ is connected iff there is a path between every pair of vertices $v, w \in V(G)$
- A connected component of a graph $G$ is a maximal connected subgraph of $G$.
- A graph $G$ is a tree iff there is a unique path between every pair of vertices $v, w \in V(G)$
- Intuition: if the path between two vertices is not unique, then there is a cycle.


## Basic concepts - morphisms

- A homomorphism from a graph $H$ into a graph $G$ is a mapping $\varphi: V(H) \rightarrow V(G)$ such that
- $\forall v, w \in V(H):(v, w) \in E(H) \Rightarrow(\varphi(v), \varphi(w)) \in E(G)$
- $\forall v \in V(H): \lambda(v)=\lambda(\varphi(v))$
- $\forall v, w \in V(H): \lambda(v, w)=\lambda(\varphi(v), \varphi(w))$
- An injective homomorphism is a subgraph isomorphism.


## Basic concepts - subgraph isomorphism vs homomorphism



A homomorphism, not an isomorphism

- If there is a homomorphism from $H$ to $G$, then we denote this $H \leq_{h} G$
- If there is a subgraph isomorphism from $H$ to $G_{\text {, }}$ then we denote this $H \leq_{i} G$
- $H \equiv G$ iff $H \leq G$ and $G \leq H$


## Basic concepts - morphisms

- An induced homomorphism from a graph $H$ into a graph $G$ is a mapping $\varphi: V(H) \rightarrow V(G)$ such that $\forall v, w \in V(H):(v, w) \in E(H) \Leftrightarrow(\varphi(v), \varphi(w)) \in E(G)$ - $\forall v \in V(H): \lambda(v)=\lambda(\varphi(v))$ - $\forall v, w \in V(H): \lambda(v, w)=\lambda(\varphi(v), \varphi(w))$
- An injective induced homomorphism is an induced subgraph isomorphism.


## Basic concepts - induced vs normal subgraph isomorphism



Subgraph isomorphism
NOT induced subgraph isomorphism


Subgraph isomorphism Induced subgraph isomorphism


Subgraph isomorphism
Induced subgraph isomorphism

## Basic concepts <br> Automorphisms

- Automorphism = isomorphism of a graph on itself.
- |aut(H)| is the size of the automorphism group aut(H).
- For bounded degree $H$, one can compute |aut(H)| in polynomial time.


Triangle "Mirror"
"Rotation"
Aut(triangle) has size $2 * 3=6$

## Basic concepts - Networks




Network =
big database graph
Pattern = small graph

## Contents

- Introduction
- Networks from different points of view
- Basic concepts
- Data mining tasks
- Relevant fields of research
- Patterns \& pattern mining
- Learning


## Network data mining tasks

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  <br> community <br> detection | 2. Pattern mining <br> 3. Edge/vertex <br> structure/feature <br> learning |
| Evolution | 5. Generative <br> models | 4. Temporal <br> learning |

## Clustering \& community

 detection- Given:
- network $D$
- Find:

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  |  |
| Evolution |  |  |

- Set of subsets (clusters, communities) $V_{1}, V_{2}, \ldots, V_{n}$ of $V(G)$
- Covering V(G) or not
- Disjoint or overlapping
- such that vertices in same cluster are close
- similar or connected
- and vertices from different clusters are distant
- dissimilar / not connected

Clusters \& communities
examples

- Find groups of people who are densily connected
- Find groups of people who have a similar opinion or behavior (and are connected)
- People in the same country, company, school, domain, ... will often cluster together.


## Pattern mining

- Given:
- network $D$
- pattern language $L$

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  |  |
| Evolution |  |  |

- interestingness criterion $I: L \times D \rightarrow\{$ true, false $\}$
- Find
- all patterns $P \in L$ for which $I(P, D)$


## Pattern mining example

Why do you (Y) watch "Jurassic park" (JP)?


Vertex/edge structure/
feature prediction

- Given:
- network D,

|  | Global | Local |
| :--- | :--- | :--- |
| Static |  |  |
| Evolution |  |  |

- example set $X=V(D) \cup E(D)$, target space $Y$
- Unknown distribution $P$ on $X \times Y$
- Training set $Z_{\text {train }} \subseteq X \times Y$
- Test set $X_{\text {test }} \subseteq X$
- Loss function $L: Y \rightarrow \mathbb{R}^{+}$
- Find
- $\hat{f}$ minimizing $\mathrm{E}\left[\sum_{i} L\left(\hat{f}\left(x_{i}^{\text {test }}\right), y_{i}^{\text {test }}\right)\right]$


## vertex/edge structure/ feature prediction example

- Predicting on existing objects:
- Social network: Given a user, his profile, his friendship relations, is this user interested in chess?
- Given a pair of friends ( $X, Y$ ). Is X planning to send a message to Y today?
- Predicting on hypothetical objects:
- Given a group of people, do they have a common friend (not yet in the network)?
- Given two people. do they know each other (even though not yet represented in the network)?


## Learning from temporal data

- Given:

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  |  |
| Evolution |  |  |

- time-dependent network $\left.\left\{D_{t}\right\}^{T}{ }^{T}=1\right\}^{\prime}$ (possibly represented by vertices and edges with time stamps etc.)
${ }^{-}$a loss function $L: \mathcal{G} \times \mathcal{G} \rightarrow \mathbb{R}^{+}$
- Find:
- Prediction $\widehat{D}_{T+\Delta t}$ of the network (or parts thereof) to minimize $\mathrm{E}\left[L\left(D_{T+\Delta t}, \widehat{D}_{T+\Delta t}\right)\right]$


## Learning from temporal data example

- Social network:
- When will X update his profile?
- Will $X$ and $Y$ become friends?
- Will a common friend of X and Y join the network?


## Learning generative models

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  |  |
| Evolution |  |  |

- Generative model = probability distribution mapping a network on a new network, i.e.
- $h: \mathcal{G} \times \mathcal{G} \rightarrow[0,1]$ s.t.
$\forall D \in \mathcal{G}, \sum_{D^{\prime} \in \mathcal{G}} h\left(D, D^{\prime}\right)=1$


## Learning generative models

- $h: \mathcal{G} \times \mathcal{G} \rightarrow[0,1]$
- Given:
- A hypothesis space $\mathcal{H}$ of generative models
- An unknown $h \in \mathcal{H}$, we assume there was some $D_{0}$ and for $t=1 . . T$, $D_{t}$ was drawn from $h\left(D_{t-1},\right)$
- time-dependent network $\left\{D_{t}\right\}\{t=1\}$
- Find:
- Given loss function $L: \mathcal{H} \times \mathcal{H} \rightarrow \mathbb{R}^{+}$, a model $\hat{h} \in \mathcal{H}$ such that $\mathrm{E}[\mathrm{L}(\mathrm{h}, \widehat{h})]$ is minimal
- Model $\hat{h} \in \mathcal{H}$ such that $h^{\mathrm{n}}\left(\mathrm{D}_{0}\right)$ has the same asymptotic properties as $D_{T}$.


## Learning generative models

- Difference with learning from temporal data":
- Not just predicting a future event, but also global properties should be right
- E.g. errors which propagate quickly should be avoided. (e.g. positive feedback loops)


## Wnat data do I need?

- Many datasets are undirected unlabeled graphs (interaction = yes/no)
- Ok for many models focussing on global \& assymptotic aspects
- How about correlations between interests of friends?
- Then you need to records people's interests
- In general, "local" models will need richer data


## Contents

- Introduction
- Networks from different points of view
- Basic concepts
- Data mining tasks
- Relevant fields of research
- Patterns \& pattern mining
- Learning


## Relevant fields of research

- Statistical physics
- Complex systems
- Multi agent systems, ants, other simulation
- Grammar induction
- (Algorithmic) graph theory
- Spectral graph theory
- pattern mining, data mining, machine learning


## Statistical physics

- If we assume a given set of laws, what will happen?
- Law = graph generation model
- Erdos-Reny model:
- Barabasi-Albert model
- Result =
- Assymptotic behavior, what happens if there are many particles?


## Statistical physics The Erdos-Reny model

- A random graph from the Erdos-Reny distribution $G_{p}(n, p)$ is constructed as follows:
- Let $G$ be a graph on $n$ vertices.
- For every pair of vertices $\{v, w\}$, connect $v$ and $w$ with an edge with probability $p$.
- A random graph from the Erdos-Reny distribution $G_{M}(n, M)$ is constructed as follows:
- Let $G$ be a graph on $n$ vertices.
- Choose randomly $M$ elements from $\{\{v, w\} \mid v, w \in V(G)\}$ and draw an edge between the two elements of these pairs.

Statistical physics
Allmost all graphs

- Let $\mathrm{G}_{\mathrm{n}}$ be a random graph drawn from $G(n, p(n))$, i.e. $p$ is a function of $n$. A predicate $q$ (i.e. a boolean function) holds for (asymptotically) allmost surely (a.a.s.) if

$$
\lim _{n \rightarrow \infty} P\left(q\left(G_{n}\right)=\text { true }\right)=1
$$

- Similar for $G(n, M)$
- If no $G(n, p)$ or $G(n, M)$ specified: $G(n, 1 / 2)$ by default ("allmost every graph").


## Statistical physics Assymptotic properties

- E.g. the "giant component"
- If $\lim _{n \rightarrow \infty} n p<1$, then the largest component of a $G(n, p)$ graph is a.a.s. not larger than $3 \cdot \log (n) /(1-n p)^{2}$
- If $\lim _{n \rightarrow \infty} n p=1$, then the largest component of a $G(n, p)$ graph is a.a.s. $\mathrm{n}^{2 / 3}$
- If $\lim _{n \rightarrow \infty} n p>1$, then the largest component of a $G(n, p)$ graph is a.a.s. close to $\beta \mathrm{n}$ with $\beta+e^{-\beta p n}=1$


## Statistical physics <br> Assymptotic properties

- E.g. connectedness:
- If $\lim _{n \rightarrow \infty} n p / \ln (n)<1$ then, $G(n, p)$ is a.a.s. disconnected
- If $\lim _{n \rightarrow \infty} n p / \ln (n)>1$ then, $G(n, p)$ is a.a.s. connected


## Statistical physics <br> 0-1 law

- Given a first order logic formula F over graphs, $\lim _{n \rightarrow \infty} P(F(G(n, 1 / 2))=$ true $)$ is either 1 or 0.
- E.g. "contains a triangle": adding vertices (and hence edges) only increases the probability of a triangle; if many vertices, the probability gets close to 1 .


# Statistical physics What can ML and DM learn? 

- Compute from the model the "expected value" of a pattern.
- An "interesting pattern" is one which deviates from the expected value according to the model.
- E.g. assume $G$ is the union of a random graph and a clique. Under what conditions can we detect the clique as an abnormally dense spot?1


## Complex systems

- Model processes in society
- systems of interacting individuals
- what (often large-scale) properties do we observe?
- Study behavior of systems with such properties
- assymptotics
- simulation of systems
- emerging patterns


## Complex systems <br> What can ML/DM learn?

- Techniques to model application domains
- Social behavior
- Economics
...


# Multi-agent systems, ants, simulation 

- How to simulate?
- Do artificial populations offer more value than artificial individuals?


## Multi-agent systems What can ML/DM learn?

- Simulation (sampling a temporal model forward in time)
- Multi-agent learning (and the effects of changing behavior due to learning)
- Game theory (~statistical physics: Nash equilibria)


## Grammar induction

- Generative model ~ probabilistic grammar
- Initial state
- (probabilistic) production rules
- Graph grammars
- Hyperedge replacement grammars
- Vertex replace grammars


## Grammar induction What can DM/ML learn?

- Graph grammars: learning generative models producing certain probability distributions over graphs


## (Algorithmic) graph theory

- Algorithms on graphs and their complexity are well-studied.
- How to solve graph problems
- Complexity of solving problems


## (Algorithmic) graph theory What can ML/DM learn?

- Pattern matching
- E.g. see part 2
- Support measures
- e.g. Maximum independent set, Lovasz theta function, ... (part 3)
- Shortest path algorithms
- Similarity, maximum common subgraph, ...


## Relational databases

- A network is a relational database where the binary "edge" relation has foreign keys to itself.
- Matching patterns = evaluating queries


## Relational databases What can ML/DM learn?

- Database theory sometimes shows how to match patterns (=evaluate queries), but is usually not optimized for "recursive" foreign keys.
- Many ideas for data structures (e.g. recently growing interest in graph database indexing)


## Spectral graph theory¹

- Study of
- the adjacency matrix of a graph
- its eigenvalues
- the Laplacian matrix: $L=D-A$ with $D$ degree matrix ( $D_{v v}$ is degree of $v$ ) and $A$ adjacency matrix.


## Spectral graph theory What can ML/DM learn?

- Laplacian describes "influence flow", used in
- semi-supervised learning
- manifold embedding
- Clustering:
- \# of zero eigenvalues = \# of connected components


## Mining, learning

- Relational learning (e.g. SRL)
- Learning in graphs (e.g. MLG)
- Using logical representations (e.g. ILP)
- (but adding logic makes problems often undecidable)


## Introduction summary (1/3) Basic concepts

- Graphs
- labels
- adjacency matrix
- Paths
- distance
- Morphisms (matching operators)


## Introduction summary (2/3) Network data mining tasks

|  | Global / <br> assymptotic | Local |
| :--- | :--- | :--- |
| Static |  <br> community <br> detection | 2. Pattern mining <br> 3.Edge/vertex <br> structure/feature <br> learning <br> Evolution5. Generative <br> models | | 4.Temporal <br> learning |
| :--- |

## Introduction summary (3/3) Domains researching networks



## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Introduction
- Pattern matching
- Frequency
- Additional remarks
- Learning


## Introduction What is a pattern?

- Pattern = collection of vertices that should satisfy some constraints (connections, labels, ...)



## Introduction

A generic pattern miner
Assume interesting is anti-monotonic $k \leftarrow 0$; $C_{0} \leftarrow$ MinimalPatterns
While $C_{k} \neq\{ \}$

$$
\begin{aligned}
& S_{k} \leftarrow\left\{P \in C_{k} \mid \text { interesting }(P)\right\} \\
& C_{k+1} \leftarrow \mathrm{U}_{P \in S_{k}} \text { extensions }(P) \\
& k \leftarrow k+1
\end{aligned}
$$

EndWhile
Solutions $\leftarrow \mathrm{U}_{k} S_{k}$

## Introduction

## A generic pattern miner

- "assume interesting is anti-monotonic" allows for pruning.

$$
\begin{aligned}
& \text { interesting }(G) \wedge G \leq H \\
& \Rightarrow \operatorname{interesting}(H)
\end{aligned}
$$

- Can also mine using non-anti-monotonic criterion (e.g. correlated patterns ${ }^{1}$ )
- Breadth-first "Apriori-style" ${ }^{2}$
- Also depth-first possible ${ }^{3}$
- To be instantiated:
- MinimalPatterns \& Extensions
- Interesting


## Introduction

## Enumeration of patterns

To be instantiated:

- MinimalPatterns \& Extensions
- Interesting
- Many approaches are generate-and-test
- How to generate all graphs subject to antimonotonic constraint?
- Practice: mining graph patterns in databases of transactions represented with graphs: AGM, gSpan, FSG, Gaston² $\rightarrow$ use canonical form
- Theory: polynomial-delay (+evaluation antimonotone criterion) ${ }^{1}$

[^0]
## Introduction <br> Complexity notions

- Enumeration/listing problems ("find all ...") may have output $O$ exponential in input size $I$.
- Polynomial delay: between solution $j-1$ and $j$ at most poly(I) time.
- Incremental polynomial time: between solution $j-1$ and $j$ at most poly $(I, j)$ time.
- Output-polynomial time: total running time at most poly $(I, O)$


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Introduction
- Pattern matching
- Frequency
To be instantiated:
$\quad$ MinimalPatterns \& Extensions
- Additional remarks
- Learning


## Pattern matching Overview

- Problem statement
- Hardness results
- Triangle counting
- Small patterns
- Larger patterns
- Cliques
- Sampling
- Fixed parameter tractability


## Pattern matching

## The problem

- Given:
- A network D
- A pattern $P$
- Find listisome/one/...) embeddings of P in D OR
- an aggregate (count, average, ...) computed over these embeddings


## Pattern matching Why do we care?

- Basic operation for both learning and mining
- There is a literature on basic pattern matching, but learning \& mining queries have specific characteristics
- Data is rich, satisfies integrity constraints, ...
- Patterns may have wildcards
- DM/ML is aimed at collecting statistics


## Pattern matching <br> Subgraph isomorphism complexity

- Pattern $P$, network $D$
- List embeddings $\pi$ : $P \rightarrow D$
- \#P-complete
- Classic algorithms (backtracking search):
$O\left(|V(D)|^{|V(P)|}\right)$

[^1]
## Pattern matching Heuristic search

Ullmann's algorithm ${ }^{1}$ (pattern $P$, network $D$ ) $\operatorname{Match}(P, D,\{ \})$

Procedure Match ( $P$, $D$, partial embedding $\pi$ ) If $V(P)=\operatorname{dom}(\pi)$
then ListSolution $(\pi)$
else
Select $v \in V(P) \backslash \operatorname{dom}(\pi)$
Let $C=\{w \in V(D) \mid w$ maybe image of $v\}$
For all $w \in C$ do
$\operatorname{Match}(P, D, \pi \cup\{(v, w)\})$

## Pattern matching <br> Avoiding worst-case complexity

- Database of transactions of graphs
- E.g. many small molecule graphs, RNA
- exploit structure of database graphs, e.g.
- atoms have max degree = 6 ¹
- molecules are often planar (or even outerplanar ${ }^{2}$ )
- bounded treewidth ${ }^{3}$

[^2]
## Pattern matching <br> Avoiding worst-case complexity

- Network
* No definite structure we can rely on
x Approximate matching may help but
- Subgraph isomorphism is hard to approximate
$\checkmark$ Structural properties of patterns
- Triangles and other small patterns
- Trees
$\checkmark$ Statistical properties of network
- Random graphs


## Pattern matching Triangle counting

- Simple problem:
- Given: a triangle $P$ and a network $D$
- List or count: all embeddings of $P$ in $D$
- but a lot of literature on solving it


## Pattern matching Triangle counting

- Idea 1: Brute force
- Check every triple of vertices of $G$
- Runtime $O\left(|V(G)|^{3}\right)$


## Pattern matching Triangle counting

- Idea 2: matrix multiplication ${ }^{1}$
- Let $A$ be the adjacency matrix of $G$
- $\left(A^{n}\right)_{u, v}=$ \# of walks of length $n$ between $u$ and $v$
- Matrix multiplication is $O\left(|V(G)|^{2.37}\right)^{2}$
- Hence, compute $\operatorname{tr}\left(A^{3}\right) / 3$ !
$\int_{2}^{1}$
$A=\left(\begin{array}{llll}0 & 1 & 1 & 1 \\ 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 \\ 1 & 0 & 1 & 0\end{array}\right)$
$A^{2}=\left(\begin{array}{llll}3 & 0 & 1 & 1 \\ 0 & 1 & 1 & 1 \\ 1 & 1 & 2 & 1 \\ 1 & 1 & 1 & 2\end{array}\right) \quad A^{3}=\left(\begin{array}{llll}2 & 3 & 4 & 4 \\ 3 & 0 & 1 & 1 \\ 4 & 1 & 2 & 3 \\ 4 & 1 & 3 & 2\end{array}\right)$
$\operatorname{tr}\left(A^{3}\right) / 3!=(2+2+2) / 6=1$

[^3]${ }^{2}$ For practical problems, the exponent will be higher

## Pattern matching Triangle counting

- Idea 3: Sparse graphs
- Iterate over edges ${ }^{1}: O\left(|E(G)|^{\frac{3}{2}}\right)$
- Nodelterator: Iterate over pairs of neighbors of vertices: $O\left(d_{\text {max }}^{2}|E(G)|\right)$ with $d_{\max }$ the maximal degree of $G$.


## Pattern matching Triangle counting

- Idea 4: Approximation
- Sparsification: delete randomly part of the graph, count triangles, adjust for sampling
- Sampling: perform a number of random attempts, adjust for sampling
- Partition graph into parts, solve them, adjust for triangles spanning several partitions.


## Pattern matching Triangle counting

| Alon et al., 1997 | Exact | $O\left(\|V(G)\|^{2.37}\right)$ |
| :--- | :--- | :--- |
| Itai \& Radeh, 1978 | Exact | $O\left(\|E(G)\|^{\frac{3}{2}}\right)$ |
| Tsourakakis ICDM2008 | Exact |  |
| Faloutsos KDD2009 | Approx | Sparsification |
| Avron 2010 | Approx | Sampling |
| Pagh\&Tsourakakis, InfProcLet 2012 | Approx | Partitioning |
| Becchetti et al. TKDD 2010 | Approx |  |

${ }^{1}$ Itai and Rodeh 1978

## Pattern matching <br> Counting motifs of size 3-5

- Motifs = not necessarily connected induced subgraph
- Brute force sometimes still works ${ }^{1}$
- Sampling strategy (e.g. GUISE²):
- Consider an induced graph
- Omit vertex or add neighbor
- Metropolis hastings: adjust for fact that higherdegree vertices are reached more easily

[^4]
## Pattern matching <br> Sampling motifs with MCMC

Motifs


## Pattern matching <br> Sampling motifs with MCMC

Motifs


Pattern matching
Sampling motifs with MCMC


Motifs
State $s 1$
$d(s 1)=9$


## Pattern matching

 Sampling motifs with MCMC
$T(u, v)=\min \left(\frac{1}{d(u)}, \frac{1}{d(v)}\right) \quad($ for $u \neq v)$
$T(u, u)=1-\sum_{v \neq u} T(u, v)$
Motifs
State s1
$d(s 1)=9$
$\underset{\square}{Z}$
$\vec{\nabla}$
$\nabla$
0
Z
Z

Pattern matching
Sampling motifs with MCMC


Motifs


## Pattern matching Overview

- Problem statement
- Hardness results
- Triangle counting
- Small patterns
- Larger patterns
- Cliques
- Sampling
- Fixed parameter tractability


## Pattern matching <br> Cliques

- Several approaches aim at finding maximal cliques
- Clique detection for non-directed graphs,


## Clusters \& communities Bicliques

- Bi-partite networks $D=\left(V_{1} \cup V_{2}, E\right)$ with $E \subseteq V_{1} \times V_{2}$.
- Bi-clique $=C_{1} \times C_{2}$ with $C_{1} \subseteq V_{1}$ and $C_{2} \subseteq V_{2}$.
- Several variants:
- Tile mining (in boolean matrices, unweighted networks)
- Non-negative matrix factorization (real-valued matrices, weighted networks)
Advertisement:
Join ECML/PKDD2013 Tue1B session on Networks (1)
J.Ramon, P.Miettinen, J.Vreeken, Detecting bicliques in GF[q],


## Pattern matching

## Larger patterns: Sampling

Pattern matching Avoiding worst-case complexity

- Network
$\times$ No definite structure we can rely on
- Approximate matching may help but
- Subgraph isomorphism is hard to approximate
$\checkmark$ Structural properties of patterns
- Triangles and other small patterns DONE
- Trees
$\checkmark$ Statistical properties of network
Sampling $\longrightarrow$. Random graphs


## Pattern matching <br> Sampling - strategies

- Partitioning
- Match pattern in one or all partitions
- Assume that network has rather uniform structure
- Scale result to full network
- Worked for triangle counting, harder for larger patterns.
- Simulation (e.g. Fürer)
- Attempt several times to find single match
- assume network is sufficiently uniform
- average over iterations


## Fürer's algorithm ${ }^{1}$ for pattern matching in random graphs

- Input:
- Network $G$ drawn from $G(n, p)$
- Pattern H with a decomposition (see later)
- Output:
- |Emb $(H, G) \mid$, the number of images of $H$ in $G$.
- Complexity:
- Exact \& worst case: \#P-complete
- Exact for "most graphs": still untractable
- Approximate for most graphs : this algorithm (FPRAS)


## Pattern matching Fürer's algorithm - FPRAS

- Fully Polynomial Randomized Approximation Schema:
- Randomized algorithms outputting for almost every graph in polynomial time a solution with relative error of $\varepsilon$.
- A little more formally:
- An FPRAS is a randomized algorithm such that there is a polynomial $p(, \cdot)$ such that for every $\delta$ there is an $\mathrm{n}_{0}$ such that for all $n>n_{o}$ and $\varepsilon$, and for at least a fraction 1- $\delta$ of the graphs of size $n$, the algorithm outputs in time at most $p(n, 1 / \varepsilon)$ a solution within a factor $1 \pm \varepsilon$ of the correct solution.


## Pattern matching Fürer: Basic algorithm

Function unbiased_estimator(H,G) [estimates |Emb $(H, G) \mid]$ $(\varphi, P(\varphi))=$ Try_to_find_embedding(H,G);
IF $\varphi=$ failed
THEN return o;[No embedding found]
ELSE return 1/P $(\varphi)$; [return inverse
EndFunction
probability of $\varphi$ ]

Function count_embeddings(H,G, $\varepsilon$ )
$\mathrm{c}=\mathrm{o} ; \mathrm{s}=\mathrm{C} / \mathrm{\varepsilon}^{2}$;
For i = 1 .. s do c=c+unbiased_estimator(H,G); EndFor
Return c/s;
EndFunction

## Pattern matching Fürer: Basic algorithm

- Count_embeddings works correctly:
- Every embedding $\varphi$ is found with probability $P(\varphi)$ by Try_to_find_embeddings(H,G). With probability $1-\Sigma_{\varphi} P(\varphi)$ Try_to_find_embeddings fails
- Every embedding $\varphi$, is found once in $1 / P(\varphi)$ calls, and in that case, unbiased_estimator returns $1 / P(\varphi)$. Hence, $\varphi$ contributes 1 to each call to unbiased_estimator, on average.
- Hence, on average, unbiased_estimator returns the number of embeddings
- Task left: find a good Try_to_find_embedding


## Pattern matching Fürer - strategy

- Decompose vertex set of pattern
- match one partition at a time until complete
- Compute probability of finding this particular solution
- Show the overall algorithm converges sufficiently fast.
- The sum of a (very) large number of identical distributions becomes Gaussian. Standard deviation goes down with square of sample
- Whatever the sample size, we can always (with very small probability) have a large error


## Pattern matching

Fürer: finding embeddings
INPUT: $G, H$, partition $\left\{V_{1}, V_{2}, \ldots, V_{l}\right\}$ of $V(G)$
$X \leftarrow 1 ; \varphi_{0} \leftarrow\{ \}$
For $i=1 . . l$
$E \leftarrow\left\{\varphi \in \operatorname{Emb}\left(\cup_{\{j=1\}}^{i} V_{j}, G\right) \mid \varphi \supset \varphi_{i-1}\right\}$
$X_{i} \leftarrow|E|$
if $X_{i}=0$ then terminate and return (failed, 0 )
pick an embedding $\varphi_{i}$ uniformly at random from $E$
$\mathrm{X} \leftarrow X . X_{i}$
EndFor
Return $\left(\varphi_{l}, 1 / X\right)$

Fürer - Sample run (step 1.1)


$$
\begin{aligned}
& \text { - } X=1 ; \varphi_{0}=\{ \} \\
& \text { - } i=1
\end{aligned}
$$

Fürer - Sample run (step 1.2)


$$
\begin{aligned}
& \text { - } X=1 ; \varphi_{0}=\{ \} \\
& \text { - } i=1 \\
& X_{1}=|E|=8
\end{aligned}
$$

Fürer - Sample run (step 1.3)


- $i=1$
- $X_{1}=8$
- $\varphi_{1}=\{(1, a)\}$
- $X=8$

Fürer - Sample run (step 2.1)


Fürer - Sample run (step 2.2)


$$
\begin{aligned}
& \text { - } i=2 \\
& \text { - } X=8 ; \varphi_{1}=\{(1, a)\} i \\
& \text { - } X_{2}=|E|=3 \cdot 2=6 \\
& \text { - } X \leftarrow X \cdot X_{2}=8 * 6=48
\end{aligned}
$$

Fürer - Sample run (step 2.3)


$$
\begin{aligned}
& \text { - } i=2 \\
& \text { - } \varphi_{1}=\{(1, a)\} i \\
& \text { - } X_{2}=|E|=3 \cdot 2=6 \\
& \text { - } X=48 \\
& \text { - } \varphi_{2}= \\
& \{(1, a),(2, h),(3, c)\}
\end{aligned}
$$

Fürer - Sample run (step 3.1)


Fürer - Sample run (step 3.2)


- $i=3$
- $\varphi_{2}=\{(1, a) ;(2, h),(3, c)\} ;$
- $X=48$
- $X_{3}=|E|=2$
- $X=X \cdot X_{3}=48 * 2=96$

Fürer - Sample run (step 3.3)


- $i=3$
- $\varphi_{2}=\{(1, a) ;(2, h),(3, c)\} ;$
- $X=48$
- $\mathrm{X}_{3}=2 ; \mathrm{X}=96$
- $\varphi_{3}=$
$\{(1, a),(2, h),(3, c),(4, f)\}$

Fürer - Sample run (step 4.1)


Fürer - Sample run (step 4.2)


Fürer - Sample run - a solution


## Fürer - Sample run bis



- $i=3$
- $\varphi_{2}=\{(1, a) ;(2, h),(3, c)\} ;$
- $X_{3}=|E|=2$
- $X=X \cdot X_{3}=48 * 2=96$


## Fürer - Sample run bis



- $i=3$
- $\varphi_{2}=\{(1, a) ;(2, h),(3, c)\} ;$
- $X=96$
- $\varphi_{3}=\{(1, a),(2, h),(3, c)$,
$(4, b)\}$

Fürer - Sample run bis


## Pattern matching Fürer's theorem

- Important: When does it work?
- The algorithm is an FPRAS (for allmost all G) if:
- Partitioning of V(H) is a "ordered bipartite decomposition"
- $\mathrm{p}(\mathrm{n}) \mathrm{n}^{2} \rightarrow \infty$
- $\mathrm{np}^{0} / \Delta^{4} \rightarrow \infty$
- $E(H)^{3} n^{-4} p^{-2} \rightarrow 0$

Where

$$
v=\max (2, \gamma), \gamma=\max \{|E(F)| /(|V(F)|-2)|F \leq H,|V(F)| \geq 3\}
$$

## Pattern matching <br> Fürer: Simplified theorem

- It works if the count is not too small.
- Intuition:
- The frequency of very rare events is hard to measure (the "interesting" part of the sample where we actually observe something, is smaller)
- The worst case is where one doesn't know whether $|\operatorname{Emb}(H, G)|=0$ or $|\operatorname{Emb}(H, G)|=1$
- If we run the algorithm and we find some embeddings, our estimate will be rather accurate.
- If no "ordered bipartite decomposition", then we can still run this algorithm, but no FPRAS guarantee.


## Pattern matching; Fürer

## Ordered bipartite decomposition

- An ordered bipartite decomposition of $H$ is a partition $\left\{V_{1 \nu} V_{21} \ldots, V_{\beta}\right.$ of $V(H)$ such that
- Each $V_{i}(i=1 . . l)$ is an independent set in $H$
- $\forall i, v \in V_{i} \exists j$ such that $N_{H}(v) \subseteq U_{k<i} V_{k} \cup V_{j}$
- So if a neighbor of a vertex $v \in V_{i}$ is in $V_{j}$ with $j>i$, then no neighbors of $v$ must be in $V_{j^{\prime}}$ with $j^{\prime}>i$ and $j \neq j^{\prime}$.

Fürer - Graphs with a decomposition


- Cycles longer than 3 (see above running example)
- Bounded degree outerplanar graphs without triangle
- Trees
- Grids
- Not: triangles (but separate proof), ...


## Pattern matching

## Fixed parameter tractability

Pattern matching

## Avoiding worst-case complexity

- Network
$x$ No definite structure we can rely on
- Approximate matching may help but
- Subgraph isomorphism is hard to approximate
$\checkmark$ Structural properties of patterns
- Triangles and other small patterns DONE
" Trees
Fixed parameter tractability
$\checkmark$ Statistical properties of network
- Random graphs DONE


## Pattern matching <br> Fixed parameter tractability

- Classic complexity classes:
- Input size: $n$
- $P$ : Polynomial time $O\left(n^{d}\right)$ for some $d$
- Are all problems not in $P$ are hard: not $O\left(n^{d}\right)$ for some $d$ ? No, some may still be easier than others
- Fixed parameter tractability:
- Input has two parts of sizes: $n$ and $k$
- For $k$ fixed, the problem is tractable: $O\left(n^{d} f(k)\right)$
- May be acceptable if $k$ is small


## Pattern matching <br> Fixed parameter tractability

- Fixed parameter tractability:
- Input has two parts of sizes: $n$ and $k$
- For $k$ fixed, the problem is tractable: $O\left(n^{d} f(k)\right)$
- May be acceptable if $k$ is small
- Pattern matching:
- Network: size $n$
- Pattern: size $k$
- $k$ is usually small, $n$ may be large.
- So fixed parameter tractability is suitable for pattern matching!


## Pattern matching <br> Fixed parameter tractability

- Matching trees in network in

$$
O\left(m k 2^{k}\right)=O^{*}\left(2^{k}\right)
$$

- $k=V(P)$ : pattern size
- $m=E(D)$ : network size
- Randomized algorithm¹ which works well for practical pattern mining ${ }^{2}$
${ }^{1}$ R. Williams, Inf Proc Lett 2009
${ }^{2}$ A. Kibriya \& J.Ramon, DMKD 2013


## Pattern matching summary

- Basic operation for mining and learning
- Networks have no hard structure we can rely on
× Approximate matching may help but
- Subgraph isomorphism is hard to approximate
$\checkmark$ Structural properties of patterns
- Triangles and other small patterns
- Trees
$\checkmark$ Statistical properties of network
- Random graphs

Pattern matching - open problems

- Many matching problems for which fixed parameter approach could help
- More complex queries
- Implicit relations
- Big data (not in RAM)
- combine with indexing
- reduce passes over data (or samples)
- distributed approaches, ...


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Introduction
- Pattern matching
- Frequency
- Additional remarks
- Learning


## Frequency / support¹ measures The problem

- How frequent is pattern $P$ in network $D$ ?
- Why assign a "frequency" to a pattern?
- Popular criterion to measure relevance of pattern
" E.g. 40\% of respondents liked both movie m1 and m2.
- Way to represent association rules
" E.g. Of all respondents liking movie $m 1$ and $m 2,50 \%$ also liked $m 3$ (i.e. $40 \% * 50 \%=20 \%$ )
- Measure of statistical power
- E.g. We rolled the dice 100 times and observed 40 times a 6. It must be biased.


## Support measures <br> What do you want?

- Counting objects
- X\% of people own a house in the same region where they work.
- network unimportant, just count people



## Support measures <br> What do you want?

- Counting objects
- Performing statistics
- We rolled the dice 100 times and observed 40 times a 6. It is biased!
- We rolled the dice. 100 people observed a 6. Would it be biased?



## Support measures <br> What do you want?

- Counting objects
- Performing statistics
- Association rules
- If $X$ has a friend $Y$ such that $Y$ smokes, then with probability a\%, X smokes too
- If $X$ and $Y$ are friends and $Y$ smokes, then with probability b\%, X smokes too
- Different quantor/aggregator placement, probably $a \neq b$


## Support measures What do you want?

- What do you want?
- Counting objects
- Performing statistics
- Association rules
- If you know what you want, you're closer to knowing what to do. No measure is good in all cases.


## Support measures Overview

- Embedding-based
- Embedding count
- Image count
- Key-based
- Key image count
- Min-image
- Overlap-based
- Maximum independent set
- Minimum clique partition
- Intermediate measures and relaxations


## Support measures <br> Embedding-based

- Embedding-count: $|\operatorname{Emb}(P, D)|$
- Image-count: |Img $(P, D) \mid$

$$
|\operatorname{Emb}(P, D)|=|\operatorname{Aut}(P)| \cdot|\operatorname{Img}(P, D)|
$$

- E.g.: Among all triples $(X, Y, Z)$ such that $X$ and $Y$ are friends and $Z$ is a family member of $X$, the fraction of triples where $Y$ knows $Z$ is $a \%$.
- Embeddings may be concentrated in small part of network, e.g. large family where everyone is friend with each other.
- Not anti-monotonic, no pattern-mining pruning


## Support measures <br> Key-based

- Decide before the start of data mining what is the type of object of interest ("the primary key")
- E.g. We are interested in 'friends' relationships
- a\% of 'friends' relations are between colleagues.
- $b \%$ of friends have the same mother tongue,
- c\% of friend pairs ( $X, Y$ ) have at least one common friend $Z$,


## Support measures <br> Key-based

- Decide before the start on the "key".
- The "key" is a common subpattern of all patterns considered.
- There is a fixed, finite set of objects (all images/embeddings of the "key"), the network relations are not considered.
- Easy to count
- Anti-monotonic (good for pattern mining)
- Not all statistics are valid (e.g. the dice example)


## Support measures Key-based: dice example

- Key = dice observation
- Performing statistics:

- We rolled the dice. 100 people observeda6. Would it be biased?


4 images of Key
eqch showing a 6

## Support measures Min-image ${ }^{1}$

- minImage $(P, D)=$

$$
\min _{v \in V(P)}|\{\pi(w) \mid \pi \in \operatorname{Emb}(P, D)\}|
$$

- Allows for choosing each vertex as (singleton) key, giving a lower bound for each vertex-key-based frequency
- Anti-monotonic:
$P \leq Q \Rightarrow \operatorname{minImage}(P, D) \geq \operatorname{minImage}(Q, D)$


# Support measures <br> Overlap-based: model dependence 

- The easiest way to perform statistics is to have independent observations.
- How do we get as much independent observations as possible out of a network?
- Model the independences in "overlap graph".
- Caution: selecting independent observations is not necessarily a sample from the original distribution!


## Support measures <br> Overlap-based: Overlap graph

- Overlap graph $G_{P}^{D}$ :
- $V\left(G_{P}^{D}\right)=\operatorname{Img}(P, D)$
- $\left(g_{1}, g_{2}\right) \in E\left(G_{P}^{D}\right)$ if images $g_{1}$ and $g_{2}$ overlap
- What is overlap?
- Two occurrences of a pattern overlap if we can't consider them independent in the context of the statistics we are doing
- Vertex-overlap: $\left(g_{1}, g_{2}\right) \in E\left(G_{P}^{D}\right) \Leftrightarrow V\left(g_{1}\right) \cap V\left(g_{2}\right) \neq \emptyset$
- Edge-overlap: $\left(g_{1}, g_{2}\right) \in E\left(G_{P}^{D}\right) \Leftrightarrow E\left(g_{1}\right) \cap E\left(g_{2}\right) \neq \varnothing$
- Other options, e.g. Harmful overlap ${ }^{1}$


## Support measures <br> Overlap-based: What is overlap?

- Dice example:
- Overlap if the Roll is the same.
- We are interested in dice1, so naturally all embeddings will contain dice1.



## Support measures <br> Overlap-based: Court example

- Vertices: case, judges, (prodeo) lawyers, party
- Edges between case-judge, case-lawyer, case-party



## Independent set

- Wis an independent set of $H$ iff
- $W \subseteq V(H)$
- There are no $v, w \in W$ such that $(v, w) \in E(H)$


Independent set of size 2


Independent set of size 1

## Support measures Overlap-based: MIS measure

- The size of a Maximum Independent Set (MIS) of the overlap graph $G_{P}^{D}$ of pattern $P$ in network $D: \operatorname{MIS}(P, D)=\operatorname{MIS}\left(G_{P}^{D}\right)$


$$
\operatorname{MIS}\left(G_{P}^{D}\right)=3
$$

- (dice1,obs=6,Person1)
(dice1,obs=6,Person2)
- (dice1,obs=6,Person2)
- (dice1,obs=3,Person2)


## Support measures Overlap-based: MIS measure

- The size of a Maximum Independent Set (MIS) of the overlap graph $G_{P}^{D}$ of pattern $P$ in network $D: \operatorname{MIS}(P, D)=\operatorname{MIS}\left(G_{P}^{D}\right)$



## Support measures <br> Overlap-based: more measures

- Maximum independent set idea:
© allows to measure overlap and extract independent observations
© Anti-monotonic
© NP-hard to compute
© Possibly ignores too much information
- Does the overlap graph allow for other measures?


## Support measures <br> Overlap-based: more measures

- Requirements for feasible measure:
- Anti-monotonic in pattern:
- $p \leq P \Rightarrow f(p, D) \geq f(P, D)$
- Monotonic in network:
- $D \leq D^{\prime} \Rightarrow f(P, D) \leq f\left(P, D^{\prime}\right)$
- Normalized
- If there are $n$ independent (non-overlapping) observations (overlap graph = $n$ isolated vertices), then support is $n$.


## Support measures

Overlap-based: more measures

- If $f$ is a function on the overlap graph, and $f$ is feasible measure, then:

$$
M I S(P, D) \leq f(P, D) \leq M C P(P, D)
$$

- where $M C P\left(G_{P}^{D}\right)$ is the minimum clique partition number of the overlap graph, another NP-hard to compute number.


## Support measures <br> Overlap-based: more measures

- Fortunately, several efficiently computable functions are between MIS and MCP.
- Lovasz theta': $\vartheta$
- feasible measure²; computable with semidefinite program (SDP), which is still rather expensive
- MIS-relaxation3: s
- Is a feasible measure3; computable with linear program (LP), hence efficiently.
- We have $M I S \leq \vartheta \leq s \leq M C P$

[^5]
## Support measures Summary

| Measure | Anti-Monotonic? | Statistics? | Efficient? |
| :--- | :---: | :---: | :---: |
| Embedding count | $x$ | $x$ | $\checkmark$ |
| Image count | $x$ | $x$ | $\checkmark$ |
| key image count | $\checkmark$ | $x$ | $\checkmark$ |
| min-image | $\checkmark$ | $x$ | $\checkmark$ |
| Overlap-MIS | $\checkmark$ | $\checkmark$ | $x$ |
| Overlap-MCP | $\checkmark$ | $?$ | $\times$ |
| Overlap- $\vartheta$ | $\checkmark$ | $\checkmark$ | $?$ |
| Overlap-s | $\checkmark$ | $\checkmark$ | $\checkmark$ |

## Frequency - open problems

- Combine pattern matching and frequency
- Exploit network structure to increase speed (methods up to now don't)


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Introduction
- Pattern matching
- Frequency
- Additional remarks
- Learning


## Expected number of embeddings

- Let $D \sim G(n, p)$, then
$|E m b(P, D)|=n^{|V(P)|} p^{|E(P)|}(1-p)^{\frac{n(n-1)}{2}-|E(P)|}$
- For small $p$ :

$$
|E m b(P, D)|=n^{|V(P)|} p^{|E(P)|}
$$

- For trees:

$$
|\operatorname{Emb}(P, D)|=(n p)^{|V(P)|} / p
$$

- Often $D$ is connected and $n p>1$


## Expected number of embeddings

- \# of expected embeddings grows with pattern size for sparse patterns
- Denser patterns may be easier to interprete
- Also patterns less frequent than expected may be of interest.
- This also happens with itemsets: items may be correlated, uncorrelate or anti-correlated


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Learning
- Introduction
- Learning from non-independent examples
- Temporal models


## Learning - introduction

- Popular learning ideas:
- connected vertices have similar target value
- correlation between features and target value
" more classic feature-to-target supervised learning
- "Dual space" idea: one feature per vertex $u$, is 1 for vertices connected to that vertex $u$ (else o).
- If individuals are important (but not many features are known)


## Learning - introduction Similar to your neighbor

- Semi-supervised learning
- E.g.: try to minimize the number of edges with on both sides different labels
- E.g. target values tend to average of neighbors.
- Manifold embedding:
- Assign all vertices a coordinate such that connected vertices are close together (and notconnected vertices are far apart)


## Learning - Introduction <br> From feature to target value

- Learning tasks¹:
- Vertices / edges / ...
- (existence)prediction / labeling / weighting /feature construction / ...
vertex/edge structure/ feature prediction example
- Supervised learning
- Input:
" vertex/edge to predict
- Neighborhood
- Output:
- Label or existence
- But how about the classic i.i.d. assumptions?


## Learning - Prediction in a fixed network

- Most common setting:
- Fixed network $D$
- training and test vertices (edges) in $D$
- But what if the network changes (e.g. an influential node is added/deleted)?
- Causal patterns remain the same
- Correlation patterns may change significantly


## Learning - Can’t distinguish individual and its features




What is the rule?
Does everyone follow the class of its neighbors? Is v1 very influential?
Is class + if there are green neighbors?

## Learning - network-specific challenges

- Suppose the same rules stay true, but new nodes/edge (same distribution). Distinction viral/features may matter
- Movie database:
- Persons from a fixed distribution
- Movies from a fixed distribution
- Persons watch movies
- A few new persons and movies are added


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Learning
- Introduction
- Learning from non-independent examples
- Temporal models


## Learning from dependent examples

- Members of a network are dependent
- Typical assumptions of learning algorithms don't hold, in particular that
- Examples are independently and identically drawn (i.i.d.)


## Movie rating example

- Movie rating
- Obj: Movie (genre, duration, actor popularity)
- Obj: Person (age, gender, ...)
- Obj: Screening (location, time, ...)
- Target: Rating
- Several ratings per person / movie / cinema


## Lawsuit example

- Lawsuits:
- Obj: Person
- Obj: Lawyer
- Obj: Judge
- Example: case
- Target: outcome
- Judges handle several cases, persons may be involved in several cases


## Learning from pattern

 features- Each example is an embedding of a pattern
- MovieRating: (Movie, Person, Cinema, Rating)
- Lawsuit: (Case, Person, Judge, Outcome)
- Examples overlap:
- See also "support measures"
- We call them networked examples


## Representing networked examples

- Several alternative equivalent representations:
- Every example is represented with a vertex connected to the participating objects
- Every example is represented with a hyperedge, containing all participating + all relevant objects.



# Learning from networked / dependent examples 

- Tasks:

1. Elementary statistics, confidence intervals, hypothesis testing, ...
2. Learning, generalization guarantees

- Models:
a. bounding covariance between dependent examples
b. modeling how examples are dependent
- Combinations: 1a and 2b


# Bounding covariance of examples and hypothesis testing 

- (Wang, Neville, Gallagher, Eliassi-Rad, ECML/PKDD-2011):
- vertices are examples
- edges indicate a bounded covariance
- safe correction for statistical significance tests
- safe upper bound for variance on sums etc.
- Upper bound for variance can be an important tool in proving generalization guarantees.


## Bounding covariance of examples

- $n$ independent random variables $\left\{X_{i}\right\}_{i=1}^{n}$ with variance $\sigma^{2}$ : Variance on $\sum_{i=1}^{n} X_{i}$ is $n \sigma^{2}$
- $n$ identical random variables $\left\{X_{i}\right\}_{i=1}^{n}$ with variance $\sigma^{2}$ : Variance on $\sum_{i=1}^{n} X_{i}$ is $n^{2} \sigma^{2}$


## Bounding covariance of examples

- $n$ networked examples
- No edge between $X_{i}$ and $X_{j}=$ independent:

$$
E\left[\left(X_{i}-E\left[X_{i}\right]\right)\left(X_{j}-E\left[X_{j}\right]\right]=0\right.
$$

- Edge $\left(X_{i}, X_{j}\right) \in E(D)$ between $X_{i}$ and $X_{j}=$ bounded covariance $E\left[\left(X_{i}-E\left[X_{i}\right]\right)\left(X_{j}-E\left[X_{j}\right]\right] \leq \gamma\right.$
- Variance on $\sum_{i=1}^{n} X_{i}$ is
$\sum_{i, j} E\left[\left(X_{i}-E\left[X_{i}\right]\right)\left(X_{j}-E\left[X_{j}\right]\right] \leq|E(D)| \gamma+n \sigma^{2}\right.$


# Learning from networked / dependent examples 

- Tasks:

1. Elementary statistics, confidence intervals, hypothesis testing, ...
2. Learning, generalization guarantees

- Models:
a. bounding covariance between dependent examples
b. modeling how examples are dependent
- Combinations: 1a and 2b


# Variance, significance, effective sample size 

- Effective sample size of a given set of networked examples is $n$ iff it contains as much information (for the task at hand, e.g. learning or hypothesis testing) as a set of $n$ i.i.d. examples ${ }^{1}$.


## Probably approximately <br> correct (PAC) structure

- PAC: with probability $1-\delta$ the loss is bounded by $\varepsilon$ where

$$
\delta=\exp \left(\frac{-m(S) \epsilon^{2}}{C_{1}+C_{2} \epsilon}\right)
$$

- with $m(S)$ the effective sample size of training set $S$. Higher $m(S)=$ better
- i.i.d. sample $S, m(S)=|S|=$ best possible


## Independence assumptions

- Weaker form of i.i.d
- But not arbitrary
- arbitrary $\Rightarrow$ no bound possible


## Independence assumptions: i.i.d. vertex features

- Edges are fixed.
- The features of every vertex are drawn i.i.d. (not even depending on the edges).


1. Choose edges (possibly very dependently)

2. Draw vertex
features (don't
look at edges)

## Independence

## assumptions applied

## YES

- Sneak preview
- Randomized trial: patients are assigned randomly to set of treatment params
- Cases are assigned randomly to judges


## NO

- Select movie based on genre, or with friends
- Patients go to closeby hospital or to hospital recommended by their friends
- Judges handle cases connected to their existing cases


## Training set measures

- Overlap (hyper)graph G
- vertices are objects
- (hyper)edges are examples
- Training set $S \subseteq E(G)$
- Measures $m(S)$ of training set $m(S)$ :

$$
\max \left(n_{E Q W}, n_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq s \leq n_{M S C}
$$

## Approach 1:

## Equal-weight (EQW)

## $\max \left(\boldsymbol{n}_{E Q W}, \boldsymbol{n}_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq \boldsymbol{s} \leq n_{M S C}$

 $\uparrow$all examples get same weight
(Janson 2004) \& (Usunier 2005)

$$
\delta \propto \exp \left(\frac{-n_{E Q W} \epsilon^{2}}{C_{1}+C_{2} \epsilon}\right)
$$

With $n_{E Q W}=\frac{|S|}{\chi^{*}(G)}$
and $\chi^{*}(G)$ fractional edge chromatic


$$
\chi^{*}(G)=3 ; n_{\text {EQW }}=\frac{6}{3}=2
$$

## Approach 2:

## Independent set (IND)

$$
\max \left(n_{E Q W}, n_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq s \leq n_{M S C}
$$ 1

We find $n_{I N D}$ independent examples

$$
\delta \propto \exp \left(\frac{-n_{I N D} \epsilon^{2}}{C_{1}+C_{2} \epsilon}\right)
$$

With $n_{I N D}=|S|$ (examples in $S$ independent!)


## Approach 3:

## Maximum independent set

$$
\max \left(n_{E Q W}, n_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq s \leq n_{M S C}
$$

$$
\uparrow
$$

Maximum independent set of examples

$$
n_{M I S}=|M I S(G)|
$$

|MIS (G)|hard to approximate!
$\Rightarrow$ no const lower bound for $\frac{n_{I N D}}{n_{M I S}}$
For some $G, \frac{n_{E Q W}}{n_{M I S}}=2 /|S|$


## Minimum clique partition number

$$
\max \left(n_{E Q W}, n_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq s \leq n_{M S C}
$$

Minimum set cover

$$
\begin{gathered}
n_{M S C}=|M S C(G)| \\
n_{M S C} \leq\left(k-1+\frac{1}{k}\right) n_{M I S}
\end{gathered}
$$

$|M S C(G)|$ too is hard to compute


$$
k-1+\frac{1}{k}=2.5
$$

## Approach 4: MIS-relaxation

$$
\max \left(n_{E Q W}, n_{I N D}\right) \leq n_{M I S} \leq \vartheta \leq s \leq n_{M S C}
$$

LP relaxation of MIS
(Wang \& Ramon, DMKD 2013)
Graph pattern support measure

- Anti-monotonic, Normalized
- Linear program $\rightarrow$ efficient



## networked PAC

- PAC: $\mathrm{P}($ Loss $\leq \epsilon) \geq 1-\delta$ where

$$
\delta=\exp \left(\frac{-\boldsymbol{s} \epsilon^{2}}{C_{1}+C_{2} \epsilon}\right)
$$

- with $\boldsymbol{s}$

$$
S=\max \left(\sum_{i=1}^{|S|} w_{i}\right)
$$

subject to

$$
\forall v \in V(G): \sum_{e_{i}: v \in e_{i}} w_{i} \leq 1
$$

## Networked PAC

- Influence of each factor is at most 1:
max $S$

$$
s=w_{1}+w_{2}+w_{3}+w_{4}+w_{5}+w_{6}
$$

s.t.

$$
\begin{array}{lr}
v_{1}: & w_{1}+w_{2} \leq 1 \\
v_{3}: & w_{1}+w_{3} \leq 1 \\
v_{2}: & w_{2}+w_{3} \leq 1 \\
v_{4}: & w_{4}+w_{5}+w_{6} \leq 1
\end{array}
$$



## Technical elaboration

- Chernoff bound for weighted sums:

For $\boldsymbol{X}_{\boldsymbol{i}}(\boldsymbol{i}=1 . . \boldsymbol{n})$ independent random variables,
$E\left[X_{i}\right]=0 ;\left|X_{i}\right| \leq a_{i}+M ; X=\sum_{i} X_{i}:$
$P\left(\sum_{i=1}^{n} X_{i} \geq n \epsilon\right) \leq \exp \left(\frac{-n \epsilon^{2}}{\operatorname{Var}(X)+\sum_{i} a_{i}+M \epsilon / 3}\right)$

- Let $X_{i}=\xi\left(\{\boldsymbol{\phi}(v)\}_{v \in e_{i}}\right)$ and $\forall v: \sum_{e_{i}: v \in e_{i}} w_{i} \leq 1$, then the above Chernoff inequality still holds


## Learning from non-independent examples: Summary

- Networks modeling relations induce dependencies between objects and examples
- Modeling such dependencies is useful to
- better understand the learning setting
- get more statistical power from the data
- Upper bound correlation between examples
- Model common factors of examples


## Learning from non-independent data: Open problems

- Can we formalize \& structure models for learning?
- How to extract most statistical value from data?
- How to take intervention into account (very important for applications)?


## Contents

- Introduction
- Networks from different points of view
- Patterns \& pattern mining
- Learning
- Introduction
- Learning from non-independent exammples
- Temporal models


## Temporal models

- Temporal model = probability distribution mapping a network on a new network, i.e. $h: \mathcal{G} \times \mathcal{G} \rightarrow[0,1]$ s.t. $\forall D \in \mathcal{G}, \sum_{D^{\prime} \in \mathcal{G}} h\left(D, D^{\prime}\right)=1$
- Results in Markov chain
- When starting from empty network: generative model
- Often one attempts to find a simple rule, when provided to all individuals of a group producing an interesting/real-life pattern
- Communities, powerlaws, emerging structures, ...


## Temporal models

- Global temporal models
- How will communities evolve?
- How will global network properties evolve?
- Models aiming at emerging behavior
- Local temporal models
- How will individual nodes/edge/local neighborhoods evolve?


## Global temporal models Examples

- How will communities evolve?
- Research topics emerging and disappearing ${ }^{1}$
- Online groups emerging and disappearing ${ }^{2}$

1 Ferlez et al. ICDE 2008
2 Kairam, Wang \& Leskovec WSDM 2012

## Models for emerging or assymptotic patterns - Examples

- Theory: Barabasi-Albert: Preferential attachment
- Complex systems:
- Economics (e.g. company fusions¹)
- Between networks and physics: Groups of animals (formations of flying birds²)
- No systematic integration with DM/ML

[^6]
## Local temporal models

- Local models:
- Link prediction: what could exist may get known as existing soon.
- Learn from temporal data
- Few combined local/global approaches
- Emerging behavior shown by simulation (Complex systems)
- Model microscopic social network evolution, show that it has powerlaw assymptotics ${ }^{1}$


## Evolving large networks: <br> Types of data

## Snapshot

- At one point in time


## Temporal data

- A log of the evolution
- hard: nobody logged all events in the creation of the internet up to now



## Temporal models:

Types of data

- Snapshot:
- Historical information may be missing
- Can we still detect traces of evolution in the network?
- Sometimes yes, e.g. phylogenetic trees


## Temporal models: Summary

- Scale of evolution:
- local
- global
- assymptotical
- Types of data:
- Snapshot
- Temporal

Temporal data:

## Open problems

- Integration of local and global/assymptotic levels?
- Can we learn dynamics from snapshots?


## Conclusions

- Several domains study networks from different points of view (and can learn from each other)
- This tutorial: local level
- Others: global level
- Progress towards integration

Questions?


[^0]:    ${ }^{1}$ Ramon \& Nijssen, JMLR2008
    ${ }^{2}$ Nijssen \& Kok 2004

[^1]:    \#P : counting problems $f$ such that there is a polynomial-time non-deterministic Turing machine for which upon input $x$ the number of accepting states equalls $f(x) . \# P \supseteq N P$

[^2]:    ${ }^{1}$ E.M.Luks, J Computer \& System Sciences 25(1), 1982
    ${ }^{2}$ Horvath \& Ramon, DMKD 21 (3), 2010
    ${ }^{3}$ Horvath \& Ramon, TCS 411, 2010

[^3]:    ${ }^{1}$ Alon et al. 1997

[^4]:    1 Shen-Orr et al. Nat. Genet. 2002
    ${ }^{2}$ Bhuiyan et al. ICDM 2012

[^5]:    ${ }^{1}$ D. Knuth, Electr. J. Combin 1994
    ${ }^{2}$ Calders et al. DMKD 2011
    ${ }^{3}$ Wang \& Ramon DMKD 2013

[^6]:    ${ }^{1}$ Garnett \& Mollan, ECCS 2012
    ${ }^{2}$ Hemelrijk \& Hildenbrandt, ECCS 2012

