

De-Preferential Attachment Random Graphs

Antar Bandyopadhyay

(Joint work with Subhabrata Sen)



**Theoretical Statistics and Mathematics Unit
Indian Statistical Institute, New Delhi and Kolkata**

<http://www.isid.ac.in/~antar>

Indo-German Workshop on Algorithms
Indian Statistical Institute, Kolkata, India
March 10, 2015

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

A Simple Predator-Prey Ecosystem

- Suppose we model an *ecosystem/food-chain* starting with one species where every new species which arrives later is a predator to the existing ones.

A Simple Predator-Prey Ecosystem

- Suppose we model an *ecosystem/food-chain* starting with one species where every new species which arrives later is a predator to the existing ones.
- It is natural to believe that given a choice, a new predator will like to choose its food from the existing species which are not eaten by many.

A Simple Predator-Prey Ecosystem

- Suppose we model an *ecosystem/food-chain* starting with one species where every new species which arrives later is a predator to the existing ones.
- It is natural to believe that given a choice, a new predator will like to choose its food from the existing species which are not eaten by many.
- In other words, a new predator will have *less incentive* or *less preference* to choose its prey from the existing species which have many predators.

A Simple Predator-Prey Ecosystem

- If we now define a graph with vertices as the species and the edges/links between vertices through the predator – prey relation, then such a graph should be modeled by

A Simple Predator-Prey Ecosystem

- If we now define a graph with vertices as the species and the edges/links between vertices through the predator – prey relation, then such a graph should be modeled by

A new vertex prefer to join to an existing vertex with less degree.

A Simple Predator-Prey Ecosystem

- If we now define a graph with vertices as the species and the edges/links between vertices through the predator – prey relation, then such a graph should be modeled by

A new vertex prefer to join to an existing vertex with less degree.

- This is opposite of the usual “*rich get richer model*”, also known as, *preferential attachment model* [Barabási and Albert (1999)].

A Simple Predator-Prey Ecosystem

- If we now define a graph with vertices as the species and the edges/links between vertices through the predator – prey relation, then such a graph should be modeled by

A new vertex prefer to join to an existing vertex with less degree.

- This is opposite of the usual “*rich get richer model*”, also known as, *preferential attachment model* [Barabási and Albert (1999)].
- We will call any such model a *de-preferential attachment model*.

A Simple Predator-Prey Ecosystem

- If we now define a graph with vertices as the species and the edges/links between vertices through the predator – prey relation, then such a graph should be modeled by

A new vertex prefer to join to an existing vertex with less degree.

- This is opposite of the usual “*rich get richer model*”, also known as, *preferential attachment model* [Barabási and Albert (1999)].
- We will call any such model a *de-preferential attachment model*.
- Our goal will be to study such a model rigorously and compare its properties with the preferential attachment model.

De-Preferential Attachment Random Graphs

- Like in the preferential attachment model we will start with an initial graph G_1 with possibly just one vertex.

De-Preferential Attachment Random Graphs

- Like in the preferential attachment model we will start with an initial graph G_1 with possibly just one vertex.
- We will then grow this graph in a random manner as follows.

De-Preferential Attachment Random Graphs

- Like in the preferential attachment model we will start with an initial graph G_1 with possibly just one vertex.
- We will then grow this graph in a random manner as follows.
- At every (discrete) time $n + 1 \geq 2$, we will add one new vertex, say v_{n+1} to the existing graph, say G_n , by letting it to join to the existing vertices $\{v_1, v_2, \dots, v_n\}$.

De-Preferential Attachment Random Graphs

- Like in the preferential attachment model we will start with an initial graph G_1 with possibly just one vertex.
- We will then grow this graph in a random manner as follows.
- At every (discrete) time $n + 1 \geq 2$, we will add one new vertex, say v_{n+1} to the existing graph, say G_n , by letting it to join to the existing vertices $\{v_1, v_2, \dots, v_n\}$.
- The mechanism in which v_{n+1} joins to the existing vertices will be random but with preference for vertices with lesser degree.

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?
 - We will initially consider the case when each new vertex will join only to one existing vertex.

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?
 - We will initially consider the case when each new vertex will join only to one existing vertex.
 - Note that this will lead to a tree (good for modeling food-chain network).

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?
 - We will initially consider the case when each new vertex will join only to one existing vertex.
 - Note that this will lead to a tree (good for modeling food-chain network).
 - We will also consider the case when each new vertex is going to join to $m \geq 1$ existing vertices where m will be a fixed positive integer.

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?
 - We will initially consider the case when each new vertex will join only to one existing vertex.
 - Note that this will lead to a tree (good for modeling food-chain network).
 - We will also consider the case when each new vertex is going to join to $m \geq 1$ existing vertices where m will be a fixed positive integer.
 - In this case we can have *multiple edges* and *self-loops* depending on the mechanism in which the m new links will be formed. Also there can be formation of cycles.

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (i) How many existing vertices are going to be joined with a new vertex?
 - We will initially consider the case when each new vertex will join only to one existing vertex.
 - Note that this will lead to a tree (good for modeling food-chain network).
 - We will also consider the case when each new vertex is going to join to $m \geq 1$ existing vertices where m will be a fixed positive integer.
 - In this case we can have *multiple edges* and *self-loops* depending on the mechanism in which the m new links will be formed. Also there can be formation of cycles.
 - None of these are good for a food-chain network, as A multiply eats B or A eats itself or even A eats B which eats C but C eats A are not suitable for such a network.

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (ii) How do we make the choice *de-preferential*, that is, not preferring vertices of higher degree?

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (ii) How do we make the choice *de-preferential*, that is, not preferring vertices of higher degree?

We will consider two types of attachment mechanisms, namely,

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (ii) How do we make the choice *de-preferential*, that is, not preferring vertices of higher degree?

We will consider two types of attachment mechanisms, namely,

Linear	Inverse
--------	---------

De-Preferential Attachment Random Graphs

- To make things rigorous we need to fix couple of issues:
 - (ii) How do we make the choice *de-preferential*, that is, not preferring vertices of higher degree?

We will consider two types of attachment mechanisms, namely,

Linear	Inverse
probab. \propto const. - degree	probab. $\propto 1/\text{degree}$

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$
- For $m \geq 1$ we will imagine that v_{n+1} comes with m *half-edges*, which we will denote by $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$.

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$
- For $m \geq 1$ we will imagine that v_{n+1} comes with m *half-edges*, which we will denote by $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$.
- We will write $d_i(n+1, k)$, for $k = 0, \dots, m$, to denote the degree of the vertex v_i , $i = 1, \dots, n$, after k half-edges of v_{n+1} have been attached.

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$
- For $m \geq 1$ we will imagine that v_{n+1} comes with m *half-edges*, which we will denote by $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$.
- We will write $d_i(n+1, k)$, for $k = 0, \dots, m$, to denote the degree of the vertex v_i , $i = 1, \dots, n$, after k half-edges of v_{n+1} have been attached.
- We will write $d_i(n+1, 0) = d_i(n)$ for any $1 \leq i \leq n$ and note $d_n(n) = m$.

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$
- For $m \geq 1$ we will imagine that v_{n+1} comes with m *half-edges*, which we will denote by $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$.
- We will write $d_i(n+1, k)$, for $k = 0, \dots, m$, to denote the degree of the vertex v_i , $i = 1, \dots, n$, after k half-edges of v_{n+1} have been attached.
- We will write $d_i(n+1, 0) = d_i(n)$ for any $1 \leq i \leq n$ and note $d_n(n) = m$.
- Let $\left\{ \mathcal{F}_{n,k} \mid 0 \leq k \leq m-1, n \geq 1 \right\}$ be the natural filtration of the random attachments.

Some Notations

- We will denote the growing random graph sequence by $(G_n)_{n \geq 1}$.
- The vertices will be labeled as $v_1, v_2, \dots, v_n, \dots$
- For $m \geq 1$ we will imagine that v_{n+1} comes with m *half-edges*, which we will denote by $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$.
- We will write $d_i(n+1, k)$, for $k = 0, \dots, m$, to denote the degree of the vertex v_i , $i = 1, \dots, n$, after k half-edges of v_{n+1} have been attached.
- We will write $d_i(n+1, 0) = d_i(n)$ for any $1 \leq i \leq n$ and note $d_n(n) = m$.
- Let $\left\{ \mathcal{F}_{n,k} \mid 0 \leq k \leq m-1, n \geq 1 \right\}$ be the natural filtration of the random attachments.
- If $m = 1$ then we will simply write the natural filtration as $\{\mathcal{F}_n\}_{n \geq 1}$.

Models for $m = 1$

- We start with G_1 which consists of one vertex with one unattached *half-edge*. So $d_1(1) = 1$.

Models for $m = 1$

- We start with G_1 which consists of one vertex with one unattached *half-edge*. So $d_1(1) = 1$.

- **Linear De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto ((2n - 1) - d_i(n)),$$

Models for $m = 1$

- We start with G_1 which consists of one vertex with one unattached *half-edge*. So $d_1(1) = 1$.

- **Linear De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto ((2n - 1) - d_i(n)),$$

that is,

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) = \frac{1}{n - 1} \left(1 - \frac{d_i(n)}{2n - 1} \right),$$

Models for $m = 1$

- We start with G_1 which consists of one vertex with one unattached *half-edge*. So $d_1(1) = 1$.

- **Linear De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto ((2n - 1) - d_i(n)),$$

that is,

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) = \frac{1}{n - 1} \left(1 - \frac{d_i(n)}{2n - 1} \right),$$

- **Inverse De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto \frac{1}{d_i(n)},$$

Models for $m = 1$

- We start with G_1 which consists of one vertex with one unattached *half-edge*. So $d_1(1) = 1$.

- **Linear De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto ((2n - 1) - d_i(n)),$$

that is,

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) = \frac{1}{n - 1} \left(1 - \frac{d_i(n)}{2n - 1} \right),$$

- **Inverse De-Preferential Model:**

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) \propto \frac{1}{d_i(n)},$$

that is,

$$\mathbf{P} \left(v_{n+1} \longrightarrow v_i \mid \mathcal{F}_n \right) = \frac{C_n}{d_i(n)},$$

where $C_n^{-1} =: D_n = \sum_{i=1}^n \frac{1}{d_i(n)}$.

Models for $m > 1$

- We start with G_1 which consists of one vertex with m unattached *half-edges*. So $d_1(1) = m$.

Models for $m > 1$

- We start with G_1 which consists of one vertex with m unattached *half-edges*. So $d_1(1) = m$.
- At time $n + 1$, the new vertex v_{n+1} comes with m *half-edges*, namely, $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$, which are joined **sequentially by updating the degrees of the existing vertices and are not allowed to join to v_{n+1}** .

Models for $m > 1$

- We start with G_1 which consists of one vertex with m unattached *half-edges*. So $d_1(1) = m$.
- At time $n + 1$, the new vertex v_{n+1} comes with m *half-edges*, namely, $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$, which are joined **sequentially by updating the degrees of the existing vertices and are not allowed to join to v_{n+1}** .
- This prevents the formation of the *self-loops*.

Models for $m > 1$

- We start with G_1 which consists of one vertex with m unattached *half-edges*. So $d_1(1) = m$.
- At time $n + 1$, the new vertex v_{n+1} comes with m *half-edges*, namely, $e_{n+1,1}, e_{n+1,2}, \dots, e_{n+1,m}$, which are joined **sequentially by updating the degrees of the existing vertices and are not allowed to join to v_{n+1}** .
- This prevents the formation of the *self-loops*.
- We still have the possibility of having *multiple edges* between two vertices.

Models for $m > 1$

- **Linear De-Preferential Model:**

$$\mathbf{P} \left(e_{n+1,k+1} = \{v_j, v_{n+1}\} \mid \mathcal{F}_{n+1,k} \right) = \frac{1}{n-1} \left(1 - \frac{d_j(n+1, k)}{k + (2n-1)m} \right)$$

Models for $m > 1$

- Linear De-Preferential Model:**

$$\mathbf{P} \left(e_{n+1,k+1} = \{v_j, v_{n+1}\} \mid \mathcal{F}_{n+1,k} \right) = \frac{1}{n-1} \left(1 - \frac{d_j(n+1, k)}{k + (2n-1)m} \right)$$

- Inverse De-Preferential Model:**

$$\mathbf{P} \left(e_{n+1,k+1} = \{v_j, v_{n+1}\} \mid \mathcal{F}_{n+1,k} \right) = C_{n+1,k} \frac{1}{d_j(n+1, k)}$$

$$\text{where } C_{n+1,k}^{-1} =: D_{n+1,k} = \sum_{j=1}^n \frac{1}{d_j(n+1, k)}.$$

Some Earlier Work

- A somewhat similar, in fact a bit more general model was considered by Sevim and Rikvold (2006, 2008).

Some Earlier Work

- A somewhat similar, in fact a bit more general model was considered by Sevim and Rikvold (2006, 2008).
- They obtain by some intuitive arguments (not quite rigorous) the asymptotic degree distribution and validated their claims by simulation results.

Some Earlier Work

- A somewhat similar, in fact a bit more general model was considered by Sevim and Rikvold (2006, 2008).
- They obtain by some intuitive arguments (not quite rigorous) the asymptotic degree distribution and validated their claims by simulation results.
- Our results support their observations.

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

Main Results: Linear Case with $m = 1$

Theorem 1 (WLLN for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n)}{\log n} \xrightarrow{P} 1.$$

Main Results: Linear Case with $m = 1$

Theorem 1 (WLLN for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n)}{\log n} \xrightarrow{P} 1.$$

Theorem 2 (CLT for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n) - \log n}{\sqrt{\log n}} \xrightarrow{d} \text{Normal}(0, 1).$$

Main Results: Linear Case with $m = 1$

Theorem 3 (Asymptotic degree distribution)

Let $P_k(n)$ be the proportion of vertices in G_n with degree $k \geq 1$. Then for any $k \geq 1$,

$$P_k(n) \longrightarrow \frac{1}{2^k} \text{ a.s.}$$

Main Results: Linear Case with $m = 1$

Theorem 3 (Asymptotic degree distribution)

Let $P_k(n)$ be the proportion of vertices in G_n with degree $k \geq 1$. Then for any $k \geq 1$,

$$P_k(n) \longrightarrow \frac{1}{2^k} \text{ a.s.}$$

Remark: The *asymptotic degree distribution* of G_n is Geometric $(\frac{1}{2})$ which has mean 2, mode 1 and exponential tail.

Main Results: Linear Case with $m = 1$

Theorem 4 (Asymptotic degree distribution of the chosen vertex)

Let U_{n+1} be the (random) selected vertex from $\{v_1, v_2, \dots, v_n\}$ where the new vertex v_{n+1} connects. Then for any $k \geq 1$,

$$\mathbf{P}(\text{degree}_{G_n}(U_{n+1}) = k) \longrightarrow \frac{1}{2^k}.$$

Main Results: Linear Case with $m \geq 1$

Theorem 5 (WLLN for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n)}{\log n} \xrightarrow{P} m.$$

Main Results: Linear Case with $m \geq 1$

Theorem 5 (WLLN for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n)}{\log n} \xrightarrow{P} m.$$

Theorem 6 (CLT for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n) - m \log n}{\sqrt{m \log n}} \xrightarrow{d} \text{Normal}(0, 1).$$

Main Results: Inverse Case with $m = 1$

Theorem 7 (SLLN for fixed vertex degree)

Fix a vertex $i \geq 1$ then

$$\frac{d_i(n)}{\sqrt{\log n}} \longrightarrow \sqrt{\frac{2}{\lambda^*}} \text{ a.s.,}$$

where $\lambda^* > 0$ is the unique positive solution of the equation

$$\sum_{n=1}^{\infty} \prod_{i=1}^n \frac{1}{1 + i\lambda^*} = 1.$$

Main Results: Inverse Case with $m = 1$

Theorem 8 (Asymptotic degree distribution)

Let $P_k(n)$ be the proportion of vertices in G_n with degree $k \geq 1$. Then for any $k \geq 1$,

$$P_k(n) \longrightarrow k\lambda^* \prod_{i=1}^k \frac{1}{i\lambda^* + 1} \text{ a.s.}$$

Main Results: Inverse Case with $m = 1$

Theorem 8 (Asymptotic degree distribution)

Let $P_k(n)$ be the proportion of vertices in G_n with degree $k \geq 1$. Then for any $k \geq 1$,

$$P_k(n) \longrightarrow k\lambda^* \prod_{i=1}^k \frac{1}{i\lambda^* + 1} \text{ a.s.}$$

Remark: The *asymptotic degree distribution* of G_n has mean 2, mode 1 and thin tail.

Main Results: Inverse Case with $m = 1$

Theorem 9 (Asymptotic degree distribution of the chosen vertex)

Let U_{n+1} be the (random) selected vertex from $\{v_1, v_2, \dots, v_n\}$ where the new vertex v_{n+1} connects. Then for any $k \geq 1$,

$$\mathbf{P}(\text{degree}_{G_n}(U_{n+1}) = k) \longrightarrow \prod_{i=1}^k \frac{1}{i\lambda^* + 1}.$$

Main Results: Inverse Case with $m > 1$

Theorem 10 (“WLLN” for fixed vertex degree)

\exists constants $0 < C_1 < C_2 < \infty$ such that for any fixed vertex i ,

$$\mathbf{P} \left(C_1 \leq \frac{d_i(n)}{m\sqrt{\log n}} \leq C_2 \right) \rightarrow 1,$$

as $n \rightarrow \infty$.

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

Techniques Used for the Linear De-Preferential Case

- This is similar to the preferential attachment random graph models and the main tools are martingale techniques.

Techniques Used for the Linear De-Preferential Case

- This is similar to the preferential attachment random graph models and the main tools are martingale techniques.
- In this case $m = 1$ and $m > 1$ are not much different.

Techniques Used for the Linear De-Preferential Case

- This is similar to the preferential attachment random graph models and the main tools are martingale techniques.
- In this case $m = 1$ and $m > 1$ are not much different.
- For the CLTs we use martingale CLT.

Techniques Used for the Inverse De-Preferential Case

- We used two different *embeddings/couplings* for this case.

Techniques Used for the Inverse De-Preferential Case

- We used two different *embeddings/couplings* for this case.
- One type of embedding for $m = 1$ and a different embedding for $m > 1$.

Techniques Used for the Inverse De-Preferential with $m = 1$

- We consider a continuous time age dependent branching process and keep all the statistics, that is, entire growing tree structure.

Techniques Used for the Inverse De-Preferential with $m = 1$

- We consider a continuous time age dependent branching process and keep all the statistics, that is, entire growing tree structure.
- Formally, let \mathcal{G} be the set of all finite rooted tree. We consider a continuous time process $\{\Upsilon(t) : t \geq 0\}$ of randomly growing trees on \mathcal{G} .

Techniques Used for the Inverse De-Preferential with $m = 1$

- We consider a continuous time age dependent branching process and keep all the statistics, that is, entire growing tree structure.
- Formally, let \mathcal{G} be the set of all finite rooted tree. We consider a continuous time process $\{\Upsilon(t) : t \geq 0\}$ of randomly growing trees on \mathcal{G} .
- $\Upsilon(0)$ is a single vertex (root) with a half-edge (so degree is 1).

Techniques Used for the Inverse De-Preferential with $m = 1$

- We consider a continuous time age dependent branching process and keep all the statistics, that is, entire growing tree structure.
- Formally, let \mathcal{G} be the set of all finite rooted tree. We consider a continuous time process $\{\Upsilon(t) : t \geq 0\}$ of randomly growing trees on \mathcal{G} .
- $\Upsilon(0)$ is a single vertex (root) with a half-edge (so degree is 1).
- Each vertex reproduces independently according to identical copies of a age dependent pure birth process $(\xi(t))_{t \geq 0}$ such that $\mathbf{P}(\xi(0) = 1) = 1$ and

$$\mathbf{P}\left(\xi(t+h) = k+1 \mid \xi(t) = k\right) = \frac{h}{k+1} + o(h).$$

Techniques Used for the Inverse De-Preferential with $m = 1$

- We consider a continuous time age dependent branching process and keep all the statistics, that is, entire growing tree structure.
- Formally, let \mathcal{G} be the set of all finite rooted tree. We consider a continuous time process $\{\Upsilon(t) : t \geq 0\}$ of randomly growing trees on \mathcal{G} .
- $\Upsilon(0)$ is a single vertex (root) with a half-edge (so degree is 1).
- Each vertex reproduces independently according to identical copies of a age dependent pure birth process $(\xi(t))_{t \geq 0}$ such that $\mathbf{P}(\xi(0) = 1) = 1$ and

$$\mathbf{P}\left(\xi(t+h) = k+1 \mid \xi(t) = k\right) = \frac{h}{k+1} + o(h).$$

- This process is an example of a *Crump-Mode-Jagers (CMJ) branching process* [Crump and Mode (1968) and Jagers (1969)].

Techniques Used for the Inverse De-Preferential with $m = 1$

Embedding Theorem for $m = 1$

Starting with $\tau_1 = 0$ consider the following sequence of stopping times

$$\tau_n := \inf\{t \geq \tau_{n-1} \mid |\Upsilon(t)| = n\}.$$

For $m = 1$, the sequence of random graphs $\{G_n\}_{n=1}^\infty$ have the same distribution as the sequence of random trees $\{\Upsilon(\tau_n)\}_{n=1}^\infty$.

Techniques Used for the Inverse De-Preferential with $m = 1$

Embedding Theorem for $m = 1$

Starting with $\tau_1 = 0$ consider the following sequence of stopping times

$$\tau_n := \inf\{t \geq \tau_{n-1} \mid |\Upsilon(t)| = n\}.$$

For $m = 1$, the sequence of random graphs $\{G_n\}_{n=1}^\infty$ have the same distribution as the sequence of random trees $\{\Upsilon(\tau_n)\}_{n=1}^\infty$.

Remarks:

- (i) This is immediate from the construction of the CMJ branching process.

Techniques Used for the Inverse De-Preferential with $m = 1$

Embedding Theorem for $m = 1$

Starting with $\tau_1 = 0$ consider the following sequence of stopping times

$$\tau_n := \inf\{t \geq \tau_{n-1} \mid |\Upsilon(t)| = n\}.$$

For $m = 1$, the sequence of random graphs $\{G_n\}_{n=1}^\infty$ have the same distribution as the sequence of random trees $\{\Upsilon(\tau_n)\}_{n=1}^\infty$.

Remarks:

- (i) This is immediate from the construction of the CMJ branching process.
- (ii) For studying preferential attachment model with non-linear weights a similar observation was made by Rudas and Tóth (2007).

Techniques Used for the Inverse De-Preferential with $m = 1$

- Let $\hat{\rho}(\lambda)$ be the expected Laplace transform of the pure birth process $(\xi(t))_{t \geq 0}$.

Techniques Used for the Inverse De-Preferential with $m = 1$

- Let $\hat{\rho}(\lambda)$ be the expected Laplace transform of the pure birth process $(\xi(t))_{t \geq 0}$.
- Then it is easy to see that

$$\hat{\rho}(\lambda) = \sum_{n=1}^{\infty} \prod_{i=1}^n \frac{1}{i\lambda + 1}.$$

Techniques Used for the Inverse De-Preferential with $m = 1$

- Let $\hat{\rho}(\lambda)$ be the expected Laplace transform of the pure birth process $(\xi(t))_{t \geq 0}$.
- Then it is easy to see that

$$\hat{\rho}(\lambda) = \sum_{n=1}^{\infty} \prod_{i=1}^n \frac{1}{i\lambda + 1}.$$

- Thus $\hat{\rho}(\lambda) = 1$ has a unique positive solution which we denote by λ^* .

Techniques Used for the Inverse De-Preferential with $m = 1$

- Let $\hat{\rho}(\lambda)$ be the expected Laplace transform of the pure birth process $(\xi(t))_{t \geq 0}$.
- Then it is easy to see that

$$\hat{\rho}(\lambda) = \sum_{n=1}^{\infty} \prod_{i=1}^n \frac{1}{i\lambda + 1}.$$

- Thus $\hat{\rho}(\lambda) = 1$ has a unique positive solution which we denote by λ^* .
- $\lambda^* > 0$ is called the *Malthusian parameter* for the (supercritical) CMJ process.

Techniques Used for the Inverse De-Preferential with $m = 1$

Theorem A of Nerman (1961)

Suppose $\{\Upsilon(t) : t \geq 0\}$ is a (supercritical) CMJ process with Multhusian parameter λ^* and let $\phi : \mathcal{G} \rightarrow \mathbb{R}$ be bounded function. Then the following limit holds almost surely

$$\lim_{t \rightarrow \infty} \frac{1}{|\Upsilon(t)|} \sum_{x \in \Upsilon(t)} \phi(\Upsilon(t)_{\downarrow x}) = \lambda^* \int_0^\infty \exp\{-\lambda^* t\} \mathbf{E}(\phi(\Upsilon(t))) dt,$$

where for a tree $\mathcal{T} \in \mathcal{G}$ and a vertex $x \in \mathcal{T}$ we define $\mathcal{T}_{\downarrow x}$ as the sub-tree rooted at x consisting of all the descendants of x .

Techniques Used for the Inverse De-Preferential with $m = 1$

Theorem A of Nerman (1961)

Suppose $\{\Upsilon(t) : t \geq 0\}$ is a (supercritical) CMJ process with Multihusian parameter λ^* and let $\phi : \mathcal{G} \rightarrow \mathbb{R}$ be bounded function. Then the following limit holds almost surely

$$\lim_{t \rightarrow \infty} \frac{1}{|\Upsilon(t)|} \sum_{x \in \Upsilon(t)} \phi(\Upsilon(t)_{\downarrow x}) = \lambda^* \int_0^\infty \exp\{-\lambda^* t\} \mathbf{E}(\phi(\Upsilon(t))) dt,$$

where for a tree $\mathcal{T} \in \mathcal{G}$ and a vertex $x \in \mathcal{T}$ we define $\mathcal{T}_{\downarrow x}$ as the sub-tree rooted at x consisting of all the descendants of x .

Remark: This proves the SLLN for the degree of a fixed vertex and also the asymptotic degree distribution in the inverse de-preferential case.

Techniques Used for the Inverse De-Preferential with $m > 1$

- For this we use a different technique similar to the *Athreya-Karlin Embedding*.

Techniques Used for the Inverse De-Preferential with $m > 1$

- For this we use a different technique similar to the *Athreya-Karlin Embedding*.
- Let $\{Z(t) : t \geq 0\}$ be a pure birth process with $\mathbf{P}(Z(0) = m) = 1$ and birth rates $\lambda_i = \frac{1}{i}$.

Techniques Used for the Inverse De-Preferential with $m > 1$

- For this we use a different technique similar to the *Athreya-Karlin Embedding*.
- Let $\{Z(t) : t \geq 0\}$ be a pure birth process with $\mathbf{P}(Z(0) = m) = 1$ and birth rates $\lambda_i = \frac{1}{i}$.
- For $i \geq 1$, let $(Z_i(t))_{t \geq 0}$ be i.i.d. copies of the pure birth process $(Z(t))_{t \geq 0}$.

Techniques Used for the Inverse De-Preferential with $m > 1$

- For this we use a different technique similar to the *Athreya-Karlin Embedding*.
- Let $\{Z(t) : t \geq 0\}$ be a pure birth process with $\mathbf{P}(Z(0) = m) = 1$ and birth rates $\lambda_i = \frac{1}{i}$.
- For $i \geq 1$, let $(Z_i(t))_{t \geq 0}$ be i.i.d. copies of the pure birth process $(Z(t))_{t \geq 0}$.
- We recursively define the following stopping times starting with $\tau_1 = 0$,

$$\tau_2 := \inf \left\{ t \geq 0 \mid Z_1(t) - m = m \right\}$$

$$\tau_3 := \inf \left(t \geq \tau_2 \mid Z_1(t) + Z_2(t - \tau_2) - 2m = m \right)$$

$$\vdots \quad \vdots \quad \vdots$$

$$\tau_{n+1} := \inf \left(t \geq \tau_n \mid Z_1(t) + Z_2(t - \tau_2) + \cdots + Z_n(t - \tau_n) - n m = m \right)$$

Techniques Used for the Inverse De-Preferential with $m > 1$

Embedding Theorem for $m > 1$

For $m \geq 1$, the two sequence of random variables, namely, $\left\{ (d_i(n))_{i=1}^n \mid n \geq 1 \right\}$ and $\left\{ (Z_i(\tau_n - \tau_i))_{i=1}^n \mid n \geq 1 \right\}$ has the same distribution.

Techniques Used for the Inverse De-Preferential with $m > 1$

Embedding Theorem for $m > 1$

For $m \geq 1$, the two sequence of random variables, namely, $\left\{ (d_i(n))_{i=1}^n \mid n \geq 1 \right\}$ and $\left\{ (Z_i(\tau_n - \tau_i))_{i=1}^n \mid n \geq 1 \right\}$ has the same distribution.

WLLN for the Pure Birth Process

Let $\{Z(t) : t \geq 0\}$ be a pure birth process with $\mathbf{P}(Z(0) = m) = 1$ and birth rates $\lambda_i = \frac{1}{i}$. Then

$$\frac{Z(t)}{\sqrt{t}} \xrightarrow{P} \sqrt{2}.$$

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

Initial Configuration

- The special initial configuration we have chosen is not necessary to derive most of the results.

Initial Configuration

- The special initial configuration we have chosen is not necessary to derive most of the results.
- For example all results in the linear case go through for starting with any finite graph.

Initial Configuration

- The special initial configuration we have chosen is not necessary to derive most of the results.
- For example all results in the linear case go through for starting with any finite graph.
- But it is necessary assumption for the results on inverse case which we prove using the embedding to CMJ branching process.

- 1 Introduction
 - A Toy Model for a Ecosystem/Food-Chain
 - De-Preferential Attachment Model
- 2 Model Description
 - Notations
 - Models for $m = 1$
 - Models for $m > 1$
 - Earlier Work
- 3 Main Results
 - For the Linear De-Preferential Case
 - For the Inverse De-Preferential Case
- 4 Techniques Used in the Proofs
 - Linear Case
 - Inverse Case
- 5 Special Initial Condition
- 6 Open Problems

Open Problems

- The main question which remains open, is the asymptotic degree distribution for $m > 1$ case. Particularly, for the inverse de-preferential model.

Open Problems

- The main question which remains open, is the asymptotic degree distribution for $m > 1$ case. Particularly, for the inverse de-preferential model.
- There is a formula derived by non-rigorous methods [Sevim and Rikvold (2008)] which can be validated by simulation but no rigorous proof is available.

Open Problems

- The main question which remains open, is the asymptotic degree distribution for $m > 1$ case. Particularly, for the inverse de-preferential model.
- There is a formula derived by non-rigorous methods [Sevim and Rikvold (2008)] which can be validated by simulation but no rigorous proof is available.
- For $m > 1$ case it seems that the Athreya-Karlin Embedding technique is fairly unsatisfactory for the inverse de-preferential case. Proofs of a complete WLLN and CLT remain open for the degree of a fixed vertex.

Open Problems

- The main question which remains open, is the asymptotic degree distribution for $m > 1$ case. Particularly, for the inverse de-preferential model.
- There is a formula derived by non-rigorous methods [Sevim and Rikvold (2008)] which can be validated by simulation but no rigorous proof is available.
- For $m > 1$ case it seems that the Athreya-Karlin Embedding technique is fairly unsatisfactory for the inverse de-preferential case. Proofs of a complete WLLN and CLT remain open for the degree of a fixed vertex.
- For $m = 1$ case one should remove the dependency on the initial configuration but it seems it is a technically very difficult problem!

Thank You