Evolution of distances in preferential attachment models

Júlia Komjáthy & Joost Jorritsma

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Scale-free property

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$$\mathbb{P}(\deg(v) = x) \asymp \frac{C}{x^{\tau}}$$

$$\log \mathbb{P}(\deg(v) = x) \asymp \log C - \tau \log x$$

log(proportion of degree *x* vertices) vs log *x* is a straight line.

Power laws



Figure 5: The outdegree plots: Log-log plot of frequency f_d versus the outdegree d.



Log-log plot of degree distribution of the router level internet network

from Faloutsos, Faloutsos, Faloutsos. 1999

Growing networks

A question:

How did the network evolve around the servers of '99?



Kevin Bacon game: Movie networks are small worlds

from Mark Robinson Writes

Definition

A network G on N vertices is a *small world* if the average distance

$$\overline{\mathsf{dist}}(G) = \frac{1}{\binom{N}{2}} \sum_{u,v \in G} d_G(u,v) = \Theta(\log N).$$

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Alternative def of ultrasmall:

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Static network models

- Configuration model
- Inhomogeneous random graphs
- Chung-Lu, Norros-Reitu models

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- Original Barabási-Albert model aka Preferential attachment model
- Variable degree Preferential attachment model
- Bianconi-Barabási model aka PA with multiplicative fitness
- PA with additive fitness
- PA with power of choice

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- Heuristically:

 $\operatorname{Prob}(v_{N+1} \longrightarrow v_i) \propto \deg_N(v_i)$

Classic variations

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- Fixed outdegree: (m, δ) : *m* edges/new vertex. **Prob** $(v_{N+1} \xrightarrow{j} v_i) \propto \deg_{N,i}(v_i) + \delta/m$
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- Variable outdegree: $\forall v_i \in \mathsf{PA}_N$: independently $\operatorname{Prob}(v_{N+1} \longrightarrow v_i) \propto \frac{f(\deg_N(v_i))}{N}$, where f(x) concave with $\lim_{x \to \infty} f(x)/x = \gamma_f \leq 1$.

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Animation

Degree distribution

Theorem (Degree sequence of PA(m,0))

Bollobás, Riordan, Spencer, Tusnády '01 Limiting degree distribution of Classic PA (m,0)

$$\lim_{N\to\infty} \operatorname{Prob}(\operatorname{deg}(V_N)=k) \asymp \frac{1}{k^3}.$$

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Generally

Limiting degree distribution is

$$\lim_{N\to\infty} \mathbf{Prob}(\deg(V_N)=k) \asymp \frac{1}{k^{\tau}},$$

where

$$\tau_{m,\delta} = 3 + \delta/m \qquad \tau_f = 1 + 1/\gamma_f.$$

Graph distances

Theorem (Distances in PAMs, $\tau \in (2,3)$ **)**

Dommers, v/d Hofstad, Hoogiemstra '10 & Dereich and Mörters '13 Let U_N, V_N be two uniformly chosen vertices (within the giant component)¹. When $\tau \in (2,3)$,

$$d_G^{(N)}(U_N, V_N) = (1 + o_{\mathbf{P}}(1)) \log \log N \cdot \frac{4}{|\log(\tau - 2)|}$$

Lower tightness holds.

 $^{{}^{1}}d_{G}^{(N)}$ graph distance within PA_{N}

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 $\tau > 3$ Dommers, v/d Hofstad, Hoogiemstra '10 Typical distance in PA(m, δ) for $m \ge 2, \tau > 3$:

$$\mathsf{d}_{\mathsf{G}}^{(\mathsf{N})}(U_N,V_N) = \Theta(\log N).$$

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Universality

For static models with power law degrees, $\tau \in (2,3)$

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Factor 2:

In PA, a high degree vertex tends to be connected via a path of 2 to a higher degree vertex.

In static network models, directly.

Back to the question:

How do distances shrink as time passes?

Theorem (Evolution of distances)

Jorritsma, Komjáthy, (AoAP, 22) Let U_N, V_N be two uniformly chosen vertices in PA_N (within the giant component). When $\tau \in (2,3)$, & $m \ge 2$, for t > N:

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$$d_G^{(t)}(U_N, V_N) \asymp \left(\log \log N - \log(\log(t/N) \lor 1)\right) \cdot \frac{4}{\log(\tau - 2)|} \lor 2$$

+tightness around the main term.

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$$X_N \coloneqq \sup_{t>N} \left(d_G^{(t)}(U_N, V_N) - \left[\left(\log \log N - \log(\log(t/N) \vee 1) \right) \cdot \frac{4}{\log(\tau - 2)|} \vee 2 \right] \right)$$

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 $(X_N)_{N \ge 1}$ is a tight sequence of random variables. Message 1: Fix $a \in (0, 1)$. Then at $t = N \exp(\varepsilon (\log N)^a) \ll N^{1+\varepsilon}$,

$$d_G^{(t)}(U_N, V_N) \approx (1-a) \log \log N \cdot \frac{4}{|\log(\tau-2)|}.$$

Message 2: $d_G^{(t)}(U_N, V_N)$ never leaves a (tight) strip around the main term.

Proofs

Lower bound

• A path counting method:

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$$\mathbb{E}\Big[\sum_{t>N} \sum_{[\pi_{U_N,V_N} \text{too short at } t]} \mathbf{1}\{\pi_{U_N,V_N} \text{ present in } \mathsf{PA}_t\}\Big] \xrightarrow{?} 0.$$

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$$\mathbb{E}\Big[\sum_{\pi_{U_N,V_N} \text{too short at arrival}} \mathbf{1}\{\pi_{U_N,V_N} \text{ present in } \mathsf{PA}_t\}\Big] \xrightarrow{?} 0.$$

• A path counting method:

A **possible path** π is a sequence of labelled vertices. The path's (potential) arrival time is the youngest vertex on the path. Among all possible labeled paths π_{U_N,V_N} ,

$$\mathbb{E}\Big[\sum_{t>N} \sum_{[\pi_{U_N,V_N} \text{too short at } t]} \mathbf{1}\{\pi_{U_N,V_N} \text{ present in } \mathsf{PA}_t\}\Big] \xrightarrow{?} 0.$$
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- A union bound not over time, but over possible paths.
- This still does not tend to zero...





• A path is bad if it reaches a too old vertex in too few steps



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- i.e., if its k vertex is older than some threshold ℓ_k



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- \Rightarrow Enough to bound:

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i.e., containing the newly added v_t .



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 This tends to zero ⇒ whp no bad paths present ever.



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Upper bound
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- They are (killed) multitype branching processes.
- We will use this info as a black box.

Proof outline for fixed *N*

y-axes=arrival time of vertices; *x*-axis: graph distance. LWL=limit distribution of the trees, depth R_N (large constant).



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Building the connecting path

y-axes=arrival time of vertices; *x*-axis: graph distance.



Greedy path to the core

Core: vertices born before time \sqrt{N} . Core has bounded diameter.



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$$K_N \le \frac{\log \log(N)}{|\log(\tau - 2)|} + C$$

- From any vertex in Layer k, there are at least $s_k^{arepsilon_k}$ young wedges to Layer k+1

- "Young" edges ensures independent weights w.r.t. to weighted LWL



Dense core has small diameter



Extension to the growing network

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- We need to track the degree-growth of the vertex that starts the wedging procedure (via a Móri martingale).
- We redefine the layers for each time t > N
- Careful union bound: summing error probs only where the event changes



Thank you for the attention!



Figure: Six instances of an infection spreading on a two-dimensional spatial scale free networks with different parameters.

Spreading processes on networks

Susceptible-Infected model:

- At time *t* = 0 the source node is infected, all other nodes susceptible.
- if, on an edge {u, v}, u is infected and v is not, then v becomes infected after a random iid transmission delay L_(u,v).

The epidemic curve

The function that counts the total number of infected nodes before time *t*:

 $I(t) = #\{ infected nodes before time t \}$



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$$d_L(u, v) = \min_{\pi: \text{path } u \leftrightarrow v} \text{sum of } L_e \text{ on edges on } \pi$$

Pre-sampling all randomness

Add *iid* weights from distribution *L* to existing edges.

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Epidemic curve:

$$I(t) = \#\{v : d_L(u, v) \le t\}.$$



$$L \equiv 1: \text{ graph distance}$$

$$d_G^{(N)}(U_N, V_N) \asymp \log \log N \cdot \frac{4}{|\log(\tau-2)|}$$

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Theorem (Explosion in PA)

1

Jorritsma, K, '20 Consider PA with $\tau \in (2,3)$, and U_N, V_N two typical vertices in the giant component of **PA**_N. Then if

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From small to mini-worlds

E(*L*) = ∞: fluctuations are tight in most cases. Lower tightness always, upper tightness under technical condition that holds for most *L*s.

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• Fix your favorite $1 \ll g(N) = O(\log \log N)$. Then one can construct a distribution *L* such that

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Shrinking of weighted distances when $I(L) = \infty$

For graph distances

$$\sup_{t' \ge t} \left(d_G^{(N')}(U_N, V_N) - 2 \left(\log \log N - \log(1 \vee \log(N'/N)) \right) \cdot \frac{2}{\log(\tau - 2)|} \vee 1 \right) = O_{\mathsf{P}}(1).$$

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E.

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