### Covariance Matrix Adaptation Evolution Strategy for Link Prediction in Dynamic Social Networks

Catherine A. Bliss, Morgan R. Frank, Christopher M. Danforth, & Peter Sheridan Dodds



Background Data Reciprocal reply networ Link prediction Similarity indices Evolutionary computatio Results

Data Recipi



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Background Data Reciprocal reply networks Link prediction Similarity indices Evolutionary computation Results Conclusions

Background Data Reciprocal reply networks



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Background Data Reciprocal reply network Link prediction Similarly indices Evolutionary computatio Results Conclusions



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Results Conclusions



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Data



40,000 tweets (100MB) / min. 50 million tweets (150GB) / day 50 billion tweets (100TB) / 5+ years

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Similarity indices Evolutionary com

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reply networks exhibit assortativity with respect to happiness, Journal of Computational Science =





- Liben-Nowell & Kleinberg (2007) author collaboration networks (N ∝ 10<sup>3</sup>)
- Use similarity indices to rank the most likely occurring top N links



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Similarity indices

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



Data Recipi

Similarity indices



Resource Allocation (Zhou, Lu, & Zhang, 2009)

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### Similarity indices

- Liben-Nowell & Kleinberg (2007) author collaboration networks (N ∝ 10<sup>3</sup>)
- Use similarity indices to rank the most likely occurring top N links C(u, v) = |Γ(u) ∩ Γ(v)|

$$R(u, v) = \sum_{z \in \Gamma(u) \cap \Gamma(v)} \frac{1}{|\Gamma(z)|}$$
$$W(u, v) = 1 - \frac{1}{2} \sum |f_{u,n} - f_{v,n}|$$



Background Data Reciprocal reply netwo Link prediction Similarity indices Evolutionary computati Results Conclusions

Common neighbors	$C(u,v) =  \Gamma(u) \cap \Gamma(v) $	Background
Jacard	$J(u,v) = \frac{ \Gamma(u) \cap \Gamma(v) }{ \Gamma(u) \cup \Gamma(v) }$	Data Beciprocal reply networks
Adamic-Adar	$A(u,v) = \sum_{z \in (v,v) \in \Gamma(v)} \frac{1}{\log( \Gamma(z) )}$	Link prodiction
Pref Attachment	$Pr(u,v) = k_{u} \times k_{v}$	Similarity indices
Hub promoted	$Hp(x,y) = \frac{ \Gamma(u) \cap \Gamma(v) }{\min\{k_{U}, k_{V}\}}$	Desults
Hub depressed	$Hd(u,v) = \frac{ \Gamma(u) \cap \Gamma(v) }{\max\{k_u, k_v\}}$	Conclusions
LHN	$L(u,v) = \frac{ \Gamma(u) \cap \Gamma(v) }{k_U k_V}$	Contractorio
Salton	$Sa(u,v) = \frac{ \Gamma(u) \cap \Gamma(v) }{\sqrt{k_U k_V}}$	
Sorenson	$So(u,v) = \frac{2 \Gamma(u)\cap\Gamma(v) }{k_U+k_V}$	
Resource Allocation	$R(u,v) = \sum_{z \in \Gamma(u) \cap \Gamma(v)} \frac{1}{ \Gamma(z) }$	
Average Path Weight	$P(u,v) = \frac{\sum\limits_{\substack{p \in \mathcal{P}_2(u,v) \cup \mathcal{P}_3(u,v)}} w_p}{ \mathcal{P}_2(u,v)  +  \mathcal{P}_3(u,v) }$	
Katz	$\mathcal{K} = \sum_{n=1}^{\infty} \beta^n A^n$	
Tweet count similarity	$T(u,v) = 1 - \frac{ T(u) - T(v) }{\max\{ T(a) - T(b) \}}_{a,b \in V}$	
Word similarity	$W(u,v) = 1 - \frac{1}{2} \sum_{n=1}^{50000}  f_{u,n} - f_{v,n} $	
Happiness similarity	$H(u, v) = 1 - \frac{ h(u) - h(v) }{\max\{ h(a) - h(b) \}_{a, b \in V}}$	
ld similarity	$I(u, v) = 1 - \frac{ Id(u) - Id(v) }{\max\{ Id(a) - Id(b) \}_{a, b \in V}}$	
	(日)	











### CMA-ES implementation<sup>2</sup>

#### Individual An individual or candidate solution is a vector, $\vec{w} \in \mathbf{R}^{16}$ . A C P Hd Hp Sa So ĸ н R w Pr L 4 .9 .5 .8 .3 2 .01 .6 .2 .04 .8 .1 -.1

### CMA-ES implementation<sup>2</sup>



<sup>&</sup>lt;sup>2</sup>Hansen, N. (2006). The CMA Evolution Strategy: A Comparing Review. In J.A. Lozano, P. Larrañga, I. Inza and E. Bengoetxea (eds.). Towards a new evolutionary computation. Advances in estimation of distribution algorithms. pp. 75-102, Springer.

## CMA-ES implementation<sup>2</sup>

Individual	⇒	Reproduction & Mutation		Population
An individual or candidate solution is a vector, $\vec{w} \in \mathbf{R}^{16}$ .		<b>CMA-ES</b> From 1 individual, generate a Gaussian cloud of candidate colutions in <b>P</b> <sup>16</sup> using		A population consists of several candidate solutions (vectors in $R^{16}$ ).
		the covariance matrix.		
Selection	-	Evaluate fitness		
The candidate solution with the best performance (min. fitness) suprives		$S = \sum_{i=1}^{16} w_i S_i .$	Noo sco nev	de-node pairs w/top res in S are predicted . v links.
selection.			Fitn	ess=# incorrect links # predicted links

<sup>2</sup>Hansen, N. (2006). The CMA Evolution Strategy: A Comparing Review. In J.A. Lozano, P. Larrañga, I. Inza and E. Bengoetxea (eds.). Towards a new evolutionary computation. Advances in estimation of distribution algorithms. pp. 75-102, Springer.







## **Receiver Operating Curve**







### Precision



Precision depicts  $\frac{TP}{TP+FP}$ . High precision is achieved for top N < 20, which is often the region of interest. The precision for predicted links in  $W4 \rightarrow W5'$  is lower than the other weeks and this may be due to missing data for those weeks



### Conclusions

- Evolutionary algorithms show promise
- Many additional questions in link prediction (e.g., prediction of weights, prediction of link decay)
- Leveraging link prediction to understand network dynamics
- Further investigation of the role of incomplete data on network inference

### Thank you

- Manuscript: In press at the Journal of Computational Science. Pre-print available at http://arxiv.org/abs/1304.6257
- Contact: www.cems.uvm.edu/~cabliss
- Lab: www.onehappybird.com



Conclusions

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