

# An Introduction to “Growing Protean Graphs”

Andy Wilson

February 7, 2012

This paper consists of a short introduction to the recent paper “Growing Protean Graphs” by Pawel Prałat and Nicholas Wormald ([4]). In this paper, Prałat and Wormald introduce a random graph model that gains vertices over time<sup>1</sup> and emulates the web graph, the real-world graph in which vertices are websites and edges are hyperlinks. The key features that the authors wish to emulate are the *power law* on vertex degrees and the unique *giant component*. The first property means that the number of vertices of degree  $d$  is approximately  $d^{-\gamma}$  for some positive constant  $\gamma$ , and the second means that over half of the vertices of the graph are in the same connected component. Many classical graphs, such as  $\mathcal{G}_{n,p}$  do not have these properties, but there are other recent models that do attain these properties, such as preferential attachment, which is analyzed by Chung and Lu in [2]. Protean graphs are unique in that “older” vertices are preferred when edges are added. We will not prove that protean graphs have these properties in this paper, but [4] and [3] contain proofs of these results.

Instead, in this paper we will present the algorithm on how to grow a protean graph and then use the differential equations method to analyze two key properties of the vertices of a protean graph. A *protean graph* is a sequence of undirected graphs  $G_t$  created by the following process:

- Choose  $p \in (0.5, 1]$ ,  $d \in \mathbb{N}$ ,  $\eta \in (0, 1)$ .
- Let  $G_0$  be the graph with one vertex,  $v_1$ .
- Given  $G_t$ , randomly add a vertex (with probability  $p$ ) or remove a vertex (with probability  $1 - p$ )
  - If adding a vertex, label the vertex  $v_i$ , where  $i$  is the smallest integer not yet used to label a vertex. Then randomly add  $d$  edges from  $v_i$  to existing vertices, where  $v_j$  gets an edge with probability  $j^{-\eta} / (\sum_{v_k} k^{-\eta})$ . (Technically, these edges are added randomly one at a time.)
  - If removing a vertex, choose an existing vertex uniformly at random. Assume we choose  $v_l$ . Delete  $v_l$  and all its edges. Then decrement the label on each vertex with label larger than  $l$  by one.

---

<sup>1</sup>In [3], the first author introduced a similar model that had a fixed number of vertices.

We will alter the above process slightly in order to do our analysis. Instead of starting with the graph on a single vertex, we will now begin with  $G_{t_0}$ , a graph on  $n$  vertices. Let  $v$  be the vertex with the largest label in  $G_{t_0}$ , i.e. the “newest” vertex in  $G_{t_0}$ . We essentially follow the same process as above, except we never remove  $v$ . We will consider the random variables

$$\begin{aligned} N_t &= \text{number of vertices of } G_t - 1 \\ J_t &= \text{number of vertices older than } v \text{ in } G_t \end{aligned}$$

Also let  $t_f = t_0 + \lfloor cn^{2p}/\log^3 n \rfloor$ .

**Theorem 1.** *With probability at least  $1 - \exp(-\theta(\log^2 n))$ , we have*

$$\begin{aligned} N_t &= n + (2p - 1)(t - t_0) + O(n^p \log n) \\ J_t &= n \left( \frac{N_t}{n} \right)^{\frac{p-1}{2p-1}} (1 + O(\log^{-1/2} n)) \end{aligned}$$

for all  $t \in [t_0, t_f]$ .

The proof will employ the differential equations method, a method used to obtain concentrations for random variables of a discrete random process. For more information, see work of Wormald ([5]) and Bohman ([1]). Essentially, the idea is to notice that

$$\begin{aligned} E(N_{t+1} - N_t | G_t) &= 2p - 1 \\ E(J_{t+1} - J_t | G_t) &= -(1 - p) \frac{J_t}{N_t} \end{aligned}$$

and then define deterministic functions that have the above as their derivatives, i.e.

$$\begin{aligned} z'(x) &= 2p - 1 \\ y'(x) &= -(1 - p) \frac{y}{z} \end{aligned}$$

with initial conditions  $z(t_0/n) = y(t_0/n) = 1$ . (We have scaled the system by a factor of  $n$  here.) We can then obtain the following solution to the differential equation involving  $z$  and  $y$ .

$$\begin{aligned} (2p - 1) \log y + (1 - p) \log z &= C_1 \\ z - (2p - 1)x &= C_2 \end{aligned}$$

We will then use the following lemma to show that  $N_t$  and  $J_t$  are concentrated around the “expected” solutions  $z(t/n)n$  and  $y(t/n)n$ , respectively.

**Lemma 1.** Let  $G_0, \dots, G_t$  be a random process and  $X_i$  a random variable determined by  $G_0, \dots, G_i$ . If there are  $b, c_i \in \mathbb{R}$  such that

$$\begin{aligned} E(X_i - X_{i-1} | G_0, \dots, G_{i-1}) &< b \\ |X_i - X_{i-1} - b| &\leq c_i \end{aligned}$$

then

$$P(\exists i \text{ such that } X_i - X_0 \geq ib + \alpha) \leq \exp\left(-\frac{\alpha^2}{2 \sum_{j=0}^i c_j^2}\right)$$

See [5] for a proof. Now we will prove (half of) the main theorem - in particular, we will show concentration for  $N_t$ .

*Proof.* Define  $F(N_t) = N_t - (2p - 1)t$ . We calculate

$$\begin{aligned} F(N_{t+1}) - F(N_t) &= N_{t+1} - N_t - (2p - 1) \\ E(F(N_{t+1}) - F(N_t)) &= E(N_{t+1} - N_t) - (2p - 1) \\ &= 0 \\ |F(N_{t+1}) - F(N_t)| &\leq |N_{t+1} - N_t| + (2p - 1) \\ &\leq 1 + 2p - 1 \\ &\leq 2p \\ &\leq 2 \end{aligned}$$

because  $N_t$  can change by at most 1 in each step. Now we apply the lemma with  $b = 0$ ,  $c_i = 2$ , and  $\alpha = n^p \log n$  to both  $F(N_t)$  and  $-F(N_t)$  to get

$$|N_t - N_{t_0} - (2p - 1)(t - t_0)| \leq n^p \log n$$

with probability greater than

$$\begin{aligned} 1 - \exp\left(-\frac{n^{2p} \log^2 n}{4(t_f - t_0)}\right) &\geq 1 - \exp(-\theta(\log^5 n)) \\ &\geq 1 - \exp(-\theta(\log^2 n)) \end{aligned}$$

by the definition of  $t_f$ . Since  $N_{t_0} = n - 1$ , this gives us the concentration of  $N_t$  that we desired.

The analysis for the other half of the proof is slightly more difficult. It requires an additional parameter, a stopping time  $T$ .  $T$  is defined so that the desired concentration occurs for all  $t$  between  $t_0$  and  $T$ . Then Prałat and Wormald show that, with high probability,  $T = t_f$ . For more details, see [4].  $\square$

## References

- [1] Tom Bohman, *The triangle-free process*, Advances in Mathematics **221** (2009), 1653–1677.
- [2] Fan Chung and Linyuan Lu, *Coupling online and offline analyses for random power law graphs*, Internet Mathematics **4** (2004), 409–461.
- [3] Pawel Prałat, *Protean graphs*, Internet Mathematics **3** (2006), 21–40.
- [4] Pawel Prałat and Nicholas Wormald, *Growing protean graphs*, Internet Mathematics **4** (2007), 1–16.
- [5] Nicholas Wormald, *The differential equation method for random graph processes and greedy algorithms*, Lectures on Approximation and Randomized Algorithms (1999), 73–155.