

Strong Bounds for Evolution in Networks

Paul G. Spirakis^{1,2}

ESRC workshop on Algorithmic Game Theory
Department of Mathematics, London School of Economics

These results have been presented in:

- **Theor. Comp. Science 2013:** Natural Models for Evolution on Networks, by *G. Mertzios*³, *S. Nikolettseas*¹, *C. Raptopoulos*¹, and *P. Spirakis*^{1,2}
- **SODA 2012; Algorithmica:** Approximating Fixation Probabilities in the Generalized Moran Process, by *J. Díaz*⁴, *L.A. Goldberg*⁵, *G. Mertzios*³, *D. Richerby*⁵, *M. Serna*⁴, and *P. Spirakis*^{1,2}
- **ICALP 2013:** Strong Bounds for Evolution in Undirected Graphs, by *G. Mertzios*³ and *P. Spirakis*^{1,2}

¹CTI & University of Patras, Greece, ²University of Liverpool, UK, ³Durham University, UK,
⁴Universitat Politècnica de Catalunya, Spain, ⁵University of Oxford, UK

Evolutionary graph theory

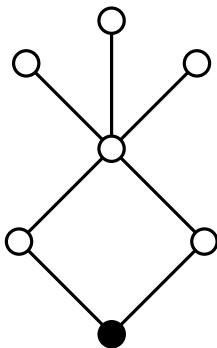
- **Evolution** in biology / **Population dynamics** have been mainly traditionally in **homogeneous** populations
- However, in reality, the **topology** / **structure** of the population can strongly **affect** the **output** of the **dynamics**.
- **Evolutionary graph theory** has been introduced in [Lieberman, Hauert, Nowak, *Nature*, 2005]
- Main idea: arrange the population on a network (i.e. graph)
- There are two types of vertices:
 - **aggressive** (“mutants”) \longleftrightarrow **fitness** $r \geq 1$,
 - **non-aggressive** (“residents”) \longleftrightarrow **fitness** 1.

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- Time is discrete $t = 1, 2, \dots$
- At every iteration $t \geq 1$,
 - **choose** a vertex u with **probability** proportional to its **fitness**;
 - **choose** randomly a **neighbor** $v \in N(u)$ (resp. an **arc** $\langle uv \rangle$);
 - **replace** v by an **offspring** of u .

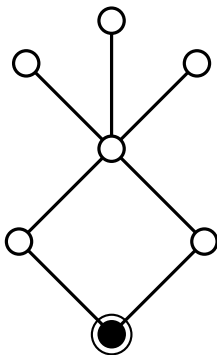
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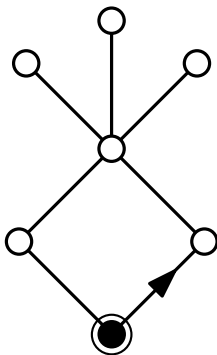
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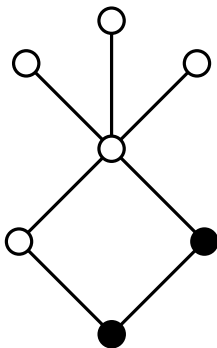
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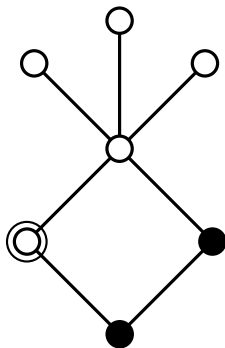
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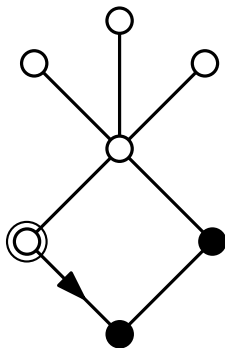
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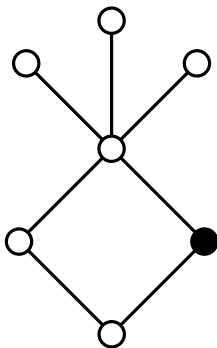
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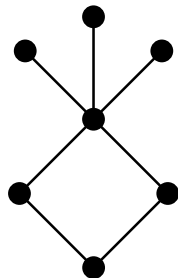
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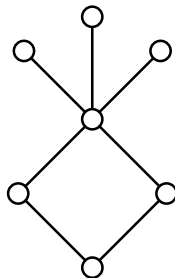
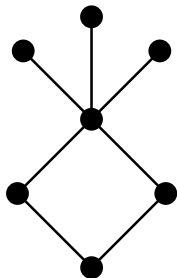
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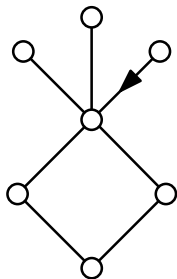
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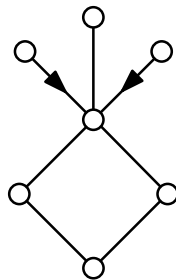
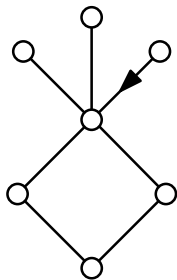
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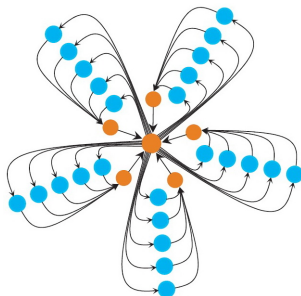
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\Rightarrow We are mainly interested in **undirected** graphs.

Evolutionary graph theory

Theorem (**Isothermal Theorem**, Lieberman et al., *Nature*, 2005)

Let $G = (V, E)$ be an **undirected** and **regular** graph (i.e. $\deg(u) = \deg(v)$ for every $u, v \in V$). If $r > 1$, then $f_r(G) = \frac{1 - \frac{1}{r}}{1 - \frac{1}{r^n}} \approx 1 - \frac{1}{r}$.

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- The complete graph acts as a “benchmark”
- A graph G is called:
 - an **amplifier** if $f_r(G) > \frac{1 - \frac{1}{r}}{1 - \frac{1}{r^n}}$, and
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- **Question 1:** Do there **exist** strong undirected **amplifiers** / **suppressors** of selection?

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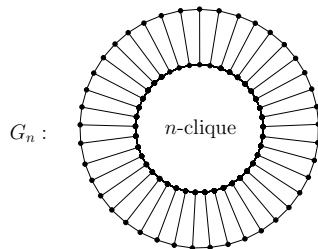
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- **Question 2:** How does the **population structure** affect the **fixation probability**?

A class of undirected suppressors of selection

- For every $n \geq 1$, we define the “clique-wheel” graph G_n with $2n$ vertices:
 - clique with n vertices
 - induced cycle with n vertices
 - perfect matching between them



Theorem (Mertzios, Nikolettseas, Raptopoulos, Spirakis, *TCS*, 2013)

For every $r \in (1, \frac{4}{3})$, the fixation probability of G_n is $f_{G_n}(r) \leq \frac{1}{2}(1 - \frac{1}{r})$, as $n \rightarrow \infty$.

Computation of fixation probabilities

Questions that were open until recently:

- How can we **compute** the **fixation/****extinction** probability for a given graph?
- Can we do this **efficiently**?
 - the resulting Markov chain implies a system of **linear equations**
 - however: **exponentially** many equations – in general one for every vertex subset
- Does the generalized Moran process **reach absorption** (i.e. fixation or extinction) **quickly**?

Nothing is known until now, except immediate results for special cases

- e.g. expected **linear** time for **regular graphs**

Computation of fixation probabilities

Our results: [Díaz, Goldberg, Mertzios, Richerby, Spirakis, *SODA*, 2012; *Algorithmica*, to appear]

- The generalized Moran process reaches **absorption** (either fixation or extinction) in **polynomial** number of steps **with high probability**.
- Two **FPRAS** (*fully polynomial randomized approximation schemes*) for the problems of:
 - computing the **fixation** probability on general graphs for $r \geq 1$
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Definition

An **FPRAS** for a function f is a **randomized algorithm** g that, given **input** X , gives an output satisfying:

$$(1 - \varepsilon)f(X) \leq g(X) \leq (1 + \varepsilon)f(X)$$

with **probability** at least $\frac{3}{4}$ and has running time **polynomial** in $|X|$ and $\frac{1}{\varepsilon}$.

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General approach for the FPRAS:

- **simulate** (polynomially many) times the generalized Moran process until **absorption** is reached
- **count** the number of simulations that reached **fixation**

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General approach for the **FPRAS**:

- **simulate** (polynomially many) times the generalized Moran process **until absorption** is reached
- **count** the number of simulations that reached **fixation**

The **correctness** of the FPRAS is based on **two points**:

- 1 expected **polynomial** time until **absorption** is reached
⇒ every simulation needs polynomial number of steps
- 2 the **fixation** probability is **polynomially upper/lower** bounded (i.e. not too big/small)
⇒ a polynomial number of simulations suffices to estimate the fixation/absorption probabilities.

Upper / lower bounds

So far, the only known general bounds for the fixation probability:

Lemma (Díaz, Goldberg, Mertzios, Richerby, Serna, Spirakis, *SODA*, 2012; *Algorithmica*, to appear)

Let $G = (V, E)$ be an *undirected* graph with n vertices. Then:

- $f_r(G) \geq \frac{1}{n}$ for any $r \geq 1$
 - $f_r(G) \leq 1 - \frac{1}{n+r}$ for any $r > 0$
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- Tighter upper / lower bounds \Rightarrow better running time of these FPRAS.

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⇒ the fixation probability of the graph G is $f_r(G) = \frac{1}{n} \sum_{v \in V} f_r(v)$

- We are interested in finding graphs with many strong / weak starts $f_r(v)$ for the mutant

Universal and selective amplifiers

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Let \mathcal{G} be an **infinite class** of undirected graphs. If for every $r > r_0$ and every graph $G \in \mathcal{G}$ with $n \geq n_0$ vertices:

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Moreover:

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- **$(\Theta(n), n)$ -selective suppressors** are called **strong selective suppressors**

First result:

Theorem

For any function $g(n) = \Omega(n^{\frac{3}{4} + \varepsilon})$, where $\varepsilon > 0$, there exists *no class \mathcal{G} of $g(n)$ -universal amplifiers* for any $r > r_0 = 1$.

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Corollary

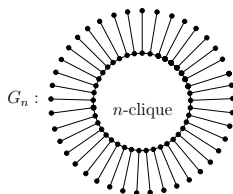
There exists *no infinite class of strong universal amplifiers*.

Our results

Second result:

Theorem

The class $\mathcal{G} = \{G_n : n \geq 1\}$ of *urchin graphs* is a class of $(\frac{n}{2}, n)$ -selective amplifiers.



Our results

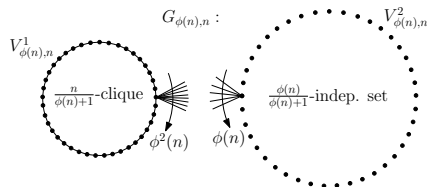
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Therefore there exist:

- strong selective amplifiers
- “quite” strong selective suppressors

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Theorem (Thermal Theorem)

Let $G = (V, E)$ be a connected undirected graph and $r > 1$.

Then $f_r(v) \geq \frac{r-1}{r + \frac{\deg v}{\deg_{\min}}}$ for every $v \in V$.

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- **Almost tight bound**:
 - for **regular** graphs: $f_r(v) \geq \frac{r-1}{r+1}$

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Corollary

In **every graph** G there exists at least one **vertex** v with $f_r(v) \geq \frac{r-1}{r+1}$ (i.e. independent of the size n).

No strong universal amplifiers

Theorem

For any function $g(n) = \Omega(n^{\frac{3}{4} + \varepsilon})$, where $\varepsilon > 0$, there exists *no class \mathcal{G} of $g(n)$ -universal amplifiers* for any $r > 1$.

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Proof sketch (by contradiction).

- Let $g(n) = \Omega(n^{1-\delta})$, where $\delta = \frac{1}{4} - \varepsilon < \frac{1}{4}$
- Suppose that \mathcal{G} is a class of $g(n)$ -universal amplifiers, i.e. for every $r > 1$ and every graph $G \in \mathcal{G}$ with $n \geq n_0$ vertices:

$$f_r(G) \geq 1 - \frac{c(r)}{g(n)} \geq 1 - \frac{c_0(r)}{n^{1-\delta}}$$

for appropriate functions $c(r)$ and $c_0(r)$.

No strong universal amplifiers

Proof sketch (by contradiction).

- We partition the vertices of G into three subsets:

$$V_1 = \left\{ v \in V : f_r(v) \geq 1 - \frac{c_0(r)}{n^{1-\delta}} \right\}$$

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- Since \mathcal{G} is a class of $g(n)$ -universal amplifiers $\Rightarrow V_1 \neq \emptyset$

No strong universal amplifiers

Proof sketch (by contradiction).

We can prove that:

- for every $v \in V_1$:

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Therefore:

- Since $\delta < \frac{1}{4} \Rightarrow 1 - \delta > 1 - 2\delta > 2\delta > \delta$
 \Rightarrow every neighbor of a vertex $v \in V_1 \cup V_2$ must belong to V_3
 $\Rightarrow V_1 \cup V_2$ is an independent set

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Proof sketch (by contradiction).

Using an upper bound from

[Mertziou, Nikolettseas, Raptopoulos, Spirakis, *Theor. Comp. Sci.*, 2013],

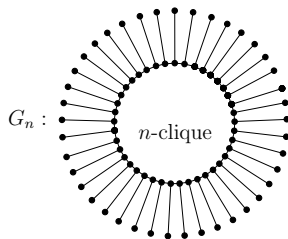
it follows:

- $\Omega(n^{-3\delta}) \leq \frac{c'''(r)}{n^{1-\delta}}$, for some function $c'''(r)$,
- **contradiction** since $\delta < \frac{1}{4}$



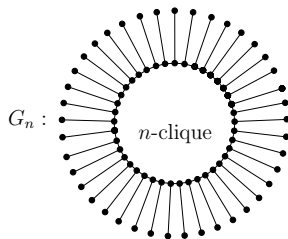
Strong selective amplifiers

- For every $n \geq 1$, we define the “urchin” graph G_n with $2n$ vertices:
 - a **clique** with n vertices
 - an **independent set** with n vertices (called “noses”)
 - a **perfect matching** between them



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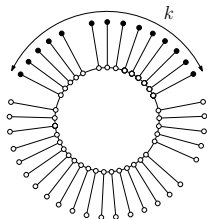
Our result:

Theorem

For every $r > 5$, the fixation probability of a **nose** v of G_n is $f_r(v) \geq 1 - \frac{c(r)}{n}$, where $c(r)$ is a function depending **only on** r .

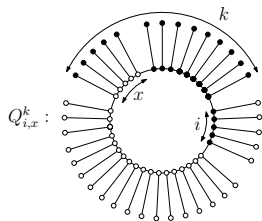
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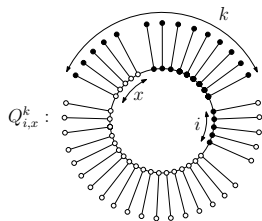


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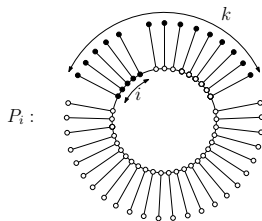
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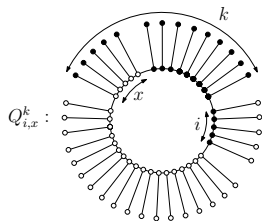
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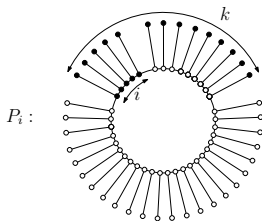
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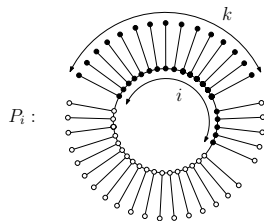


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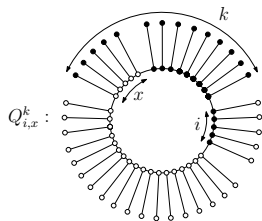
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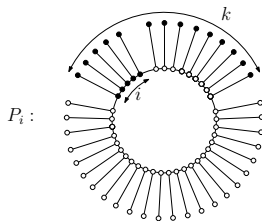
Strong selective amplifiers

- If $i = 0 \Rightarrow Q_{0,x}^k = P_{k-x}^k$

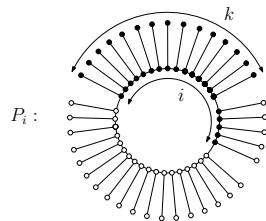


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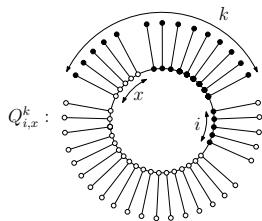
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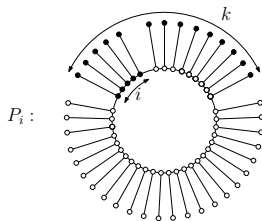
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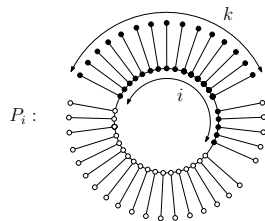


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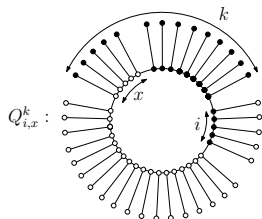
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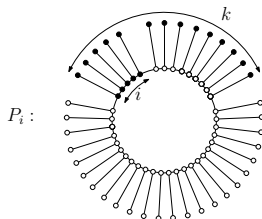
For any i and x , denote by $q_{i,x}^k$ (resp. p_i^k) the probability that:

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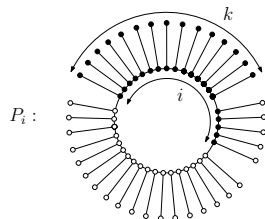


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For all appropriate values of k, i, x : $q_{i,x}^k > p_{k+i-x}^k$.

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\Rightarrow to compute a **lower bound** on the fixation probability $f_r(v)$ of a **nose** v :

- whenever we have **k infected noses** and **i infected clique vertices**,
- we assume that we are at **state P_i^k**
- denote this relaxed Markov chain by \mathcal{M}

Strong selective amplifiers

- We compute a **lower bound** for the fixation probability of state P_0^1

Strong selective amplifiers

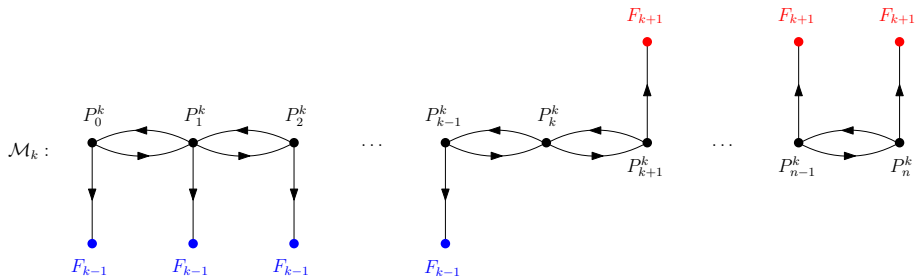
- We compute a **lower bound** for the fixation probability of state P_0^1
- To analyze the Markov chain \mathcal{M} :
 - we **decompose** \mathcal{M} into $n - 1$ Markov chains $\mathcal{M}_1, \mathcal{M}_2, \dots, \mathcal{M}_{n-1}$
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- \mathcal{M}_k has two **absorbing** states:
 - F_{k+1} (arbitrary state with $k + 1$ **infected noses**) \Rightarrow switch to \mathcal{M}_{k+1}
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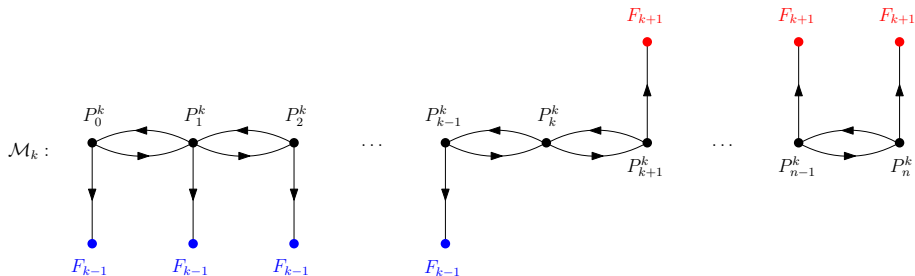
Strong selective amplifiers

Since we need to compute a lower bound of the fixation probability:

- whenever we arrive at state F_{k+1} or state F_{k-1} ,
- we assume that we have the smallest number of infected clique vertices

Therefore:

- $F_{k-1} = P_0^{k-1}$ (no infected clique nodes)



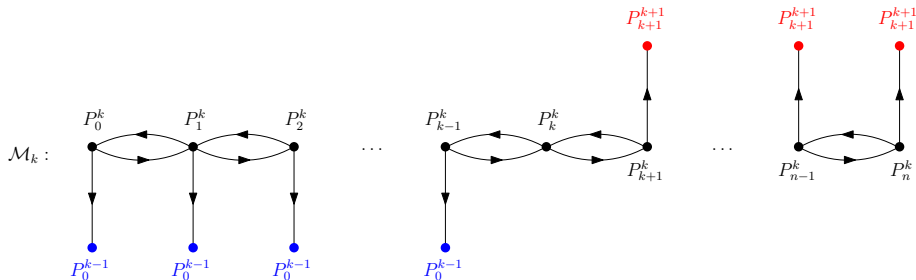
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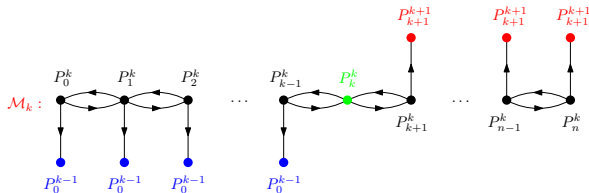
- $F_{k-1} = P_0^{k-1}$ (no infected clique noses)
- $F_{k+1} = P_{k+1}^{k+1}$
(we need at least $k + 1$ infected clique vertices to infect another nose)



Strong selective amplifiers

To analyze the Markov chains \mathcal{M}_k , $k = 1, 2, \dots, n-1$:

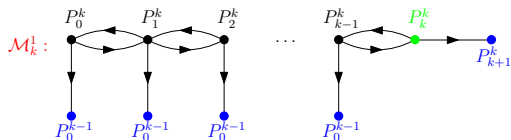
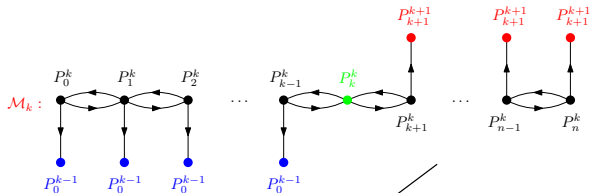
- we **decompose** every \mathcal{M}_k into two Markov chains \mathcal{M}_k^1 and \mathcal{M}_k^2



Strong selective amplifiers

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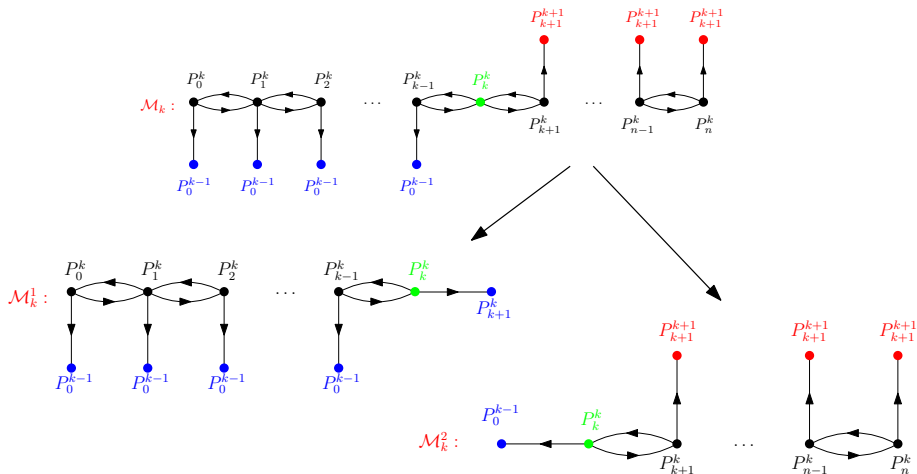
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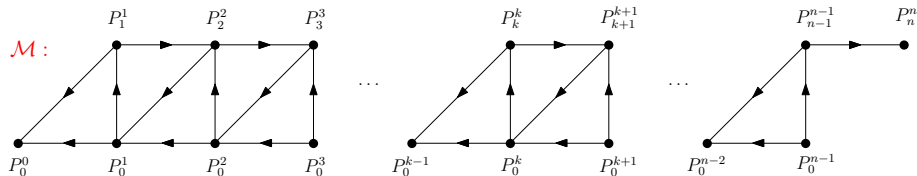
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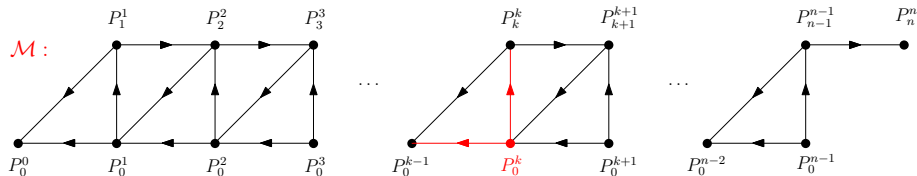
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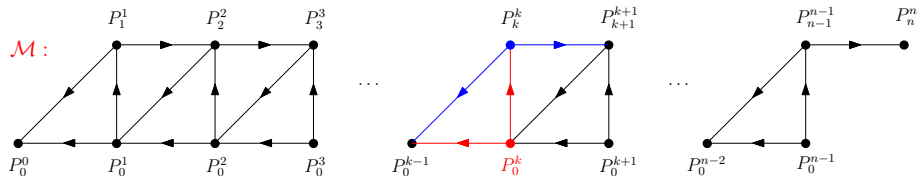
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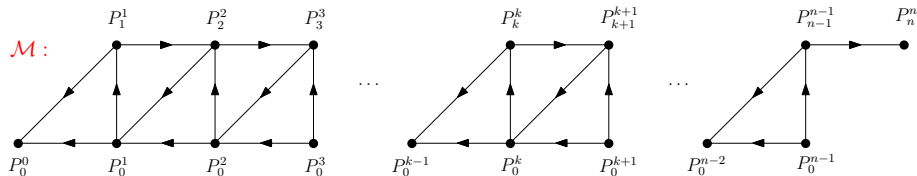


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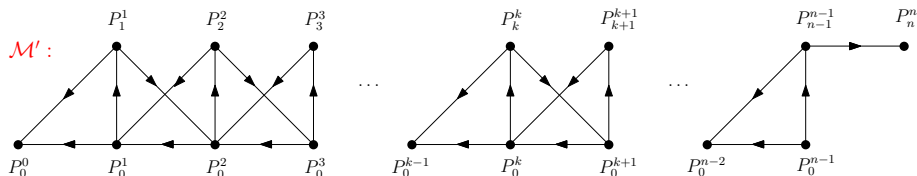
transitions from P_k^k : through the Markov chain \mathcal{M}_2

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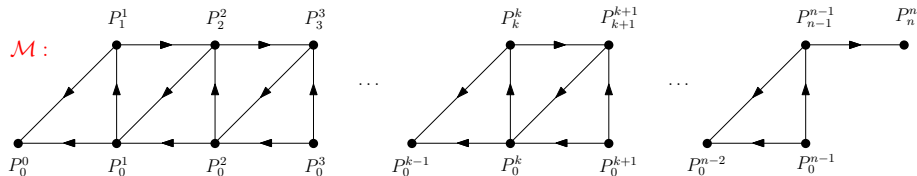


Relax \mathcal{M} further: the **infected** vertices at P_0^k are a **subset** of those at P_k^k

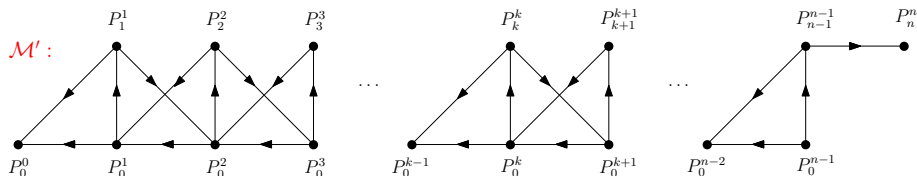


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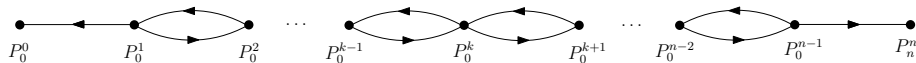
Eliminate from \mathcal{M}' the states $P_k^k \Rightarrow$ a **birth-death** process \mathcal{B}

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In the **birth-death** process \mathcal{B} :

- we can compute a **lower bound** for the probability that, starting at P_0^1 , we arrive at P_n^n before arriving at P_0^0

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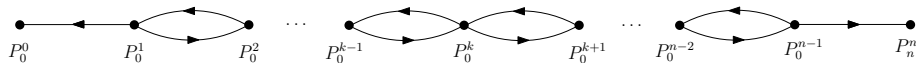


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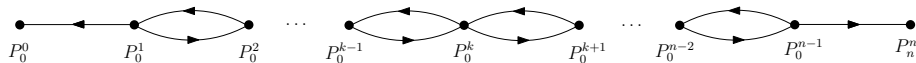


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Using these decompositions, we prove that:

Theorem

For every $r > 5$, the fixation probability of a **nose v** of G_n is $f_r(v) \geq 1 - \frac{c(r)}{n}$, where $c(r)$ is a function depending **only on r**.

\Rightarrow **urchin** graphs are $(\frac{n}{2}, n)$ -amplifiers

The Thermal Theorem

Theorem (Thermal Theorem)

Let $G = (V, E)$ be a connected undirected graph and $r > 1$.

Then $f_r(v) \geq \frac{r-1}{r + \frac{\deg v}{\deg_{\min}}}$ for every $v \in V$.

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- for every $i = 1, 2, \dots, n-1$: **relax** \mathcal{M}_{i-1}^* into the chain \mathcal{M}_i^*
- \mathcal{M}_{n-1}^* provides the desired **lower bound**

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For every vertex subset $S \subseteq V$:

- the fixation probability $f_r(S)$ of S is computed by:

$$f_r(S) = \frac{\sum_{xy \in E, x \in S, y \notin S} \left(r \frac{1}{\deg x} f_r(S + y) + \frac{1}{\deg y} f_r(S - x) \right)}{\sum_{xy \in E, x \in S, y \notin S} \left(\frac{r}{\deg x} + \frac{1}{\deg y} \right)}$$

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For every such edge $xy \in E$ (where $x \in S$ and $y \notin S$):

- x “infects” y with probability proportional to $\frac{1}{\deg x}$
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\Rightarrow for every vertex $v \in V$:

- we define $\frac{1}{\deg v}$ as the **temperature of v**
- a “**hot**” vertex affects more often its neighbors than a “**cold**” vertex

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Furthermore:

- for every set $S \notin \{\emptyset, V\}$ there exists a vertex $x(S) \in S$ and a vertex $y(S) \notin S$ such that $x(S)y(S) \in E$ and:

$$f_r(S) \geq \frac{\left(r \frac{1}{\deg x(S)} f_r(S + y(S)) + \frac{1}{\deg y(S)} f_r(S - x(S)) \right)}{\left(\frac{r}{\deg x(S)} + \frac{1}{\deg y(S)} \right)}$$

The Thermal Theorem

Therefore:

- by replacing all “ \geq ” with “ $=$ ”, we obtain a **lower bound** for all $f_r(S)$

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Definition (the linear system L_0)

Let $G = (V, E)$ be a graph and $r > 1$. Every vertex $v \in V$ has a **weight (temperature)** $d_v > 0$. The **linear system L_0** on the variables $p_r(S)$, where $\emptyset \subset S \subset V$, is:

$$p_r(S) = \frac{r \cdot d_{x(S)} \cdot p_r(S + y(S)) + d_{y(S)} \cdot p_r(S - x(S))}{r \cdot d_{x(S)} + d_{y(S)}}$$

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The **system L_0** defines naturally the **Markov chain \mathcal{M}_0** :

- one state for every vertex subset $S \subseteq V$
- states \emptyset and V are **absorbing**
- every **non-absorbing** state S has exactly two transitions to the states $S + y(S)$ and $S - x(S)$, with transition probabilities

$$q_S = \frac{rd_{x(S)}}{rd_{x(S)} + d_{y(S)}} \text{ and } 1 - q_S, \text{ respectively}$$

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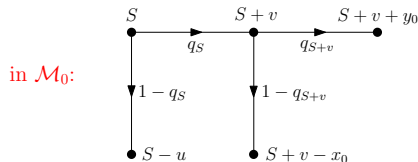
Observation

By setting $d_v = \frac{1}{\deg v}$ for every $v \in V$, it follows that $f_r(S) \geq p_r(S)$ for every set $S \subseteq V$.

The Thermal Theorem

We construct now the chain \mathcal{M}_0^* from the chain \mathcal{M}_0 as follows:

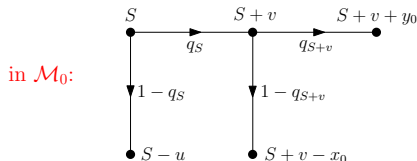
- for every set S in \mathcal{M}_0 :



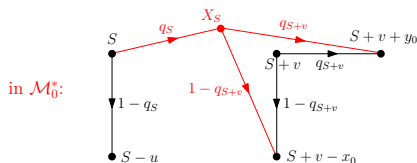
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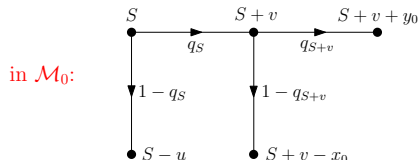
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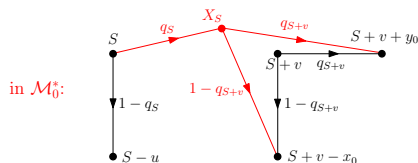
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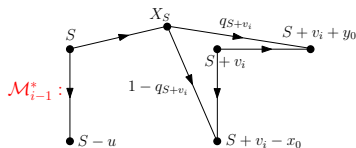
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\Rightarrow All values of $p_r(S)$ in \mathcal{M}_0^* remain the same as in \mathcal{M}_0

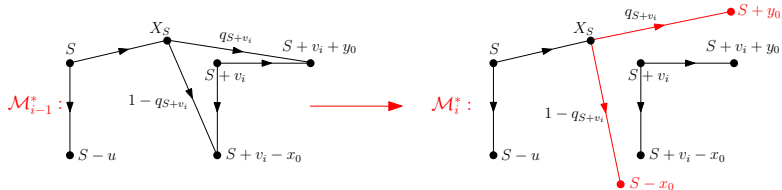
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The Thermal Theorem

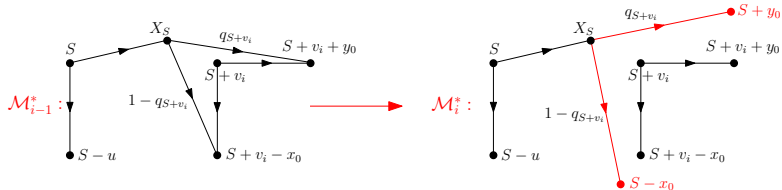
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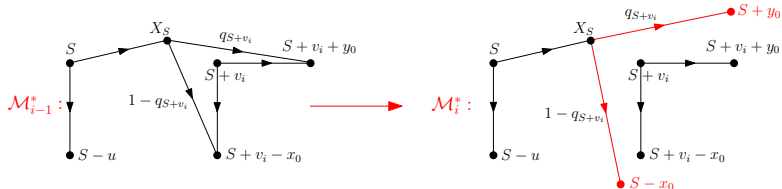


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Lemma

For all these states S , the *forward probability* of S in \mathcal{M}_i^* is a monotone *decreasing* function of the *temperature* d_{v_i} of v_i .

The Thermal Theorem

- We **increase** the **temperature** d_{v_i} in \mathcal{M}_i^* to d_{\max}
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At the end, in the chain \mathcal{M}_{n-1}^* :

- $d_{v_1} = d_{v_2} = \dots = d_{v_{n-1}} = d_{\max} = \frac{1}{\deg_{\min}}$
- $d_{v_0} = \frac{1}{\deg_{v_0}}$
- for every set S , the values of $p_r(S)$ are **not larger** than in \mathcal{M}_0^*

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- for every set S , the values of $p_r(S)$ are **not larger** than in \mathcal{M}_0^*
- We use techniques similar to the **Isothermal Theorem** in [Lieberman et al., *Nature*, 2005] to prove that:

$$f_r(v_0) \geq \frac{(r-1)}{r + \frac{d_{\max}}{d_{v_0}}} = \frac{(r-1)}{r + \frac{\deg v_0}{\deg_{\min}}}$$

- v_0 is chosen arbitrarily \Rightarrow the **Thermal Theorem**

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- We **refined** the notion of **fixation probability** to specific **vertices v**
- We proved:
 - there exist **no strong universal amplifiers**
 - there exist **strong selective amplifiers**
 - there exist **“quite” strong selective suppressors**
 - the **Thermal Theorem** (lower bound)

Summary and open problems

- Do there exist **stronger suppressors** / **amplifiers** of selection?
 - the fixation probability of the strongest known **amplifiers** of natural selection is $1 - \frac{1}{r^2}$ (“star”)
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- More **types** of **mutants** (**many colors**)?

Thank you for your attention!