

Prediction

Analysis of complex interconnected data







Quick Notes

- Second assignment is due on Oct 4th
 - Submit 2 files (report.pdf, code.zip) as a Group (pairs or two or individual) in Mycourses
- Tue., Oct. 19, 2021: Project Proposal Presentations
 - Why & What: Introduction and Motivation, Related Work, Problem Definition, Dataset Description
 - Writeup: 2 pages, due Oct 20th [8pt]
 - Presentation: 2 mins (2-3 slides), slides due Oct 18th [2pt]
 - Email the slides to the course email, use Google Slides
 - We will merge them all together, and you will go over it in cla
- Any questions?

Deadlines

- assignment 1 due on Sep. 20th
- assignment 2 due on Oct. 4th
- assignment 3 due on Oct. 18th
- project proposal slides due on Oct. 18th
- project proposal due on Oct. 20th
- Reviews (first round) due on Oct. 27th
- project proposal slides due on Nov. 3rd
- project progress report due on Nov. 5th
- Reviews (second round) due on Nov. 12th
- project final report slides due on Nov. 29th
- project final report due on Dec. 7th
- Reviews (third round) due on Dec. 14th
- project revised report and rebuttal due on Dec. 20th

WHAT

note: dates are tentative, subject to change

Common prediction tasks

- Link Prediction
- Node Classification
- Graph Classification

What is unsupervised node classification?

Examples:

https://paperswithcode.com/task/link-prediction

https://paperswithcode.com/task/node-classification

https://paperswithcode.com/task/graph-classification

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Given a graph G(V,E) predict future/missing links between nodes

- Modeling of network evolution
- Predict likely interactions, not explicitly observed (e.g. terrorist network monitoring)

See all

Mahdi Tayakoli

Professor (Robotics) at

the University of Alberta

O 19 mutual connections

Connect

Masoud Ardakani

Professor of Electrical

Engineering (Universi..

OD 14 mutual connections

Connect

• Link recommendation: "friend" suggestion in social networks



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Predicting missing links

Only a subset of edges are observed

Sparse \Rightarrow searching for O(n) needles in a $\Theta(n^2)$ haystack



From Clauset's Slides

Score(i,j)

local topological predictors

- Number of common neighbors
- Number of shortest paths
- Product of degree
- Same cluster
- etc.



From Clauset's Slides



Score(i,j)

local topological predictors

- Number of common neighbors
- Number of shortest paths
- Product of degree
- Same cluster
- etc.



From Clauset's Slides



Jaccard coefficient

 $score(i, j) = Jaccard(i, j) + Uniform(0, \epsilon)$

What happens to the network if we add edges based on this?

 $\operatorname{Jaccard}(i,j) = \frac{|\nu(i) \cap \nu(j)|}{|\nu(i) \cup \nu(j)|}$

From Clauset's Slides



Jaccard coefficient

score(i, j) = Jaccard(i, j) + Uniform(0, ε)

What happens to the network if we add edges based on this? Closes triangles



Example: Jaccard(i, j), what is it for this example?

From Clauset's Slides

 $\operatorname{Jaccard}(i,j) = \frac{|\nu(i) \cap \nu(j)|}{|\nu(i) \cup \nu(j)|}$



Jaccard coefficient

 $\operatorname{Jaccard}(i,j) = \frac{|\nu(i) \cap \nu(j)|}{|\nu(i) \cup \nu(j)|}$

 $score(i, j) = Jaccard(i, j) + Uniform(0, \epsilon)$

What happens to the network if we add edges based on this? Closes triangles



Example: Jaccard(i, j) of **0.50** (3/6) vs **0.091** (1/11)

Score(i,j)

local topological predictors

- Number of common neighbors
- Number of shortest paths
- Product of degree
 - nodes with high degrees are likely themselves to be connected 0
- Same cluster
- etc.



From Clauset's Slides

degree product: score(i, j) = $d_i d_i$ + Uniform(0, ϵ)





actual	G =	(V, E)
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observed $G_\circ = (V, E_\circ)$

i	j	Jaccard $score(i, j)$	i	j	$\begin{array}{c c} \text{degree product} \\ \text{score}(i, j) \end{array}$
4	5	1/2 + r	1	4	4+r
2	3	1/2 + r	1	6	4+r
3	6	1/3 + r	3	6	4+r
1	4	1/3 + r	1	5	2+r
1	5	r	2	3	2+r
1	6	r	2	6	2+r
2	6	r	2	4	2+r
2	4	r	3	5	2+r
2	5	r	4	5	2+r
3	5	r	2	5	1+r

From Clauset's Slides



degree product: score(i, j) = $d_i d_i$ + Uniform(0, ϵ)



From Clauset's Slides

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Measuring performance for ranked output

We can consider a threshold and convert it to a binary classification

{every i, j is either a missing link or not}

Accuracy = $(TP+TN) / (P+N)$)
Precision = TP / RP	
Recall = TP / P	{also called sensitivity}
F1 score = 2. Precision x Reca	ll / (Precision + Recall)
	{Harmonic mean}
Miss rate = FN / P	
Fallout = FP / N	{also called false positive rate}
False discovery rate = FP / R	P
Selectivity = TN / N	{also called specificity}
False omission rate = FN / RN	N
Negative predictive value = T	N / RN

	1	Σ	
	TP	FP	RP
Results	FN	TN	RN
Σ	Р	N	



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Measuring performance for ranked output

We can consider a threshold and convert it to a binary classification

{every i, j is either a missing link or not}

	ר	Σ	
	TP	RP	
Results	FN	TN	RN
Σ	Р	N	

Measures depend on the threshold

The tradeoff between precision and recall



Threshold invariant: ROC & AUC

Receiver Operating Characteristic (ROC) as a function of prediction threshold

TPR(t) = TP(t)/P (recall, sensitivity at t) FPR(t) = FP(t)/N (fallout, false alarm at t)

Area Under the Curve (AUC) of ROC

gives the probability of ranking a random positive edge higher than a random negative edge



Threshold invariant: ROC & AUC, example

binary classification \Rightarrow every candidate i, j is either a missing link or not



Measuring performance

binary classification \Rightarrow every candidate i, j is either a missing link or not

TPR = TP/P (**recall**, sensitivity) Links above threshold / Total Positives

FPR = FP/N (**fallout**, false alarm) Non-links above threshold / Total Negatives

TPR = ?, FPR = ?

TPR = ?, FPR = ?

	Truth	Σ	
results	TP = 2	FP = 2	RP = 4
	FN = 0	TN = 6	RN = 6
Σ	P = 2	N = 8	

	Truth	Σ				
results	TP = 0	FP = 4	RP = 4			
1030113	FN = 2	RN = 6				
Σ	P = 2	N = 8				
degree product						

Jaccard

		-i	$_{j}$	Jaccard $score(i, j)$	i	j	degree product $score(i, j)$
		4	5	1/2 + r	1	4	4+r
× ×	4	2	3	1/2 + r	1	6	4+r
<u>.=</u>		3	6	1/3 + r	3	6	4+r
threshold	_	-1	4	1/3 + r	1	5	2 + r
	- 1	-1	5	r	2	3	2+r
		1	6	r	2	6	2+r
×		2	6	r	2	4	2+r
i	_	2	4	r	3	5	2+r
Q		2	5	r	4	5	2+r
C		3	5	r	2	5	1+r

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Measuring performance

binary classification \Rightarrow every candidate i, j is either a missing link or not

TPR = TP/P (**recall**, sensitivity) Links above threshold / Total Positives

FPR = FP/N (**fallout**, false alarm) Non-links above threshold / Total Negatives

TPR = 2/2 = 1, FPR = 2/8=0.25

	Truth	Σ				
results	TP = 2	FP = 2	RP = 4			
resolts	FN = 0	TN = 6	RN = 6			
Σ	P = 2	N = 8				
Jaccard						

TPR = 0/2 = 0, FPR = 4/8=0.5

	Truth	Σ				
results	TP = 0	FP = 4	RP = 4			
1630113	FN = 2	RN = 6				
Σ	P = 2	N = 8				
degree product						

			Jaccard			degree product
	i	j	score(i, j)	i	j	$\operatorname{score}(i, j)$
	4	5	1/2 + r	1	4	4+r
×	2	3	1/2 + r	1	6	4+r
<u> </u>	3	6	1/3 + r	3	6	4+r
threshold	-1	4	1/3 + r	1	5	2 + r
	1	5	r	2	3	2+r
	1	6	r	2	6	2+r
×	2	6	r	2	4	2+r
i	2	4	r	3	5	2+r
ō	2	5	r	4	5	2+r
2	3	5	r	2	5	1+r

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Threshold invariant: ROC & AUC, example

binary classification \Rightarrow every candidate i, j is either a missing link or not



Threshold invariant: ROC & AUC, example

binary classification \Rightarrow every candidate i, j is either a missing link or not

rate .



actual G = (V, E)





observed $G_{\circ} = (V, E_{\circ})$

i	j	Jaccard $score(i, j)$	i	j	$\frac{\text{degree product}}{\text{score}(i, j)}$
4	5	1/2 + r	1	4	4+r
2	3	1/2 + r	1	6	4+r
3	6	1/3 + r	3	6	4+r
1	4	1/3 + r	1	5	2+r
1	5	r	2	3	2+r
1	6	r	2	6	2+r
2	6	r	2	4	2+r
0.6 0.2 0.2 0.2 0.2 0.2 0.4 False positi	0.6 ve rate (0.8 1	True positive rate (TPR) 90 000		0.2 0.4 0.6 0.8 False positive rate (FPR)
Jaccard coefficie	ent. A	UC = 1.00	c	legre	e product. $AUC = 0.31$

From Clauset's Slides

Topological Link Predictors

	CN	common neighbors of i, j					
	SP	shortest path between i, j					
	LHN	Leicht-Holme-Newman index of neighbor sets of i, j					
	PPR	j-th entry of the personalized page rank of node i					
	PA	preferential attachment (degree product) of i, j					
	JC	Jaccard's coefficient of neighbor sets of i, j					
	AA	Adamic/Adar index of <i>i</i> , <i>j</i>					
	RA	resource allocation index of i, j					
	LRA	entry i, j in low rank approximation (LRA) via singular value decomposition (SVD)					
	dLRA	dot product of columns i and j in LRA via SVD for each pair of nodes i, j					
	mLRA	average of entries i and j 's neighbors in low rank approximation					
	LRA-approx	an approximation of LRA					
	dLRA-approx	an approximation of dLRA					
mLRA-approx an approximation of r		an approximation of mLRA					
	LCC_i, LCC_j	local clustering coefficients for i and j					
	AND_i, AND_j	average neighbor degrees for i and j					
	$SPBC_i, SPBC_j$	shortest-path betweenness centralities for i and j					
	CC_i, CC_j	closeness centralities for i and j					
	DC_i, DC_j	degree centralities for i and j					
	EC_i, EC_j	eigenvector centralities for i and j					
KC _i , KC _j Katz cenLNT _i , LNT _j local nun		Katz centralities for i and j					
		local number of triangles for i and j					
	PR_i, PR_j	Page rank values for i and j					
	LC_i, LC_j	load centralities for i and j					

There are many **alternatives** to give a score to a given pair of nodes (i,j)

Many also used as similarity measures since homophily is a strong force in link creation

Ghasemian A, Hosseinmardi H, Galstyan A, Airoldi EM, Clauset A. Stacking Models for Nearly Optimal Link Prediction in Complex Networks. PNAS 2020.

See also A Survey of Link Prediction in Complex Networks, 2015 & Link prediction in complex networks: A survey, 2011

Common neighbour is a decent predictor

Random

Graph distance

Common neighbor



Sums over size inverse of log of degree of common neighbours

 $\sum_{z \in \Gamma(x) \cap \Gamma(y)} \frac{1}{\log |\Gamma(z)|}$

Liben-Nowell D, Kleinberg J. The link-prediction problem for social networks. Journal of the American society for information science and technology. 2007 May;58(7):1019-31.

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We can learn to link

Wang P, Xu B, Wu Y, Zhou X. Link prediction in social networks: the state-of-the-art. Science China Information Sciences. 2015 Jan 1;58(1):1-38.



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Learn the features as well

Instead of explicit features, automatically learn a "heuristic" that suits the current network e.g. by extracting a local enclosing subgraph around it, and uses a GNN to learn general graph structure features for link prediction



Table 1: Comparison with heuristic methods (AUC).

Data	CN	Jaccard	PA	AA	RA	Katz	PR	SR	ENS	WLK	WLNM	SEAL
USAir	93.80±1.22	89.79±1.61	88.84 ± 1.45	95.06±1.03	95.77±0.92	92.88±1.42	94.67±1.08	78.89 ± 2.31	88.96 ± 1.44	96.63±0.73	95.95 ± 1.10	96.62±0.72
NS	94.42±0.95	94.43±0.93	68.65 ± 2.03	94.45 ± 0.93	94.45±0.93	94.85 ± 1.10	94.89 ± 1.08	94.79±1.08	97.64±0.25	98.57±0.51	98.61±0.49	98.85±0.47
PB	92.04±0.35	87.41±0.39	90.14 ± 0.45	92.36 ± 0.34	92.46±0.37	92.92 ± 0.35	93.54 ± 0.41	77.08 ± 0.80	90.15 ± 0.45	93.83±0.59	93.49 ± 0.47	94.72±0.46
Yeast	89.37±0.61	89.32±0.60	82.20 ± 1.02	89.43±0.62	89.45±0.62	92.24±0.61	92.76±0.55	91.49 ± 0.57	82.36 ± 1.02	95.86±0.54	95.62±0.52	97.91±0.52
C.ele	85.13±1.61	80.19 ± 1.64	74.79 ± 2.04	86.95 ± 1.40	87.49 ± 1.41	86.34 ± 1.89	90.32±1.49	77.07 ± 2.00	74.94 ± 2.04	89.72±1.67	86.18 ± 1.72	90.30±1.35
Power	58.80 ± 0.88	58.79 ± 0.88	44.33 ± 1.02	58.79 ± 0.88	58.79 ± 0.88	65.39±1.59	66.00 ± 1.59	76.15 ± 1.06	79.52 ± 1.78	82.41 ± 3.43	84.76 ± 0.98	87.61±1.57
Router	56.43±0.52	56.40±0.52	47.58 ± 1.47	56.43±0.51	56.43±0.51	38.62±1.35	38.76±1.39	37.40±1.27	47.58 ± 1.48	87.42 ± 2.08	94.41±0.88	96.38±1.45
E.coli	93.71±0.39	81.31 ± 0.61	91.82 ± 0.58	95.36 ± 0.34	95.95±0.35	93.50 ± 0.44	95.57 ± 0.44	62.49 ± 1.43	91.89 ± 0.58	96.94±0.29	97.21±0.27	97.64±0.22

Table 2: Comparison with latent feature methods (AUC).

Data	MF	SBM	N2V	LINE	SPC	VGAE	SEAL
USAir	94.08±0.80	94.85±1.14	91.44±1.78	81.47±10.71	74.22 ± 3.11	89.28±1.99	97.09±0.70
NS	74.55 ± 4.34	92.30 ± 2.26	91.52 ± 1.28	80.63 ± 1.90	89.94 ± 2.39	94.04 ± 1.64	97.71±0.93
PB	94.30±0.53	93.90 ± 0.42	85.79 ± 0.78	76.95 ± 2.76	83.96±0.86	90.70±0.53	95.01±0.34
Yeast	90.28 ± 0.69	91.41 ± 0.60	93.67±0.46	87.45 ± 3.33	93.25 ± 0.40	93.88±0.21	97.20±0.64
C.ele	85.90 ± 1.74	86.48 ± 2.60	84.11±1.27	69.21 ± 3.14	51.90 ± 2.57	81.80 ± 2.18	89.54±2.04
Power	50.63 ± 1.10	66.57 ± 2.05	76.22 ± 0.92	55.63 ± 1.47	91.78±0.61	71.20 ± 1.65	84.18 ± 1.82
Router	78.03±1.63	85.65 ± 1.93	65.46±0.86	67.15 ± 2.10	68.79 ± 2.42	61.51 ± 1.22	95.68±1.22
E.coli	93.76±0.56	93.82 ± 0.41	90.82 ± 1.49	82.38 ± 2.19	94.92 ± 0.32	90.81±0.63	97.22±0.28

Zhang M, Chen Y. Link prediction based on graph neural networks. NeurIPS 2018



Link prediction approaches

- local topological predictors
- Global predictors, learning
 - Model-based
 - Fit a model to data by maximizing likelihood which is defined in terms of edge probabilities \Rightarrow Pr (i \rightarrow j | θ)
 - E.g. stochastic block model
 - Optimization-based
 - Adding an edge increases a measure, e.g. Q modularity
 - Embedding-based
 - Proximity in the embedded space {put connected nodes close together}

Model based link prediction



Infer Pr (i \rightarrow j | θ) for each candidate pair based on maximizing likelihood of the observed graph or other optimizations

SBM variants, Q-modularity and Infomap, e.g. Number of edges between module node i and j belong to divided by maximum possible edges between those modules

Assume links are within modules

Abbreviation	Description			
Q	modularity, Newman-Girvan	:		
Q-MR	modularity, Newman's multiresolution			
Q-MP modularity, message passing		•		
B-NR (SBM)	Bayesian stochastic block model, Newman and Reinert	,		
B-NR (DC-SBM)	Bayesian degree-corrected stochastic block model, Newman and Reinert			
B-HKK (SBM)	Bayesian stochastic block model, Hayashi, Konishi and Kawamoto			
cICL-HKK (SBM)	Corrected integrated classification likelihood, stochastic block model			
Infomap	Map equation			
MDL (SBM)	Minimum description length, stochastic block model			
MDL (DC-SBM) Minimum description length, degree-corrected stochastic block		•		
S-NB Spectral with non-backtracking matrix				



Evaluating Overfit and Underfit in Models of Network Community Structure, TKDE 2020

Embedding based link prediction

$G \rightarrow$ embed in vector space \rightarrow distance in the embedded space

Node embedding methods derive a vector representation per each node in the graph so that connected nodes have similar vectors ⇒ close in the embedded space means more likely to be linked



Matrix factorization based e.g. svd ⇒ Deep learning methods e.g. deepwalk

What embedding to use? Graph Representation Learning

See<u>A Tutorial on Network Embeddings</u>, 2018



Which one is the best?

no one predictor or family is best, or worst, across all realistic inputs

550 structurally diverse networks from six scientific domains



Ghasemian A, Hosseinmardi H, Galstyan A, Airoldi EM, Clauset A. Stacking Models for Nearly Optimal Link Prediction in Complex Networks. PNAS 2020.

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Which one to choose?

Stack the models

learns to apply the best individual predictor according to the input's characteristics

algorithm	AUC	precision	recall
Q	0.7 ± 0.14	0.14 ± 0.17	0.67 ± 0.15
Q-MR	0.67 ± 0.15	0.12 ± 0.17	0.63 ± 0.13
Q-MP	0.64 ± 0.15	0.09 ± 0.11	0.59 ± 0.17
B-NR (SBM)	0.81 ± 0.13	0.13 ± 0.12	0.65 ± 0.22
B-NR (DC-SBM)	0.7 ± 0.2	0.12 ± 0.12	0.61 ± 0.24
cICL-HKK	0.79 ± 0.13	0.14 ± 0.14	0.58 ± 0.25
B-HKK	0.77 ± 0.13	0.11 ± 0.1	0.51 ± 0.26
Infomap	0.73 ± 0.14	0.12 ± 0.12	0.68 ± 0.13
MDL (SBM)	0.79 ± 0.15	0.14 ± 0.13	0.57 ± 0.3
MDL (DC-SBM)	0.84 ± 0.1	0.13 ± 0.11	0.78 ± 0.12
S-NB	0.71 ± 0.19	0.12 ± 0.13	0.66 ± 0.17
mean model-based	0.74 ± 0.16	0.12 ± 0.13	0.63 ± 0.21
mean indiv. topol.	0.6 ± 0.13	0.09 ± 0.16	0.53 ± 0.35
mean indiv. topol. & model	0.63 ± 0.15	0.09 ± 0.16	0.55 ± 0.33
emb-DW	0.63 ± 0.23	0.17 ± 0.19	0.42 ± 0.35
emb-vgae	0.69 ± 0.19	0.05 ± 0.05	0.69 ± 0.21
all topol.	0.86 ± 0.11	0.42 ± 0.33	0.44 ± 0.32
all model-based	0.83 ± 0.12	0.39 ± 0.34	0.3 ± 0.29
all embed.	0.77 ± 0.16	0.32 ± 0.32	0.32 ± 0.31
all topol. & model	0.87 ± 0.1	0.48 ± 0.36	0.35 ± 0.35
all topol. & embed.	0.84 ± 0.13	0.4 ± 0.34	0.39 ± 0.33
all model & embed.	0.84 ± 0.13	0.36 ± 0.32	0.36 ± 0.31
all topol., model & embed.	0.85 ± 0.14	0.42 ± 0.34	0.39 ± 0.33

550 structurally diverse networks from six scientific domains

Ghasemian A, Hosseinmardi H, Galstyan A, Airoldi EM, Clauset A. Stacking Models for Nearly Optimal Link Prediction in Complex Networks. arXiv preprint arXiv:1909.07578. 2019 Sep 17.

Which one to choose?

Stack the models

learns to apply the best individual predictor according to the input's characteristics

$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Algorithm	AUC	Precision	Recall
Q-MR 0.87 ± 0.07 0.38 ± 0.16 0.78 ± 0.07 Q-MP 0.86 ± 0.08 0.25 ± 0.07 0.83 ± 0.09 B-NR (SBM) 0.93 ± 0.06 0.3 ± 0.08 0.85 ± 0.12 B-NR (DC-SBM) 0.93 ± 0.07 0.28 ± 0.08 0.88 ± 0.08 clCl-HKK 0.93 ± 0.08 0.34 ± 0.1 0.85 ± 0.14 B-HKK 0.93 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.12 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all topol. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.11 0.86 ± 0.22 0.83 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. model & embed. <th< td=""><td>Q</td><td>0.89 ± 0.07</td><td>0.42 ± 0.13</td><td>0.85 ± 0.08</td></th<>	Q	0.89 ± 0.07	0.42 ± 0.13	0.85 ± 0.08
Q-MP 0.86 ± 0.08 0.25 ± 0.07 0.83 ± 0.09 B-NR (SBM) 0.93 ± 0.06 0.3 ± 0.08 0.85 ± 0.12 B-NR (DC-SBM) 0.93 ± 0.07 0.28 ± 0.08 0.88 ± 0.08 clCL-HKK 0.93 ± 0.08 0.34 ± 0.1 0.85 ± 0.14 B-HKK 0.88 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.12 0.84 ± 0.12 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.22 ± 0.25 0.62 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.45 ± 0.16 0.92 ± 0.13 emd-DW 0.95 ± 0.07 0.76 ± 0.22 0.68 ± 0.17 all topol. 0.95 ± 0.07 0.76 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.76 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.78 ± 0.21 0.74 ± 0.23 all topol. & model 0.96 ± 0.11 0.86 ± 0.22 0.83 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 </td <td>Q-MR</td> <td>0.87 ± 0.07</td> <td>0.38 ± 0.16</td> <td>0.78 ± 0.07</td>	Q-MR	0.87 ± 0.07	0.38 ± 0.16	0.78 ± 0.07
B-NR (SBM) 0.93 ± 0.06 0.3 ± 0.08 0.85 ± 0.12 B-NR (DC-SBM) 0.93 ± 0.07 0.28 ± 0.08 0.88 ± 0.08 clCL-HKK 0.93 ± 0.08 0.34 ± 0.1 0.85 ± 0.14 B-HKK 0.88 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.94 ± 0.07 0.3 ± 0.12 0.87 ± 0.16 MDL (DC-SBM) 0.94 ± 0.07 0.3 ± 0.12 0.84 ± 0.12 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.22 ± 0.25 0.62 ± 0.33 mean indiv. topol. 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emd-DW 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all topol. 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all topol. & model 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.83 ± 0.25 all topol. & model 0.96 ± 0.11 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.19 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0	Q-MP	0.86 ± 0.08	0.25 ± 0.07	0.83 ± 0.09
B-NR (DC-SBM) 0.93 ± 0.07 0.28 ± 0.08 0.88 ± 0.08 clCL-HKK 0.93 ± 0.08 0.34 ± 0.1 0.85 ± 0.14 B-HKK 0.88 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.12 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. 0.64 ± 0.19 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all topol. & model 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.17 0.86 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	B-NR (SBM)	0.93 ± 0.06	0.3 ± 0.08	0.85 ± 0.12
cICL-HKK 0.93 ± 0.08 0.34 ± 0.1 0.85 ± 0.14 B-HKK 0.88 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.12 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all topol. & model 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.96 ± 0.09 0.89 ± 0.21 0.88 ± 0.19 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.23 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	B-NR (DC-SBM)	0.93 ± 0.07	0.28 ± 0.08	0.88 ± 0.08
B-HKK 0.88 ± 0.07 0.17 ± 0.05 0.79 ± 0.17 Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.1 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.19 0.78 ± 0.21 0.74 ± 0.23 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	cICL-HKK	0.93 ± 0.08	0.34 ± 0.1	0.85 ± 0.14
Infomap 0.91 ± 0.04 0.29 ± 0.08 0.83 ± 0.05 MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.1 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.96 ± 0.19 0.86 ± 0.22 0.83 ± 0.25 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	B-HKK	0.88 ± 0.07	0.17 ± 0.05	0.79 ± 0.17
MDL (SBM) 0.94 ± 0.07 0.31 ± 0.09 0.87 ± 0.16 MDL (DC-SBM) 0.93 ± 0.09 0.26 ± 0.09 0.89 ± 0.11 S-NB 0.94 ± 0.07 0.3 ± 0.1 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22	Infomap	0.91 ± 0.04	0.29 ± 0.08	0.83 ± 0.05
$\begin{array}{llllllllllllllllllllllllllllllllllll$	MDL (SBM)	0.94 ± 0.07	0.31 ± 0.09	0.87 ± 0.16
S-NB 0.94 ± 0.07 0.3 ± 0.1 0.87 ± 0.08 mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.78 ± 0.22 0.83 ± 0.25 all topol. & model 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	MDL (DC-SBM)	0.93 ± 0.09	0.26 ± 0.09	0.89 ± 0.11
mean model-based 0.91 ± 0.08 0.3 ± 0.12 0.84 ± 0.12 mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.96 ± 0.11 0.78 ± 0.21 0.83 ± 0.25 all topol. & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol. & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	S-NB	0.94 ± 0.07	0.3 ± 0.1	0.87 ± 0.08
mean indiv. topol. 0.64 ± 0.19 0.2 ± 0.27 0.56 ± 0.33 mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.11 0.76 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.96 ± 0.11 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	mean model-based	0.91 ± 0.08	0.3 ± 0.12	0.84 ± 0.12
mean indiv. topol. & model 0.7 ± 0.21 0.22 ± 0.25 0.62 ± 0.32 emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	mean indiv. topol.	0.64 ± 0.19	0.2 ± 0.27	0.56 ± 0.33
emd-DW 0.95 ± 0.1 0.45 ± 0.16 0.92 ± 0.13 emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & model 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	mean indiv. topol. & model	0.7 ± 0.21	0.22 ± 0.25	0.62 ± 0.32
emb-vgae 0.95 ± 0.08 0.09 ± 0.02 0.96 ± 0.09 all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	emd-DW	0.95 ± 0.1	0.45 ± 0.16	0.92 ± 0.13
all topol. 0.97 ± 0.08 0.89 ± 0.21 0.88 ± 0.2 all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	emb-vgae	0.95 ± 0.08	0.09 ± 0.02	0.96 ± 0.09
all model-based 0.95 ± 0.07 0.76 ± 0.2 0.68 ± 0.17 all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all topol.	0.97 ± 0.08	0.89 ± 0.21	0.88 ± 0.2
all embed. 0.95 ± 0.11 0.75 ± 0.23 0.74 ± 0.23 all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all model-based	0.95 ± 0.07	0.76 ± 0.2	0.68 ± 0.17
all topol. & model 0.98 ± 0.06 0.89 ± 0.22 0.88 ± 0.19 all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all embed.	0.95 ± 0.11	0.75 ± 0.23	0.74 ± 0.23
all topol. & embed. 0.96 ± 0.1 0.86 ± 0.22 0.83 ± 0.25 all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all topol. & model	0.98 ± 0.06	0.89 ± 0.22	0.88 ± 0.19
all model & embed. 0.96 ± 0.09 0.78 ± 0.21 0.74 ± 0.22 all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all topol. & embed.	0.96 ± 0.1	0.86 ± 0.22	0.83 ± 0.25
all topol., model & embed. 0.97 ± 0.09 0.86 ± 0.23 0.84 ± 0.23	all model & embed.	0.96 ± 0.09	0.78 ± 0.21	0.74 ± 0.22
	all topol., model & embed.	0.97 ± 0.09	0.86 ± 0.23	0.84 ± 0.23

Near perfect in social networks

Ghasemian A, Hosseinmardi H, Galstyan A, Airoldi EM, Clauset A. Stacking Models for Nearly Optimal Link Prediction in Complex Networks. arXiv preprint arXiv:1909.07578. 2019 Sep 17.

Link Prediction in Attributed Graphs

Individual characteristics or activity (attributes) & relations (graph) Annotated networks, **metadata** on nodes, **side** information



characteristics

age, occupation, salary, sex, etc.



activity & interest

check-ins, page-likes, group memberships, movies

Attributed Graphs

Interplay between attributes and relations, a positive feedback loop derived by two social theories:

- Social Selection
 - Similarity of individuals' characteristics motivates them to form relations

⇒ Similarity of node's attributes is a link predictors in addition to structure proximity

- Social Influence
 - Characteristics of individuals may be affected by the characteristics of their relations
 - ⇒ Your neighbours' attributes can reveal yours



Link Prediction in Attributed Graphs

Graph Neural Networks get attributed graphs as the input and can be used for many tasks including link prediction



numerous methods, multiple surveys, <u>e.g. this one (2019)</u>, two of the notable works: <u>GCN</u> (2016), <u>GAT</u> (2018), an excellent <u>course last term</u> on this & a <u>new book</u>!

Approach	Category	Inputs	Pooling	Readout	Time Complexity
GNN* (2009) [15]	RecGNN	A,X,X^e		a dummy super node	
GraphESN (2010) [16]	RecGNN	A, X		mean	
GGNN (2015) [17]	RecGNN	A, X	142 C	attention sum	-
SSE (2018) [18]	RecGNN	A, X	-	-	-
Spectral CNN (2014) [19]	Spectral-based ConvGNN	A, X	spectral clustering+max pooling	max	$O(n^3)$
Henaff et al. (2015) [20]	Spectral-based ConvGNN	A, X	spectral clustering+max pooling		$O(n^3)$
ChebNet (2016) [21]	Spectral-based ConvGNN	A, X	efficient pooling	sum	O(m)
GCN (2017) [22]	Spectral-based ConvGNN	A, X	-	-	O(m)
CayleyNet (2017) [23]	Spectral-based ConvGNN	A, X	mean/graclus pooling		O(m)
AGCN (2018) [40]	Spectral-based ConvGNN	A, X	max pooling	sum	$O(n^2)$
DualGCN (2018) [41]	Spectral-based ConvGNN	A, X		a.	O(m)
NN4G (2009) [24]	Spatial-based ConvGNN	A, X		sum/mean	O(m)
DCNN (2016) [25]	Spatial-based ConvGNN	A, X	~	mean	$O(n^2)$
PATCHY-SAN (2016) [26]	Spatial-based ConvGNN	A, X, X^e	×	concat	
MPNN (2017) [27]	Spatial-based ConvGNN	A, X, X^e		attention sum/ set2set	O(m)
GraphSage (2017) [42]	Spatial-based ConvGNN	A, X	×	2	÷
GAT (2017) [43]	Spatial-based ConvGNN	A, X		-	O(m)
MoNet (2017) [44]	Spatial-based ConvGNN	A, X		-	O(m)
PGC-DGCNN (2018) [46]	Spatial-based ConvGNN	A, X	sort pooling	attention sum	$O(n^3)$
CGMM (2018) [47]	Spatial-based ConvGNN	A, X		concat	-
LGCN (2018) [45]	Spatial-based ConvGNN	A, X		-	-
GAAN (2018) [48]	Spatial-based ConvGNN	A, X	21.	-	O(m)
FastGCN (2018) [49]	Spatial-based ConvGNN	A, X	×	-	÷
StoGCN (2018) [50]	Spatial-based ConvGNN	A, X		e	
Huang et al. (2018) [51]	Spatial-based ConvGNN	A, X		-	-
DGCNN (2018) [52]	Spatial-based ConvGNN	A, X	sort pooling	-	O(m)
DiffPool (2018) [54]	Spatial-based ConvGNN	A, X	differential pooling	mean	$O(n^{2})$
GeniePath (2019) [55]	Spatial-based ConvGNN	A, X	~	-	O(m)
DGI (2019) [56]	Spatial-based ConvGNN	A, X	<i>v</i>	147 1	O(m)
GIN (2019) [57]	Spatial-based ConvGNN	A, X	-	concat+sum	O(m)
ClusterGCN (2019) [58]	Spatial-based ConvGNN	A, X		-	

More on this later

Common prediction tasks

- Link Prediction
- Node Classification
- Graph Classification

What is unsupervised node classification?

Examples:

https://paperswithcode.com/task/link-prediction

https://paperswithcode.com/task/node-classification

https://paperswithcode.com/task/graph-classification

Attributed Graphs

Interplay between attributes and relations, a positive feedback loop derived by two social theories:

- Social Selection
 - Similarity of individuals' characteristics motivates them to form relations
 - ⇒ Similarity of node's attributes is a link predictors in addition to structure proximity
- Social Influence
 - Characteristics of individuals may be affected by the characteristics of their relations
 - ⇒ Your neighbours' attributes can reveal yours



Similar nodes tend to link to each other

How to measure age homophily in a given friendship graph when you know age of every node?

birds of the same feather flock together





Similar nodes tend to link to each other

How to measure age homophily in a given friendship graph when you know age of every node? Similar to degree assortativity, measure the correlation of age across all edges

How to measure occupation homophily? categorical attribute instead of a numeric one

birds of the same feather flock together





Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

What indicates homophily in this matrix? How does homophily look like?



Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

What indicates homophily in this matrix? How does homophily look like? **dominant diagonal** **e**_{ij}: ratio of edges between each pair of values

Assortativity index

$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.} e_{.i}}{1 - \sum_{i} e_{i.} e_{.i}} = \frac{Tr[e] - ||e^{2}|}{1 - ||e^{2}||}$$

Is normalized Q-modularity assuming attributes partition the graph

$$Q = \sum_{i} e_{ii} - e_{i.}^{2} = \text{Tr}[e] - ||e^{2}||$$



Look at the mixing patterns

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What indicates homophily in this matrix? How does homophily look like? **dominant diagonal** **e**_{ij}: ratio of edges between each pair of values

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Is there other mixing patterns?



Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

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Is there other mixing patterns?









e.g. opposites attract



Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

Other mixing patterns, such as **heterophily** can also be reflected in the mixing matrix

How does heterophily look like in the mixing matrix? How does correlation look like in the mixing matrix?

e_{ij}: ratio of edges between each pair of values

$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.}e_{.i}}{1 - \sum_{i} e_{i.}e_{.i}} = \frac{Tr[e] - ||e^{2}||}{1 - ||e^{2}||}$$



Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

Other mixing patterns, such as heterophily can also be reflected in the mixing matrix

How does heterophily look like in the mixing matrix? How does **correlation** look like in the mixing matrix? A **dominant cell** in each row and column

e_{ij}: ratio of edges between each pair of values

$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.} e_{.i}}{1 - \sum_{i} e_{i.} e_{.i}} = \frac{Tr[e] - ||e^{2}||}{1 - ||e^{2}||}$$





Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

Other mixing patterns, such as heterophily can also be reflected in the mixing matrix

How to quantify the overall correlation?

Does it resemble anything?

e_{ij}: ratio of edges between each pair of values

$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.}e_{.i}}{1 - \sum_{i} e_{i.}e_{.i}} = \frac{Tr[e] - ||e^{2}||}{1 - ||e^{2}||}$$



Look at the mixing patterns

Mixing matrix shows the number of edges connecting each pair of attribute values

Other mixing patterns, such as heterophily can also be reflected in the mixing matrix

How to quantify the overall correlation?

Does it resemble anything? confusion matrix where pairwise cluster overlaps are changed to edges between pair of values

e_{ij}: ratio of edges between each pair of values

$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.} e_{.i}}{1 - \sum_{i} e_{i.} e_{.i}} = \frac{Tr[e] - ||e^{2}||}{1 - ||e^{2}||}$$



Look at the mixing patterns

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Other mixing patterns, such as heterophily can also be reflected in the mixing matrix **e**_{ij}: ratio of edges between each pair of values

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$$r = \frac{\sum_{i} e_{ii} - \sum_{i} e_{i.}e_{.i}}{1 - \sum_{i} e_{i.}e_{.i}} = \frac{Tr[e] - ||e^{2}||}{1 - ||e^{2}||}$$



How to quantify the overall correlation?

measure the total dispersion similar to clustering agreement indexes

Beyond Assortativity: Proclivity Index for Attributed Networks, PAKDD (2017)



Structural Correlation of Attributes Proclivity

"inclination or predisposition	similar homophily	different heterophily	random
toward a particula thing"			
Assortativity	1.0	-0.33	-0.33
Prone	1.0	1.0	0.11

Beyond Assortativity: Proclivity Index for Attributed Networks, PAKDD (2017)



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Structural Correlation of Attributes Proclivity Shape and Color

Cross-proclivity in addition to self-proclivity



no correlation







Correlation between your income and occupations of your friends

کی کی ا

Structural Correlation of Attributes



Even if you don't put your information online, that information can be inferred/predicted based on what your friends reveal about themselves



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Predicting missing node attributes

Most graphs are incomplete, and often attributes of some nodes are missing

We can use structural correlations to predict a missing attribute to be e.g. the average (scalar) or most common (categorical) value of its neighbors' non-missing attributes

What does this local smoothing assume about the mixing patterns?





Predicting missing node attributes

Most graphs are incomplete, and often attributes of some nodes are missing

We can use structural correlations to predict a missing attribute to be e.g. the average (scalar) or most common (categorical) value of its neighbors' non-missing attributes

What does this local smoothing assume about the mixing patterns? mean (scalar) & mode (categorical) \Rightarrow assortative







Predicting missing node attributes, example

missing = mean (scalar) & mode (categorical)

what is the prediction in these two cases for node i?





Predicting missing node attributes, example

missing = mean (scalar) & mode (categorical)

what is the prediction in these two cases for node i?



what is the predictions for node x_h ?



Predicting missing node attributes, example

missing = mean (scalar) & mode (categorical)

what is the prediction in these two cases for node i?



what is the predictions for node x_b^2 ? repeat given current predictions

Label Propagation Algorithm

Was proposed for semi-supervised classification of iid data by defining a fully connected distance graph

Zhu X, Ghahramani Z. Learning from Labeled and Unlabeled Data with Label Propagation.





Label Smoothing

° (

Node classification

- Unsupervised learning
 - clustering, only graph is given, classes/clusters are not predefined
- Supervised learning
 - classifying, input is graph and labels on all nodes
 - You mask some nodes (labels and their connections) for training [inductive]
 - You mask some nodes (only labels) for training [transductive]
- Semi-supervised learning
 - input is graph and labels on some nodes
 - You mask some node labels for training (seeing the whole graph: transductive)
- Active learning
 - Input is graph and a budget that determines how many nodes you can query for labels
 - \circ \quad labels come in sequence and can be queried based on the current set





Semi-Supervised Node classification

- Traditional
 - label propagation & belief propagation
- Recent end-to-end methods (Feature Smoothing)
 - GCN and variants, which use a classification loss
- Embedding based
 - Unsupervised embedding extraction (e.g. node2vec) then apply a classifier



More on this later

Unifying Graph Convolutional Neural Networks and Label Propagation, 2020

Scalar values \Rightarrow correlation of predicted & actual values (r² correlation)

Categorical values \Rightarrow confusion matrix & average accuray

 C_{ii} = the number of nodes with predicted label i and actual label j

Accuracy = 1/34 Tr(C)



What is the accuracy?





Scalar values \Rightarrow correlation of predicted & actual values (<u>r² correlation</u>)

Categorical values \Rightarrow confusion matrix & average accuray

 C_{ii} = the number of nodes with predicted label i and actual label j

Accuracy = Tr(C)/Sum(C)



What is the accuracy? 90% but is always guessing the majority class and never getting the minority class correct

Class imbalance problem





Example

Truth: left circle all blue, right circle all red

Observed: 6 missing values





observed



What is the accuracy?





Example

Truth: left circle all blue, right circle all red

Observed: 6 missing values





observed



What is the accuracy?

Accuracy = % = 0.83





Active Search of Connections

What if you can query with some cost or given a budget?

Semi-supervised \Rightarrow Active learning

labels are local and depend on the given seed

(a) Local Clustering

(b) Active Search on Graph





Active Search of Connections for Case Building and Combating Human Trafficking, 2018

(c) Active Exploration

(d) Active Search of Connections