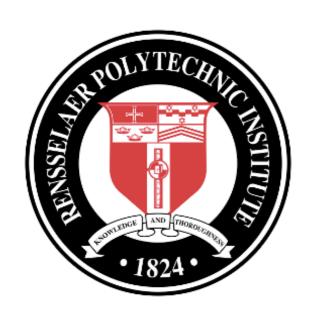
Inferring Degrees from Incomplete Networks and Nonlinear Dynamics

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Problem

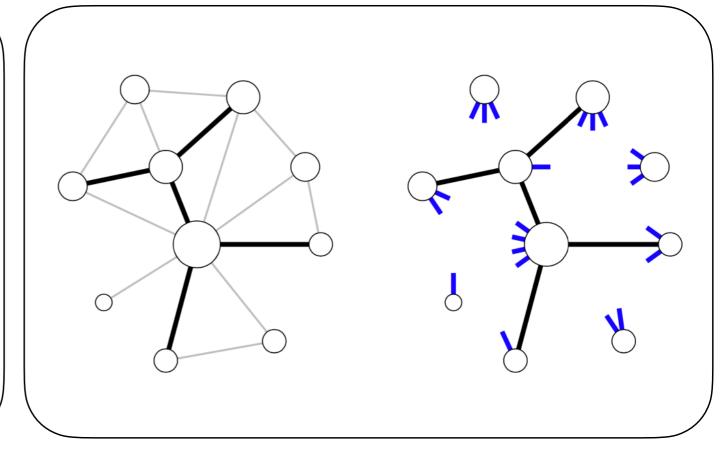
Given:

- Incomplete network
- Noisy equilibrium states
- Dynamical system

Goals:

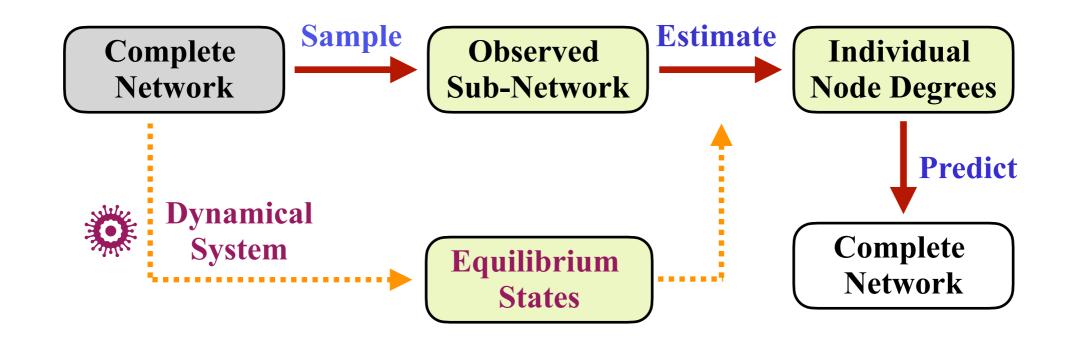
- Infer missing degrees
- Link prediction
- Robustness (sampling, measurement error, model misspecification)

- Ecology Networks: abundances of plants
- Regulatory Networks: expression levels of genes
- Epidemic Networks: infection rates of person



Prior Works

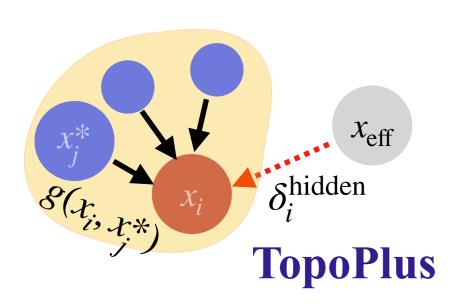
- Degree distribution, average degree, network size (BF1993; STS2008; KLS2011; KBM2012; GR2008; DKS2014; F1980; SW2005; AKM2009; RT2012; ZKS2015)
- Individual node degrees (GK2017)



Main idea: connecting topology to equilibria

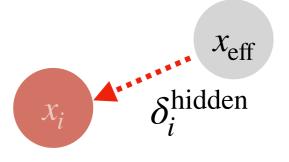
GBB2016 (Nature) JGM2020 (AAAI)





Equilibrium Condition

$$f(x_i) + \sum_{j \in V} A_{ij}^{(s)} g(x_i, x_j^*) + \delta_i^{\text{hidden}} g(x_i, x_{\text{eff}}) = 0$$



ZeroTopo



Mean-field Equilibrium Condition

$$f(x_i) + \delta_i^{\text{hidden}} g(x_i, x_{\text{eff}}) = 0$$

Experimental Setting

Dynamics	Network	n	m	$\langle oldsymbol{\delta} angle$	
(a) Ecological (b) Regulatory	Plant Pollinator Genes Network	97 662	972 1062	20.04 3.21	
© Epidemic	Facebook	4039	88234	43.69	
(a) $\dot{x}_i = B + x_i(1 - \frac{x_i}{2})(\frac{x_i}{2} - 1) + \sum_i \frac{x_i x_j A_{ij}}{2}$					

(a)
$$\dot{x}_i = B + x_i (1 - \frac{x_i}{K})(\frac{x_i}{C} - 1) + \sum_j \frac{x_i x_j A_{ij}}{D + E x_i + H x_j}$$

(b) $\dot{x}_i = -B x_i^f + \sum_j A_{ij} R \frac{x_j^h}{x_j^h + 1}$
(c) $\dot{x}_i = -B x_i + \sum_j A_{ij} R (1 - x_i) x_j$

$$\hat{\mathbf{b}} \, \dot{x}_i = -B x_i^f + \sum_j A_{ij} R \frac{x_j^h}{x_i^h + 1}$$

$$\hat{\mathbf{C}} \dot{x}_i = -Bx_i + \sum_j A_{ij} R(\hat{1} - x_i) x_j$$

Inferring Individual Node Degrees

	Method (sampling fraction)			
Network	ZeroTopo	_	TopoPlus+Round	
	(0%)	(10%)	(10%)	
Plant Pollinator	57.7	62.0	62.7	
Genes	86.2	87.1	87.8	
Facebook	59.5	64.1	65.4	

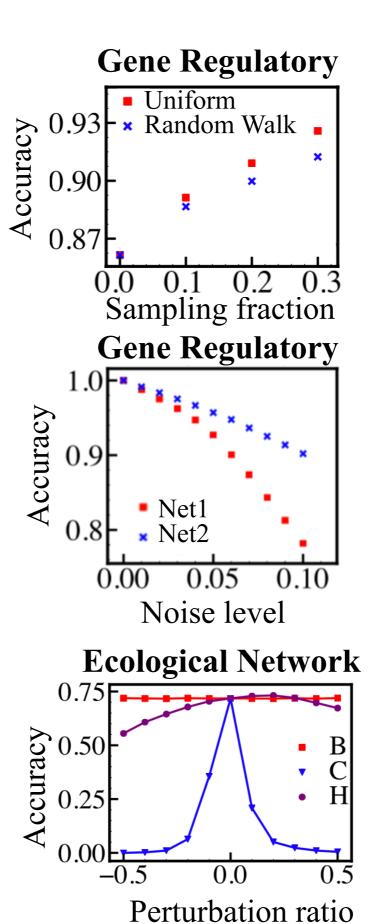
Accuracy: fraction of nodes with error $|\log(\hat{\delta}_i/\delta_i)| < 5\%$

Robustness

Sampling Method

• Measurement Error

Model Error



Comparison: degree estimation

On average 157% improvement over [GK2017]

Network Method	Plant Pollinator	Genes	Facebook
[GK2017]	5.10	19.96	9.00
ZeroTopo (w/ noise)	12.63 (10%)	84.27(10%)	9.19 (9%)

Degree estimation for link prediction

- •Our estimates: 41% improvement over observed degrees
- True degrees: 44% improvement over observed degrees

	AUC			
Method	Plant Pollinator	Genes	Facebook	
[BA1999] + Observed degrees	59.90	58.00	69.44	
[BA1999] + Our estimates	87.11	91.28	83.78	
[BA1999] + True degrees	87.42	92.45	83.83	

- All methods sample 1% of edges
- [BA1999] uses $s_{u,v} = \delta_u \delta_v$ for link prediction
- Our degree estimates with **TopoPlus** assumes no measurement error

Conclusions

- Our degree estimates use equilibrium states
- Accurate estimates even without topology, unlike existing methods
- Robust to sampling method, measurement errors, and model errors
- Our degree estimates significantly enhance link prediction

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