

# A survey of fitness-based models for biological evolution

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# Introduction

Toy models for time evolution of a system consisting of a population of “species”.

Common features

- ▶ Population is asymptotically large.
- ▶ Fitness-based models:
  - ▶ “At birth” each species is assigned a random “fitness” independent of past.
  - ▶ Time evolution eliminates species with lowest fitness from the system.

**What is the asymptotic fitness distribution ?**

The Models

- ▶ Bak-Sneppen model ('93)
- ▶ A model presented by Guiol Machado and Schinazi ('11)
- ▶ Variations of the above.

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# Bak Sneppen

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One of the first models claimed through numerical simulations to exhibit self-organized criticality.

A discrete time ergodic Markov processes with

- ▶  $N$  species arranged on the vertices of a cycle (or any finite connected graph).
- ▶ Each is assigned an initial fitness, IID  $U[0, 1]$ .
- ▶ Evolution: at each time, the species with lowest fitness and its neighbors are replaced by new species with IID  $U[0, 1]$  fitnesses.

Watch simulation

Simulations suggest

$$\pi_N \xrightarrow{N \rightarrow \infty} \text{IID } U[p_c, 1], \text{ where } p_c \sim 2/3,$$

and  $\pi_N$  is the stationary distribution.

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## Bak Sneppen Avalanches

An **avalanche** from threshold  $p$  is a part of the path from time all fitnesses are  $\geq p$  until next time this happens.

The avalanches provide a natural regenerative structure for the process.

- ▶ Evolution of avalanche depends on the past only through the location of site with lowest fitness when started.
- ▶ As a result, the sequence of durations of avalanches are IID, and so is the number of vertices affected during each avalanche, AKA the range of the avalanche.

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# Bak-Sneppen Avalanche Statistics

## Notation

$D_N(p)$  = Duration of avalanche from threshold  $p$

$R_N(p)$  = Range of avalanche from threshold  $p$

$P_N(p) = P(R_N(p) = N)$

Consider an avalanche from threshold  $p$  on  $\mathbb{Z}$  with initial fitness configuration

$$\dots, 1, 1, \dots, \underset{\substack{\uparrow \\ \text{origin}}}{p}, 1, 1, \dots$$

As before, let

$D_\infty(p)$  = Duration of avalanche

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## Theorem 1 (Meester-Znamenski '04)

$ED_N(p) \rightarrow ED_\infty(p)$ ,  $ER_N(p) \rightarrow ER_\infty(p)$ ,  $P_N(p) \rightarrow P_\infty(p)$ .

- ▶ Asymptotic properties can be studied by considering the infinite system.
- ▶ Main idea: embedding in and coupling of finite system in infinite system.

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# Critical Thresholds

Define

$$p_D = \inf\{p : ED_\infty(p) = \infty\}$$

$$p_R = \inf\{p : ER_\infty(p) = \infty\}$$

$$p_P = \inf\{p : P_\infty(p) > 0\}$$

Theorem 2 (Meester-Znamenski '03, Meester-Znamenski '04)

1.  $0 < p_D = p_R \leq p_P < 1 - e^{-68}$ .
2. If  $p_R = p_P$ , then  $\pi_N \xrightarrow{N \rightarrow \infty} \text{IID } U[p_P, 1]$ .

Letting  $F$  be the fitness at some distinguished site 0, then

Proposition 1

1.  $\pi_N(F \leq p_D) \rightarrow 0$ .
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This was not stated in the paper, but follows from the proofs.

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## Bak Sneppen, a little more

### Proposition 2 (B. WIP)

Let

$$\rho = \inf_{p,d} \sum_{k=1}^{\infty} \frac{1}{d_k p_k},$$

where the infimum is taken over all probability distributions  $(p_1, p_2, \dots)$  on  $\mathbb{N}$  and all-integer nondecreasing valued sequences  $(d_k)_{k \in \mathbb{N}}$  with the growth constraint  $d_1 = 1, d_{k+1} < 2d_k$  for  $k > 1$ . Then

$$p_P \leq 1 - e^{-\rho},$$

- ▶ Simulations give  $\rho < 11.3$ .
- ▶ We need to get to  $-\ln \frac{1}{3} = 1.09861228867$ .

### Theorem 3 (B. WIP)

If  $P(R_{\infty}(p) > r) \geq cr^{-\alpha}$  for some  $\alpha < 1$ , then  $p_P \leq p$ .

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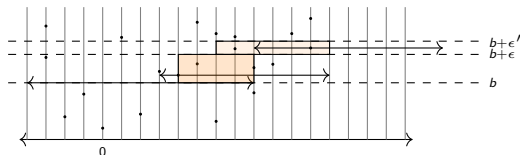
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## Bak Sneppen – on proofs

- ▶ Tool: Graphical representation of avalanches on  $\mathbb{Z}$ , due to Meester and his coauthors.
- ▶ Switch from uniform fitnesses to  $\text{Exp}(1)$ . This allows for Poisson process techniques.

**At end of avalanche from threshold  $b$ , fitness of sites in its range IID  $b + \text{Exp}(1)$ .**

- ▶ To each site attach a rate-1 Poisson process, processes are independent.
- ▶ Suppose the avalanche from threshold  $b$  starting from the origin has the range given by the arrow.
- ▶ Fitness distribution of sites in range coincides with the first arrivals of the Poisson processes above  $b$ .
- ▶ The range of avalanche from threshold  $b + \epsilon$  will be at least  $\frac{3}{2} \times R_b$ , if at least one of the avalanches in the orange region extends to the right at least as  $R_b$  did.
- ▶ Allows to approach through thinning of a Poisson Point Process.
- ▶ For large enough  $b$ , one can show that exists an infinite cascade of such avalanches below fitness  $b + \epsilon$ .





# Local Bak-Sneppen

Joint with R.C. Silva

## Two Geometries

### What would be a “proper” tractable analog for Bak-Sneppen ?

The difficulty in the Bak-Sneppen model stems from the following:

- ▶ Use **complete graph** geometry to locate the global minimum.
- ▶ Use “**nearest neighbor**” geometry to determine at what vertices species will be replaced.

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Consider a finite, connected (undirected) graph  $G = (V, E)$ .

## Initially

- ▶ Assign IID  $U[0, 1]$  fitnesses to each  $v \in V$ .
- ▶ Set  $X_0$  as the vertex with lowest fitness.

## Time evolution

- ▶ Given  $X_n$ , set  $X_{n+1}$  to be the vertex with minimal fitness among  $u \sim X_n$  and  $X_n$  itself.
- ▶ Set fitness of all elements in neighborhood of  $X_{n+1}$  as IID  $U[0, 1]$ , independent of past.

## Observe

- ▶ Markov chain on state space = product of  $V$  and  $[0, 1]$ -valued functions on  $V$ .
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What can we say about this new process ?



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# Stationary Distribution for Local Bak-Sneppen

## Notation

- ▶ For  $v \in V$ ,  $A_v = \{u \in V : \{u, v\} \in E \text{ or } u = v\}$ .
- ▶ **Random walk on  $G$** : from  $v \in V$  move to uniformly sampled  $u \in A_v$ .
- ▶ Stationary distribution:  $\mu(v) = \frac{|A_v|}{\sum_{u \in V} |A_u|}$ .
- ▶ If  $U_1, \dots, U_n$  are IID  $U[0, 1]$ , then set  $U(n, [0, 1])$  as the distribution

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## Theorem 4 (Silva-B.)

- ▶ Let  $(Z_n^u : n \in \mathbb{Z}_+)$  be independent random walks on  $G$  with  $Z_0^u = u$ .
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$$\tau_v = \inf\{n \in \mathbb{Z}_+ : Z_n^X \in A_v\}, \text{ and } V_i = \{v \in V : \tau_v = i\}.$$

Given  $V_0, V_1, \dots$ , assign fitnesses at each  $v \in V$ , which are independent and are

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# Local Bak-Sneppen for Regular Graphs

## Corollary 1

*If  $G$  is  $d$ -regular, then the stationary distribution for the local Bak-Sneppen satisfies:*

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Now send size to infinity

## Corollary 2

*Suppose that  $(V_n : n \in \mathbb{N})$  is an increasing sequence of finite sets. For each  $n$ , let  $G_n = (V_n, E_n)$  be a  $d$ -regular connected graph.*

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Guiol-Machado-Schinazi

## Why ?

- ▶ Have population growth be part of model, not external parameter.
- ▶ Tractability.

## Construction

- ▶ The population size is a reflected random walk on  $\mathbb{Z}_+$  (that is random walk minus its running minimum).
- ▶ When population increases, AKA birth (possibly multiple), new individuals are assigned IID  $U[0, 1]$  fitnesses.
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## What is the asymptotic fitness distribution ?

More precisely, letting  $\hat{F}_n(f)$  denote the empirical fitness distribution

$$\hat{F}_n(f) = \begin{cases} \text{prop. with fitness} \leq f & \text{if pop. size is} > 0 \\ \text{CDF of } \delta_0 & \text{otherwise.} \end{cases}$$

Understand limit (LLN) and fluctuations (CLT) of  $\hat{F}_n$  as  $n \rightarrow \infty$ .

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More precisely, letting  $\hat{F}_n(f)$  denote the empirical fitness distribution

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# GMS Law of Large Numbers

## Notation.

- ▶  $I \stackrel{\text{dist}}{=} \text{Incement of random walk.}$
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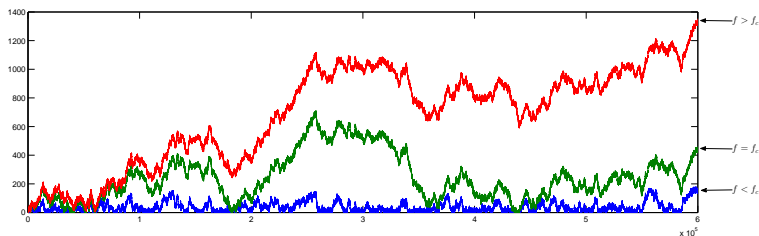
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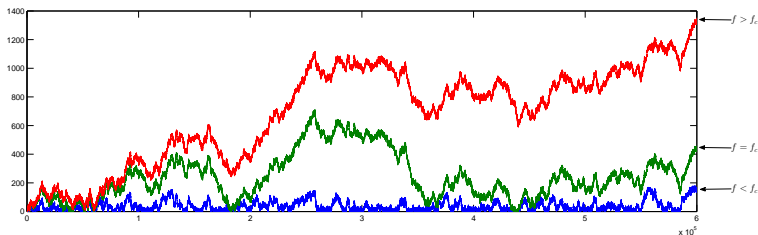
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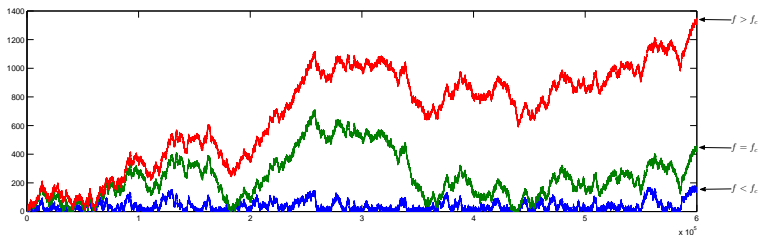
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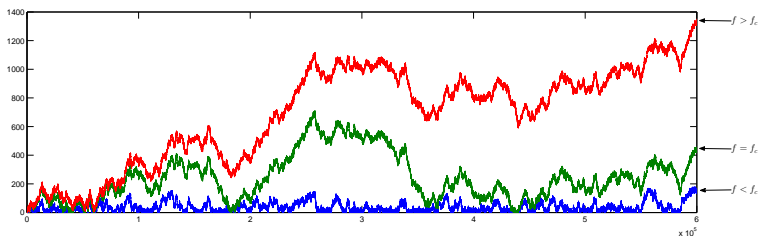
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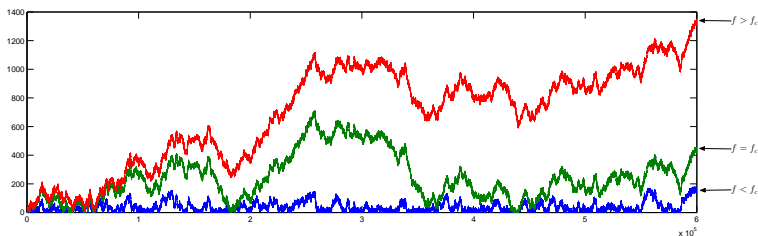
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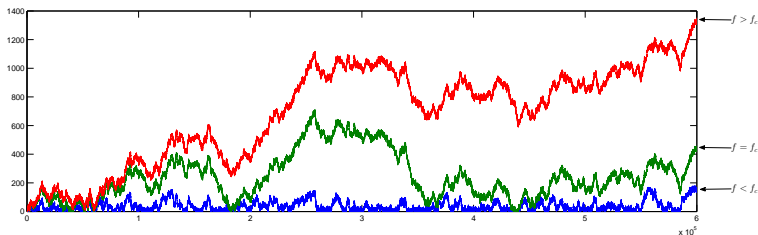
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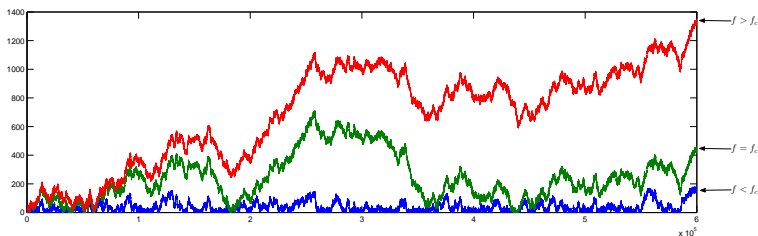
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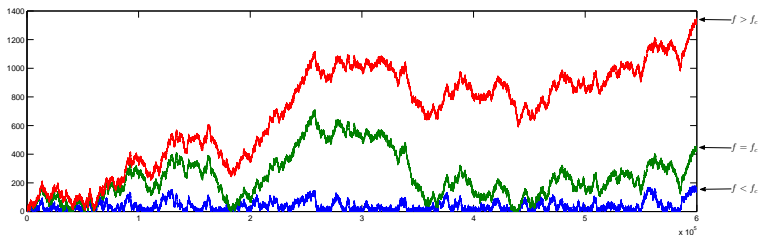
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# GMS Central Limit Theorem

Let

$$\hat{\Delta}_n = \hat{F}_n - F_\infty.$$

Then we know that  $\hat{\Delta}_n$  converges to 0, uniformly, a.s.

Next, we look at fluctuations.

Assumption

$$E(I^2) < \infty.$$

Processes appearing in limit

- ▶  $W_1$  standard BM, and the corresponding bridge  $Br_1$ :

$$Br_1(f) := W_1(f) - fW_1(1).$$

- ▶ If  $f_c = 0$ , choose  $\widetilde{W}_1 \equiv 0$ .
- ▶ If  $f_c > 0$ :  $\widetilde{W}_1$  standard BM derived from  $W_1$  as follows:

- ▶  $U \sim U[f_c, 1]$ , independent of  $W_1$ .
- ▶ An "interval"  $\widetilde{A}_t$  of length  $(1 - f_c)t$ , shifted by  $U$ .
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$$\widetilde{W}_1(t) := \frac{1}{\sqrt{f_c(1 - f_c)}} \left( (1 - f_c)W_1(f_c t) + f_c \int \mathbf{1}_{\widetilde{A}_t}(s) dW_1(s) \right).$$



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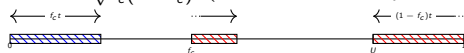
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For a path  $\omega \in D[0, 1]$ , let  $\Psi(\omega) := \omega(1) - \inf_{0 \leq t \leq 1} \omega(t)$

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$$\sqrt{n} \begin{pmatrix} \widehat{\Delta}_n(\cdot)|_{(f_c, 1]} \\ \widehat{\Delta}_n(f_c) \end{pmatrix} \Rightarrow \frac{1}{EI_+} \begin{pmatrix} \overbrace{\sigma_1 B r_1 + \sigma_2 W_2(1)(1 - F_\infty)}^{\text{Gaussian process}} \\ \underbrace{\Psi(\tilde{\sigma}_1 \tilde{W}_1 + \sigma_2 W_2)}_{\text{Positive RV}} \end{pmatrix},$$

with  $\sigma_1 = \sqrt{EI_+}$ ,  $\tilde{\sigma}_1 = \sqrt{f_c(1 - f_c)EI_+}$ ,  $\sigma_2 = \sqrt{f_c^2 E(I_+^2) + E(I_-^2)}$ , and the convergence is  $D(f_c, 1] \times \mathbb{R}$ .

Marginals

$$\sqrt{n} \widehat{\Delta}_n(f) \Rightarrow \begin{cases} \sigma(f \wedge 1) N(0, 1) & f > f_c; \\ \sigma(f_c) |N(0, 1)| & f = f_c; \\ 0 & f < f_c, \end{cases}$$

$$\text{where } \sigma(f) := \frac{1}{E(I_+)} \sqrt{f(1 - f)E(I_+) + \left(\frac{1 - f}{1 - f_c}\right)^2 (f_c^2 E(I_+^2) + E(I_-^2))}$$

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with  $\sigma_1 = \sqrt{EI_+}$ ,  $\tilde{\sigma}_1 = \sqrt{f_c(1 - f_c)EI_+}$ ,  $\sigma_2 = \sqrt{f_c^2 E(I_+^2) + E(I_-^2)}$ , and the convergence is  $D(f_c, 1] \times \mathbb{R}$ .

Marginals

$$\sqrt{n} \widehat{\Delta}_n(f) \Rightarrow \begin{cases} \sigma(f \wedge 1) N(0, 1) & f > f_c; \\ \sigma(f_c) |N(0, 1)| & f = f_c; \\ 0 & f < f_c, \end{cases}$$

where  $\sigma(f) := \frac{1}{E(I_+)} \sqrt{f(1 - f)E(I_+) + \left(\frac{1 - f}{1 - f_c}\right)^2 (f_c^2 E(I_+^2) + E(I_-^2))}$

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# GMS with selection

Assume

$$P(I = 1) = p = 1 - P(I = -1).$$

New feature

- ▶ At birth the individual obtains
  - ▶ w/prob  $r$  new  $U[0, 1]$  fitness.
  - ▶ w/prob  $1 - r$ , an existing fitness, uniformly among existing fitnesses, or new one if population is zero.
- ▶ At death, eliminate all species with lowest fitness.

We refer to the population with fixed fitness as a **site**.

Observation

- ▶ Probability of new site is  $pr$ .
- ▶ Probability of eliminating a site is  $1 - p$ .

Conclusion

1. Number of sites coincides with GMS with  $P(I = 1) = pr$ ,  $P(I = -1) = 1 - p$  and  $P(I = 0) = 1 - pr - (1 - p)$ .
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# GMS w/Selection

What is site size distribution ?

Let  $\hat{H}_n$  denote the empirical distribution of sites and their respective fitness:

$$\hat{H}_n(A \times B) = \frac{\# \text{ sites whose size is in } A \text{ and whose fitness is in } B}{\# \text{ sites}}.$$

Theorem 7 (Schinazi-B. '15)

$$\hat{H}_n \rightarrow \text{Geom} \left( \frac{pr - (1 - p)}{p - (1 - p)} \right) \otimes U[f_c, 1], \text{ a.s.}$$

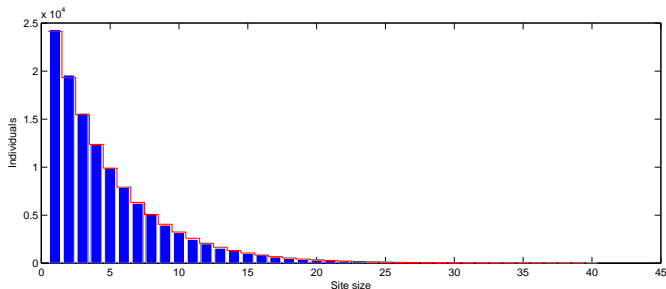


Figure: Empirical dist of site sizes ( $p = 0.8, r = 0.4, n = 10^6$ ) and corresponding Geom.

# GMS w/selection

## Why Geometric ?

Fix site size  $k > 1$ . Consider number of sites of size  $k$  with fitness  $> f_c$ .

Assume the proportion of such sites converges to  $H_\infty(k)$ .

- ▶ Number of such sites grows at speed

$$p(1-r)(H_\infty(k-1) - H_\infty(k)) + o(1) = H_\infty(k) * (pr - (1-p))$$

Then change in the number of sites of size  $k$  occurs only at

- ▶ At birth
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This equation guarantees geometric decay.

## The problem

Proving that the assumption actually holds.

- ▶ Easy calculus exercise if  $pr > \frac{1}{2}$ .
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Assume the proportion of such sites converges to  $H_\infty(k)$ .

- ▶ Number of such sites grows at speed

$$p(1-r)(H_\infty(k-1) - H_\infty(k)) + o(1) = H_\infty(k) * (pr - (1-p))$$

Then change in the number of sites of size  $k$  occurs only at

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**This equation** guarantees geometric decay.

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Proving that the assumption actually holds.

- ▶ Easy calculus exercise if  $pr > \frac{1}{2}$ .
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# GMS w/selection

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*Thank you.*

Ad: Markov chains REU at UConn this summer.

Details on our [mathprograms.org](https://mathprograms.org) page or on [markov-chains-reu.math.uconn.edu](https://markov-chains-reu.math.uconn.edu)