

# Stochastic Models for Large Interacting Systems in the Sciences

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*Interacting Particle Systems*, Springer, 1985

*Stochastic Interacting Systems: Contact, Voter and Exclusion Processes*, Springer, 1999

## (Biased) Voter Models

M. Kimura, "*Stepping stone*" model of population, Ann. Rep. Nat. Inst. Genetics Japan **3** (1953).

T. Williams and R. Bjercknes, *Stochastic model for abnormal clone spread through epithelial basal layer*, Nature **236** (1972).

## Stochastic Ising Models

R. J. Glauber, *Time-dependent statistics of the Ising model*, J. Mathematical Physics **4** (1963).

J. M. Keith, *Segmenting eukaryotic genomes with the generalized Gibbs sampler*, J. Computational Biology **13** (2006).

**Kimura** won the 1992 Darwin Medal of the Royal Society;  
**Glauber** won the 2005 Nobel Prize in Physics.

## Contact Processes

P. Grassberger and A. de la Torre, *Reggeon field Theory (Schlögl's first model) on a lattice: Monte Carlo calculations of critical behaviour*, Annals of Physics **122** (1979).

R. Schinazi, *On an interacting particle system modeling an epidemic*, J. Mathematical Biology **34** (1996).

## Exclusion Processes

C. T. MacDonald, J. H. Gibbs and A. C. Pipkin, *Kinetics of biopolymerization on nucleic acid templates*, Biopolymers **6** (1968).

H-W. Lee, V. Popkov and D. Kim, *Two-way traffic flow: Exactly solvable model of traffic jam*, J. Physics A **30** (1997).

## The Voter Model

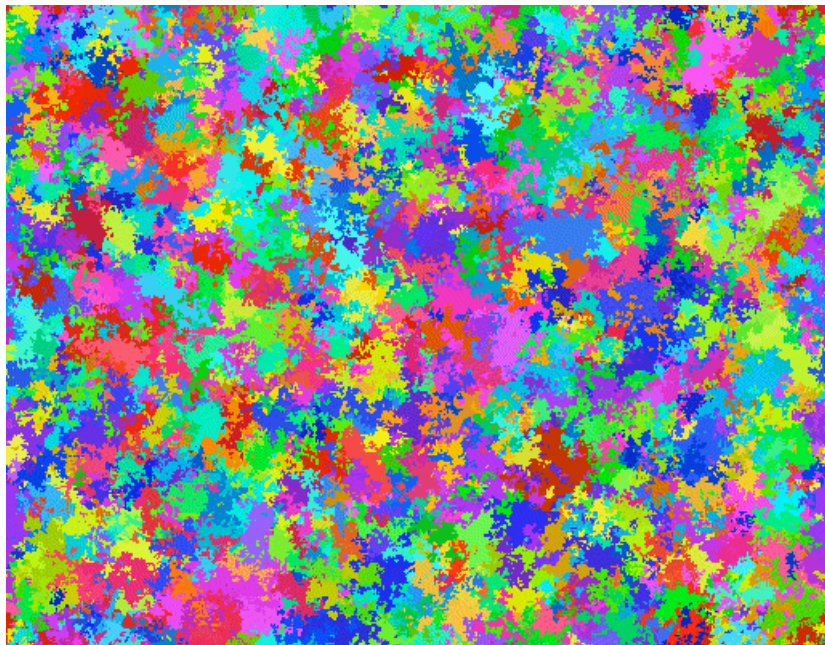
(Holley and Liggett, 1975)

State of the system at any given time: An infinite array of individuals that change opinions at random times:

|          |          |          |          |          |          |          |          |
|----------|----------|----------|----------|----------|----------|----------|----------|
| <i>R</i> | <i>D</i> | <i>R</i> | <i>R</i> | <i>R</i> | <i>D</i> | <i>D</i> | <i>D</i> |
| <i>R</i> | <i>D</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>R</i> |
| <i>R</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>D</i> | <i>D</i> |
| <i>R</i> | <i>R</i> | <i>D</i> | <i>R</i> | <i>R</i> | <i>R</i> | <i>D</i> | <i>R</i> |
| <i>D</i> | <i>D</i> | <i>R</i> | <i>R</i> | <i>R</i> | <i>D</i> | <i>D</i> | <i>R</i> |
| <i>R</i> | <i>R</i> | <i>R</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>D</i> | <i>R</i> |
| <i>R</i> | <i>D</i> | <i>D</i> | <i>D</i> | <i>R</i> | <i>D</i> | <i>D</i> | <i>R</i> |
| <i>D</i> | <i>D</i> | <i>R</i> | <i>R</i> | <i>R</i> | <i>D</i> | <i>R</i> | <i>D</i> |

At exponential times, individuals choose a neighbor at random and adopt his/her opinion.

A simulation with many opinions:



Back to two opinions  $R$  and  $D$ . Each individual flips a fair coin to determine his/her initial opinion.

**Questions:**

(a) What is  $P(\text{individual } x \text{ has opinion } R \text{ at time } t)$ ?

(b) What can be said about

$P(\text{individuals } x \text{ and } y \text{ have the same opinion at time } t)$

for large times  $t$ ?

**Answers:**

(a)  $P(\text{individual } x \text{ has opinion } R \text{ at time } t) = \frac{1}{2}$  for all  $x, t$ .

For question (b), consider the model on a  $d$ -dimensional array.

If  $d = 1$  or  $d = 2$ ,

$$\lim_{t \rightarrow \infty} P(\text{individuals } x \text{ and } y \text{ have the same opinion at time } t) = 1.$$

However, if  $d \geq 3$ , this limit, call it  $g(x, y)$ , satisfies

$$\frac{1}{2} < g(x, y) < 1 \text{ for } x \neq y$$

and

$$\lim_{y-x \rightarrow \infty} g(x, y) = \frac{1}{2}.$$

In fact,

$$g(x, y) = \frac{1}{2} + \frac{1}{2}P(\text{a random walk starting at } y - x \text{ ever hits } 0).$$

## The Biased Voter Model, or The Williams-Bjerknes Tumor Growth Model (1972)

Differences:

(a) It is more likely that health cells become cancerous than the other way around.

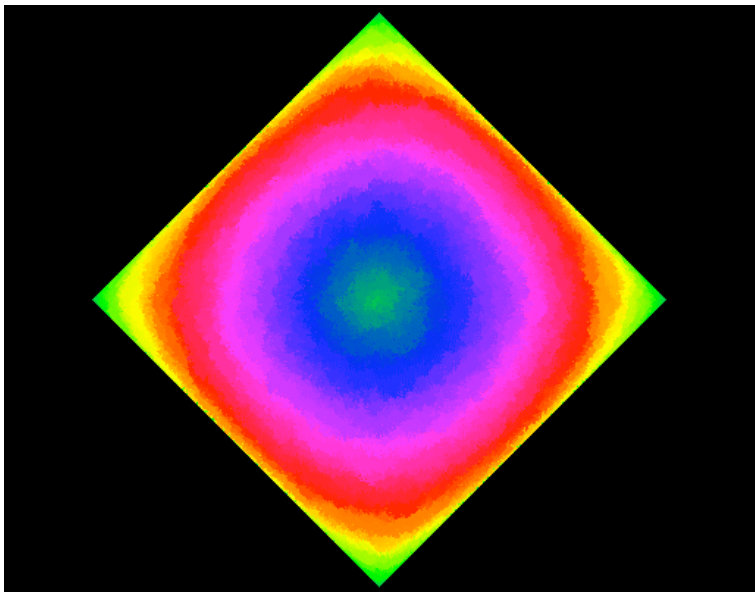
(b) Start with a single cancerous cell. Given that the tumor does not disappear, it has an asymptotic shape.

Here is a simpler discrete time model:

### Richardson's Model (1973)

Cancerous cells remain cancerous. Healthy cells become cancerous at time  $n + 1$  with probability  $p$  if at least one neighbor was cancerous at time  $n$ .





**Question:** What is the asymptotic shape?

**One answer:** It has a straight edge if  $p > p_c$ , where  $p_c$  is the critical value of a discrete time contact process. (Durrett and Liggett, 1981)

Nonrigorously,  $p_c \sim .64$ . Rigorously,  $p_c \leq \frac{2}{3}$ .

**Question:** What is the limiting asymptotic shape as  $p \downarrow 0$ ?

Richardson speculated, based on simulations, that the limiting shape is a circle. Kesten proved that in high dimensions, it is not a sphere.

**Question:** Why consider infinite systems, when real systems are finite?

**Answer:** Infinite models at infinite times are better models for large finite systems at large finite times than are finite models at infinite times.

## The Contact Process

(Harris, 1974)

- (a) Infected individuals become healthy at rate 1.
- (b) Healthy individuals become infected at rate

$$\lambda \times \#(\text{infected neighbors}).$$

On a **finite set**, the infection dies out eventually for any  $\lambda$ .

On a  $d$ -dimensional grid, there is a critical value  $\lambda_d$ ,

$$\frac{1}{2d} \leq \lambda_d \leq \frac{2}{d},$$

so that

(a) the infection **dies out** eventually if  $\lambda \leq \lambda_d$ ,

and

(b) **survives** with positive probability if  $\lambda > \lambda_d$ .

On an  $N \times N \times \cdots \times N$  grid, the extinction time  $\tau_N$  satisfies:

$$\tau_N \sim \log N \quad \text{if} \quad \lambda < \lambda_d$$

and

$$\log \tau_N \sim N^d \quad \text{if} \quad \lambda > \lambda_d.$$

$$(\log 1000 \sim 7; \log t = 1000 \Rightarrow t \sim 10^{434}.)$$

## Exclusion Processes

(Spitzer, 1970)

From a 2009 paper by N. Bogoliubov (Steklov Mathematical Institute, St. Petersburg, Russia):

“The totally asymmetric simple exclusion process (TASEP) ... is connected with the ‘crystalline limit’ of the XXZ R-matrix. It is one of the most studied models of low dimensional non-equilibrium physics.”

In one dimension, the state of the system is:

... 1 1 0 1 0 1 1 1 0 ...

At random (exponential) times, particles try to move to the right with probability  $p$  and to the left with probability  $q = 1 - p$ . Moves to occupied sites are not allowed.

Suppose  $p = 1$  and the initial configuration is

$\dots 1 1 1 1 1 0 0 0 0 0 \dots$

**Question:** What is the limiting distribution as  $t \rightarrow \infty$ ?

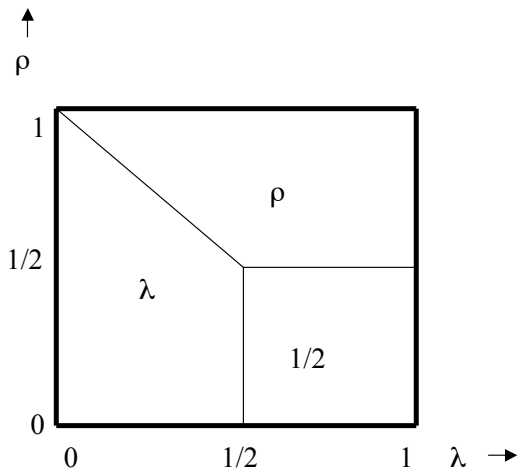
**Answer:** Independent fair coin tossing, i.e.,

$$\lim_{t \rightarrow \infty} P(1 \text{ at time } t) = \frac{1}{2}, \quad \lim_{t \rightarrow \infty} P(11 \text{ at time } t) = \frac{1}{4}, \dots$$

**Question:** What if  $p$  is general, and the initial distribution is

$\dots \lambda \lambda \lambda \lambda \lambda \rho \rho \rho \rho \rho \dots?$

**Answer:** If  $p = \frac{1}{2}$ , the limiting distribution is independent coin tossing with parameter  $\frac{\lambda + \rho}{2}$ . If  $p > \frac{1}{2}$ , the parameter is



## The PDE Connection

Rescale space and time, pretending that the states of sites are independent. The result, letting

$u(t, x) =$  the probability that site  $x$  is occupied at time  $t$ :

1. If  $p = \frac{1}{2}$ , the heat equation,

$$\frac{\partial u}{\partial t} = \frac{1}{2} \frac{\partial^2 u}{\partial x^2}.$$

2. If  $p \neq \frac{1}{2}$ , Burger's equation,

$$\frac{\partial u}{\partial t} + (p - q) \frac{\partial}{\partial x} [u(1 - u)] = 0.$$



## Rates of Convergence for Symmetric Markov Chains

(Key words: Gibbs sampler; Markov Chain Monte Carlo — (MC)<sup>2</sup>)

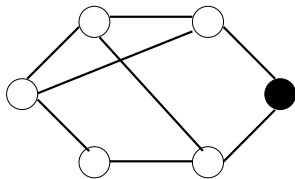
Consider a finite graph, with nonnegative weights  $c_e$  attached to the edges.

Labels are places on the vertices.

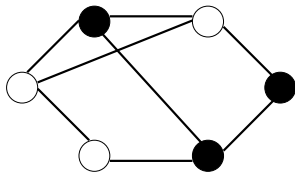
At exponential times of rate  $c_e$ , interchange the labels at the vertices joined by  $e$ .

Depending on the nature of the labels, various Markov chains can be defined:

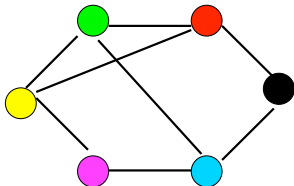
One particle MC



Symmetric exclusion



MC on permutations



For the Markov chain on permutations, think in terms of

## Card Shuffling

**Vertices = positions in a deck**

**labels = cards**

|    |                  |
|----|------------------|
| 1  | Three of Clubs   |
| 2  | Two of Diamonds  |
| ⋮  | ⋮                |
| i  | Five of Diamonds |
| ⋮  | ⋮                |
| j  | Ten of Spades    |
| ⋮  | ⋮                |
| 52 | Jack of Diamonds |

At rate  $c_{i,j}$ , interchange **Five of Diamonds** and **Ten of Spades**.

Note:  $52! \sim 10^{68} \gggggg 52$ .

For a finite state Markov chain  $X(t)$ , the transition probabilities are

$$p_t(i, j) = P(X(t) = j \mid X(0) = i).$$

If  $P(t)$  is the matrix with entries  $p_t(i, j)$ , then

$$P(t) = e^{Qt}.$$

The smallest positive eigenvalue of  $\lambda$  of  $-Q$  determines the rate of convergence to the equilibrium  $\pi$ :

$$p_t(i, j) \sim \pi(j) + a(i, j)e^{-\lambda t}, \quad t \uparrow \infty.$$

Let  $\lambda$  be this eigenvalue for the one particle process, and  $\lambda^*$  be this eigenvalue for the process on permutations. Then

$$\lambda \geq \lambda^*.$$

**Aldous' Conjecture** (1992):  $\lambda^* = \lambda$ .

### Why guess this?

1. True for  $c_e \equiv 1$  on complete graph – Diaconis and Shahshahani (1981).
2. True for  $c_e \equiv 1$  on star graphs – Flatto, Odlyzko and Wales (1985).

### Why do we care?

1. It is MUCH easier to compute the eigenvalues of an  $n \times n$  matrix than of an  $n! \times n!$  matrix.
2. The main eigenvalue for the symmetric exclusion process lies between  $\lambda$  and  $\lambda^*$ , so it would follow that it agrees with the common value of  $\lambda$  and  $\lambda^*$ .

## Recent supporting results:

1. True for **general**  $c_e$  on trees – Handjani and Jungreis (1996).
2. Other related results by Koma and Nachtergele (1997), Morris (2008), Starr and Conomos (2008), Cesi (2009), and Dieker (2009).

**Theorem.** (Caputo, Liggett and Richthammer, 2009)

On a **general** finite graph with **arbitrary** rates,

$$\lambda^* = \lambda.$$