I Spatial Models in Population Biology

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Lecture 1: Interacting Particle Systems

Jan M. Swart Spatial Models in Population Biology

Outline

- Poisson construction of continuous-time Markov chains
- Poisson construction of interacting particle systems
- Unique ergodicity
- Examples

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Continuous-time Markov chains

Let S be a countable set. A probability kernel on S is a function $K: S^2 \rightarrow [0,1]$ such that $\sum_y K(x,y) = 1$. We calculate with kernels as with matrices:

$$KL(x,z) := \sum_{y} K(x,y)L(y,z)$$
 and $Kf(x) := \sum_{y} K(x,y)f(y).$

Let $X = (X_t)_{t\geq 0}$ be a stochastic process taking values in a countable set S. We assume that the sample paths of X are *piecewise constant* and *right-continuous*. By definition, X is a (time-homogenous) *Markov process* if

$$\mathbb{P}\big[X_u \in \cdot \ \big| \ (X_s)_{0 \leq s \leq t}\big] = P_{u-t}(X_t, \cdot) \quad \text{a.s.} \qquad (0 \leq t \leq u),$$

where the *transition kernels* $(P_t)_{t\geq 0}$ form a collection of probability kernels on S such

$$P_sP_t = P_{s+t}$$
 and $\lim_{t\downarrow 0} P_t = P_0 = 1.$

We would like to define $(P_t)_{t\geq 0}$ in terms of its generator

$$G(x,y) := \lim_{t \downarrow 0} t^{-1} \big(P_t(x,y) - 1(x,y) \big),$$

where $1(x, y) := 1_{\{x=y\}}$ denotes the identity matrix. Such a generator must satisfy

$$G(x,y) \ge 0$$
 $(x \ne y)$ and $\sum_{y} G(x,y) = 0.$

We interpret G(x, y) $(x \neq y)$ as the *rate* of transitions $x \mapsto y$. We should have

$$\mathbb{P}^x[X_t=y]=P_t(x,y)=1(x,y)+tG(x,y)+O(t^2)$$
 as $t\downarrow 0,$

where \mathbb{P}^{x} denotes the law of the process started in $X_{0} = x$.

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If S is finite, then we can define

$$P_t = e^{tG} := \sum_{n=0}^{\infty} \frac{1}{n!} t^n G^n \qquad (t \ge 0).$$

This works more generally if G is bounded in the sense that

$$\sup_{x}\sum_{y:\,y\neq x}G(x,y)<\infty.$$

Generator construction of continuous-time Markov chains

Let G be a Markov generator on a countable set S. Then, for each $z \in S$, there exists a unique *minimal* solution to the *backward* equations

$$\frac{\partial}{\partial t}P_t(x,z) = \sum_y G(x,y)P_t(y,z) \qquad (t \ge 0, x \in S).$$

Moreover, $(P_t)_{t\geq 0}$ is a semigroup of subprobability kernels. For each initial state $X_0 = x \in S$, there exists a process $X = (X_t)_{t\geq 0}$ with values in $S \cup \{\infty\}$ such that

$$\begin{split} X_t &= \infty \quad \forall t \geq \tau := \inf\{t \geq 0 : X_t = \infty\},\\ \lim_{t \uparrow \tau} X_t &= \infty \quad \text{if} \quad \tau < \infty,\\ \mathbb{P}^x \Big[X_u \in y \, \Big| \, (X_s)_{0 \leq s \leq t} \Big] &= P_{u-t}(X_t, y) \quad \text{a.s.} \quad (y \in S, \ t \leq u). \end{split}$$

If $\mathbb{P}^{x}[\tau = \infty] = 1$ for all $x \in S$, then X is *nonexplosive*.

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Example: a finite contact process

Let (Λ, E) be a finite directed graph with vertex set Λ and set of directed edges $E \subset \Lambda \times \Lambda$. Let $S = \{0, 1\}^{\Lambda}$ be the space of all functions $x : \Lambda \to \{0, 1\}$. We interpret $x = (x(i))_{i \in \Lambda}$ as a particle configuration where for $i \in \Lambda$,

x(i) = 0 means the site *i* is empty, x(i) = 1 means there is a particle at *i*.

The contact process with infection rate $\lambda \ge 0$ is a continuous-time Markov chain $X = (X_t)_{t\ge 0}$ with state space S. In each jump, the number of occupied sites can increase or decrease by one. Let $x \in S$ and $j \in \Lambda$ such that x(j) = 0. Then

$$G(x,x+\delta_j):=\lambda\sum_{(i,j)\in E}x(i) \quad ext{and} \quad G(x+\delta_j,x):=1.$$

We set G(x, y) := 0 if x and y differ in more than one site and choose G(x, x) such that $\sum_{y} G(x, y) = 0$.

Example: a finite contact process

For each $(i,j) \in E$, define a branching map $\mathtt{bra}_{ij}: S o S$ by

$$ext{bra}_{ij}x(k) := \left\{ egin{array}{cc} x(i) \lor x(j) & ext{if } k=j, \ x(k) & ext{otherwise.} \end{array}
ight.$$

For each $i \in \Lambda$, define a *death map* death_i : $S \rightarrow S$ by

$$death_i x(k) := \begin{cases} 0 & \text{if } k = i, \\ x(k) & \text{otherwise} \end{cases}$$

Fix x. Assume that X_t satisfies for all $(i,j) \in E$ resp. $i \in \Lambda$,

$$X_t = \left\{ egin{array}{ll} {
m bra}_{ij} x & {
m with \ probability \ } \lambda t + O(t^2), \ {
m death}_i x & {
m with \ probability \ } t + O(t^2), \ {
m x} & {
m with \ probability \ } 1 - \lambda t |E| - t |\Lambda| + O(t^2). \end{array}
ight.$$

Then
$$\mathbb{P}[X_t = y] = 1(x, y) + tG(x, y) + O(t^2)$$
.

Poisson construction of continuous-time Markov chains

Idea: construct a continuous time Markov chain by applying maps $m: S \rightarrow S$ at times of a Poisson process.

Let $(\Omega, \mathcal{F}, \mu)$ be a measurable space with σ -finite, nonatomic measure μ . Recall that a *Poisson point set* with *intensity* μ is a random subset $\omega \subset \Omega$ such that

 $|\omega \cap A|$ is Poisson distributed with mean $\mu(A)$

whenever $A\in \mathcal{F}$, $\mu(A)<\infty$, and

 $ig|\omega\cap A_1ig|,\ldots,ig|\omega\cap A_nig|$ are independent

whenever A_1, \ldots, A_n are disjoint. In particular, if $\varepsilon := \mu(A)$ is small, then

$$\mathbb{P}ig[ig|\omega\cap Aig|=1ig]=arepsilon+O(arepsilon^2),\quad \mathbb{P}ig[ig|\omega\cap Aig|\geq 2ig]=O(arepsilon^2).$$

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Each generator G has a random mapping representation

$$G(x,y) = \sum_{m \in \mathcal{G}} r_m (1_{\{m(x) = y\}} - 1_{\{x = y\}}),$$

where $(r_m)_{m \in \mathcal{G}}$ are nonnegative rates and \mathcal{G} is a collection of maps $m: S \to S$. Let ω be a Poisson point set on $\mathcal{G} \times \mathbb{R}$ with intensity

$$\mu(\{m\} \times A) = r_m \ell(A) \qquad (A \in \mathcal{B}(\mathbb{R})),$$

where $\mathcal{B}(\mathbb{R})$ is the Borel- σ -field on \mathbb{R} and ℓ denotes Lebesgue measure. If $\sum_{m \in \mathcal{G}} r_m < \infty$, then we may order the elements of

$$\omega_{s,t} := \omega \cap \mathcal{G} \times (s,t] = \{(m_1,t_1),\ldots,(m_n,t_n)\}$$

with $t_1 < \cdots < t_n$.

Poisson construction of continuous-time Markov chains

Define random maps $\mathbf{X}_{s,t}:S
ightarrow S$ $(s\leq t)$ by

$$\mathbf{X}_{s,t} := m_n \circ \cdots \circ m_1.$$

(Poisson construction of Markov processes) Define maps $(\mathbf{X}_{s,t})_{s \leq t}$ as above in terms of a Poisson point set ω . Let X_0 be an S-valued random variable, independent of ω . Then

$$X_t := \mathbf{X}_{0,t}(X_0) \qquad (t \ge 0)$$

is a Markov process with generator G.

Remark The sample paths of X are right-continuous. We get left-continuous paths by defining

$$\mathbf{X}_{s,t-}$$
 in terms of $\omega_{s,t-} := \omega \cap \mathcal{G} imes (s,t).$

Example: a finite contact process



Example: a finite contact process



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Stochastic flows

The random maps $(\mathbf{X}_{s,t})_{s \leq t}$ form a *stochastic flow*

$$\mathbf{X}_{s,s} = 1$$
 and $\mathbf{X}_{t,u} \circ \mathbf{X}_{s,t} = \mathbf{X}_{s,u}$

with independent increments, in the sense that

$$X_{t_0,t_1},\ldots,X_{t_{n-1},t_n}$$

are independent for each $t_0 < \cdots < t_n$.

Different stochastic flows can define the same Markov process, as there may be many different ways of writing down a random mapping representation

$$Gf(x) = \sum_{m \in \mathcal{G}} r_m \{f(m(x)) - f(x)\},\$$

for a given generator G.

Compare: Different SDE's can define the same diffusion process.

Form of the generator

Here we have used that for any function $f: S \to \mathbb{R}$,

$$Gf(x) = \sum_{y} G(x, y)f(y)$$

= $\sum_{y} \sum_{m \in \mathcal{G}}^{y} r_m (1_{\{m(x) = y\}} - 1_{\{x = y\}})f(y)$
= $\sum_{m \in \mathcal{G}} r_m \{f(m(x)) - f(x)\}.$

Then, as $t \downarrow 0$,

$$\mathbb{E}^{x}[f(X_{t})] = f(x) + tGf(x) + O(t^{2})$$
$$= \left(1 - t\sum_{m \in \mathcal{G}} r_{m}\right)f(x) + t\sum_{m \in \mathcal{G}} r_{m}f(m(x)) + O(t^{2}).$$

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Example: the contact process

Let (Λ, E) be a finite directed graph. The *contact process* on (Λ, E) with *infection rate* $\lambda \ge 0$ is the continuous-time Markov chain $X = (X_t)_{t \ge 0}$ with state space $\{0, 1\}^{\Lambda}$ and generator

$$egin{aligned} & {\it Gf}(x)\! :=\! \lambda \sum_{\substack{(i,j)\in E}} ig\{fig({\tt bra}_{ij}xig) - f(x)ig\} \ &+ \sum_{i\in \Lambda}ig\{fig({\tt death}_ixig) - f(x)ig\}. \end{aligned}$$

We would like to define contact processes on infinite lattices, e.g., $\Lambda = \mathbb{Z}^d$ with $E = \{(i, j) : |i - j| = 1\}$. If Λ is infinite, then we equip $\{0, 1\}^{\Lambda}$ with the *product topology* which corresponds to *pointwise convergence*

$$x_n \to x$$
 iff $x_n(i) \to x(i)$ $\forall i \in \Lambda$.

By Tychonoff's theorem, $\{0,1\}^{\Lambda}$ is a compact space.

Let S be a compact metrizable space. Let $\mathcal{C}(S)$ denote the space of all continuous functions $f: S \to \mathbb{R}$, equipped with the supremumnorm $\|\cdot\|_{\infty}$, and let $\mathcal{M}_1(S)$ denote the space of all probability measures on S, equipped with the topology of weak convergence.

For any probability kernel K(x, dy) on S and measurable function $f: S \to \mathbb{R}$, we define

$$\mathcal{K}f(x) := \int_{\mathcal{S}} \mathcal{K}(x, \mathrm{d}y)f(y) \qquad (x \in \mathcal{S}).$$

By definition, a *Feller semigroup* is a collection of probability kernels $(P_t)_{t\geq 0}$ on S such that:

(i)
$$P_0 = 1$$
 and $P_s P_t = P_{s+t}$ $(s, t \ge 0)$,
(ii) $S \times [0, \infty) \ni (x, s) \mapsto P_s(x, \cdot) \in \mathcal{M}_1(S)$ is continuous.

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The generator of a Feller semigroup is the operator G defined as

$$Gf := \lim_{t \downarrow 0} t^{-1} (P_t f - f)$$
 (*),

with domain

$$\mathcal{D}(G) := \{ f \in \mathcal{C}(S) : \text{ the limit in } (*) \text{ exists in the norm } \| \cdot \|_{\infty} \}.$$

The *Hille-Yosida theorem* gives necessary and sufficient conditions for an operator G to generate a Feller semigroup.

The domain $\mathcal{D}(G)$ is a dense subspace of $\mathcal{C}(S)$. For each $f \in \mathcal{D}(G)$, the function $t \mapsto P_t f$ is given by the unique solution to the *backward equation*

$$\frac{\partial}{\partial t}P_t f = GP_t f$$
 $(t \ge 0)$ with $P_0 f = 1$.

Let S be a compact metrizable space and let $(P_t)_{t\geq 0}$ be a Feller semigroup of transition kernels on S. Then, for each probability law μ on S, there exists a process $(X_t)_{t\geq 0}$ such that

► The sample paths t → X_t are a.s. cadlag, i.e., right-continuous with left limits.

$$\blacktriangleright \mathbb{P}[X_0 \in \cdot] = \mu.$$

$$\blacktriangleright \mathbb{P}^{x} \left[X_{u} \in \cdot \mid (X_{s})_{0 \leq s \leq t} \right] = P_{u-t}(X_{t}, \cdot) \text{ a.s. } (t \leq u).$$

Such a process is called a Feller process.

For any map $m: S^{\wedge} \to S^{\wedge}$, let

$$\mathcal{D}(m) := \left\{ i \in \Lambda : \exists x \in S^{\Lambda} \text{ s.t. } m(x)(i) \neq x(i)
ight\}$$

denote the set of lattice points whose values can possibly be changed by m. By definition, m is a *local map* if and only if (i) $\mathcal{D}(m)$ is finite,

(ii) *m* is continuous w.r.t. the product topology.

Say that a site $j \in \Lambda$ is *m*-relevant for some $i \in \Lambda$ if

$$\exists x,y \in S^{\Lambda} ext{ s.t. } m(x)(i)
eq m(y)(i) ext{ and } x(k) = y(k) ext{ } orall k
eq j,$$

and define

$$\mathcal{R}_i(m) := \{j \in \Lambda : j \text{ is } m \text{-relevant for } i\}.$$

Remark A map $m : S^{\Lambda} \to S^{\Lambda}$ is continuous w.r.t. the product topology if and only if for each $i \in \Lambda$:

(i) $\mathcal{R}_i(m)$ is finite, (ii) x(j) = y(j) for all $j \in \mathcal{R}_i(m)$ implies m(x)(i) = m(y)(i).

Trivially

$$\mathcal{R}_i(m) = \{i\} \text{ if } i \notin \mathcal{D}(m).$$

In our example:

$$\mathcal{D}(\texttt{bra}_{ij}) = \{j\}$$
 $\mathcal{R}_j(\texttt{bra}_{ij}) = \{i, j\},$
 $\mathcal{D}(\texttt{death}_i) = \{i\}$ $\mathcal{R}_i(\texttt{death}_i) = \emptyset.$

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Let \mathcal{G} me a collection of local maps $m: S^{\Lambda} \to S^{\Lambda}$ and let $(r_m)_{m \in \mathcal{G}}$ be nonnegative constants. As before, we let ω be a Poisson point set on $\mathcal{G} \times \mathbb{R}$ with intensity

$$\mu(\{m\} \times A) = r_m \,\ell(A) \qquad (A \in \mathcal{B}(\mathbb{R})).$$

For infinite lattices (e.g., for the contact process on \mathbb{Z}^d) we typically have $\sum_{m \in \mathcal{G}} r_m = \infty$ which means

$$\omega_{s,t} := \omega \cap \mathcal{G} \times (s,t]$$

has a.s. infinitely many elements for any s < t. As a result, we cannot order the elements of $\omega_{s,t}$ by their time coordinates. Nevertheless, for each finite subset $\tilde{\omega} \subset \omega_{s,t}$, we can still define

Theorem Assume that

$$\sup_{i\in\Lambda} \sum_{\substack{m\in\mathcal{G}\\\mathcal{D}(m)\ni i}} r_m |\mathcal{R}_i(m)| < \infty.$$

Then a.s., for each $x \in S^{\Lambda}$ the limit

$$\mathbf{X}_{s,t}(x) := \lim_{\tilde{\omega}_n \uparrow \omega_{s,t}} \mathbf{X}_{s,t}^{\tilde{\omega}_n}(x)$$

exists, does not depend on the sequence of finite sets $\tilde{\omega}_n \uparrow \omega_{s,t}$, and defines a stochastic flow $(\mathbf{X}_{s,t})_{s \leq t}$. If X_0 is an S^{Λ} -valued random variable, independent of ω , then

$$X_t := \mathbf{X}_{0,t}(X_0) \qquad (t \ge 0)$$

defines a Feller process with generator

$$Gf(x) = \sum_{m \in \mathcal{G}} r_m \{f(m(x)) - f(x)\}.$$

Lemma 1 One has

$$\mathbb{E}[|\mathcal{R}_i(\mathbf{X}_{s,t})|] \leq e^{R(t-s)} \qquad (s \leq t),$$

where

$$R := \sup_{i \in \Lambda} \sum_{\substack{m \in \mathcal{G} \\ \mathcal{D}(m) \ni i}} r_m (|\mathcal{R}_i(m)| - 1).$$
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Generator construction of Interacting Particle Systems

For any
$$f \in \mathcal{C}(S^{\Lambda})$$
 and $i \in \Lambda$, we define
 $\delta f(i) := \sup \{ |f(x) - f(y)| : x, y \in S^{\Lambda}, x(j) = y(j) \ \forall j \neq i \},$
 $\mathcal{C}_{sum} := \{ f \in \mathcal{C}(S^{\Lambda}) : |||f||| := \sum_{i \in \Lambda} \delta f(i) < \infty \}$

We call $f \in \mathcal{C}_{sum}$ a function of *summable variation*. One has

$$\left|f(x)-f(y)\right|\leq \sum_{i:\,x(i)\neq y(i)}\delta f(i)\leq |||f|||.$$

Lemma 2 If $(P_t)_{t\geq 0}$ is the Feller semigroup of an interacting particle system with generator G, then $\mathcal{C}_{summ} \subset \mathcal{D}(G)$ and

$$|||P_t f||| \le e^{Rt} |||f||| \qquad (t \ge 0),$$

where R is the constant in (1).

Generator construction of Interacting Particle Systems

For any probability kernel κ on S^{Λ} , set

 $\mathcal{D}(\kappa) := \big\{ i \in \Lambda : \exists x \text{ s.t. } \kappa \big(x, \{ y : y(i) \neq x(i) \} \big) > 0 \big\}.$

By definition, a *local probability kernel* is a probab. kernel such that

(i) $\mathcal{D}(\kappa)$ is finite, (ii) $x \mapsto \kappa(x, \cdot)$ is continuous w.r.t. weak convergence. Liggett (1985) writes generators in the form

$$Gf(x) = \sum_{\kappa \in \mathcal{K}} r_{\kappa} \{ \int \kappa(x, \mathrm{d}y) f(y) - f(x) \},$$

where \mathcal{K} is a collection of local probability kernels and $(r_{\kappa})_{\kappa \in \mathcal{K}}$ are nonnegative rates. Under suitable conditions on the rates, he proves the closure of G generates a Feller semigroup and

$$|||P_t f||| \le e^{R't} |||f||| \quad (t \ge 0)$$

for some constant $R' < \infty$.

Proofs

Proof of Lemma 1 By a cut-off argument, w.l.o.g. we assume that Λ is finite. Then we can apply our previous result about the Poisson construction of continuous-time Markov chains to conclude that

$$\left(\mathcal{R}_i(\mathbf{X}_{-t,0})\right)_{t\geq 0}$$

is a set-valued Markov process that can be bounded from above by a process that jumps as

$$A\mapsto (Aackslash \mathcal{D}(m))\cup igcup_{i\in\mathcal{D}(m)\cap A}\mathcal{R}_i(m)$$
 with rate r_m

for each $m \in \mathcal{G}$. Let $(P_t)_{t\geq 0}$ denote the semigroup of the set-valued process and let G be its generator. Let f be the function f(A) := |A|. Then

$$Gf(A) = \sum_{m \in \mathcal{G}} r_m \Big\{ f\big((A \setminus \mathcal{D}(m)) \cup \bigcup_{i \in \mathcal{D}(m) \cap A} \mathcal{R}_i(m) \big) - f(A) \Big\}$$

Proofs

It follows that

$$Gf(A) \leq \sum_{m \in \mathcal{G}} r_m \sum_{i \in \mathcal{D}(m) \cap A} (|\mathcal{R}_i(m)| - 1)$$

=
$$\sum_{i \in A} \sum_{m: \mathcal{D}(m) \ni i} r_m (|\mathcal{R}_i(m)| - 1) \leq R|A|,$$

and hence

$$\frac{\partial}{\partial t} \left(e^{-Rt} P_t f \right) = -Re^{-Rt} P_t f + e^{-Rt} P_t G_n f = e^{-Rt} P_t (G_n f - Rf) \le 0$$

and therefore $e^{-Rt}P_tf \leq e^{-R0}P_0f = f,$ which means in particular that

$$\mathbb{E}\big[|\mathcal{R}_i(\mathbf{X}_{-t,0})|\big] = P_t f(\{i\}) \le e^{Rt} f(\{i\}) = e^{Rt} \qquad (t \ge 0).$$

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Proofs

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Proof of Lemma 2 We want to prove that $||P_{t}f|| \leq e^{Rt} |||f||| \quad \text{with} \quad |||f||| := \sum \delta f(i)$

$$i \in \Lambda \text{ and let } x, y \in S^{\Lambda} \text{ with } x(j) = y(j) \forall j \neq i. \text{ Then}$$

$$|P_t f(x) - P_t f(y)| = |\mathbb{E}[f(\mathbf{X}_{0,t}(x))] - \mathbb{E}[f(\mathbf{X}_{0,t}(y))]|$$

$$\leq \mathbb{E}[|f(\mathbf{X}_{0,t}(x)) - f(\mathbf{X}_{0,t}(y))|]$$

$$\leq \sum_j \mathbb{P}[\mathbf{X}_{0,t}(x)(j) \neq \mathbf{X}_{0,t}(y)(j)]\delta f(j)$$

$$\leq \sum_j \mathbb{P}[i \in \mathcal{R}_j(\mathbf{X}_{0,t})]\delta f(j)$$

It follows that

$$\sum_{i} \delta P_t f(i) \leq \sum_{ij} \mathbb{P} \big[i \in \mathcal{R}_j(\mathbf{X}_{0,t}) \big] \delta f(j) \leq e^{Rt} \sum_{j} \delta f(j).$$

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Unique ergodicity

For any probability measure μ on S^{Λ} , write

$$\mu P_t := \int \mu(\mathrm{d} x) P_t(x, \cdot) \qquad (t \ge 0).$$

If $X=(X_t)_{t\geq 0}$ has initial law $\mathbb{P}[X_0\in\,\cdot\,]=\mu$, then,

$$\mathbb{P}[X_t \in A] = \int \mathbb{P}[X_0 \in \mathrm{d}x] \mathbb{P}[X_t \in A \mid X_0 \in \mathrm{d}x] = \int \mu(\mathrm{d}x) P_t(x, A),$$

i.e., X_t has law μP_t . By definition, a probability measure ν on S^{Λ} is an *invariant law* if

$$\nu P_t = \nu \qquad (t \ge 0).$$

The process $X = (X_t)_{t \ge 0}$ is stationary if and only if $\mathbb{P}[X_0 \in \cdot]$ is an invariant law.

Unique ergodicity

Proposition If the constant *R* from (1) satisfies R < 0, then the interacting particle system has a unique invariant law ν , and

$$\mu P_t \underset{t \to \infty}{\Longrightarrow} \nu$$

for any initial law μ .

Proof If R < 0, then

$$\mathbb{P}\big[\mathcal{R}_i(\mathbf{X}_{0,t}) = \emptyset\big] \xrightarrow[t \to \infty]{} 1.$$

As a consequence, the limit

$$X_t(i) := \lim_{s \to \infty} \mathbf{X}_{t-s,t}(x)(i) \qquad (i \in \Lambda)$$

exists a.s. for each $i \in \Lambda$ and $t \in \mathbb{R}$, and does not depend on the choice of $x \in S^{\Lambda}$.

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One can check that this defines a stationary process $(X_t)_{t \in \mathbb{R}}$. Let Y have law μ and be independent of the Poisson processes used to construct the flow $(\mathbf{X}_{s,t})_{s \leq t}$. Then $\mathbf{X}_{-t,0}(Y)$ has law μP_t and

$$\mathbf{X}_{-t,0}(Y) \underset{t \to \infty}{\longrightarrow} X_0$$
 a.s.,

proving that $\mu P_t \underset{t \to \infty}{\Longrightarrow} \nu$.

Remark 1 This method is called *coupling from the past*.

Remark 2 The proof works more generally if

$$\mathbb{P}\big[\mathcal{R}_i(\mathbf{X}_{0,t}) = \emptyset\big] \underset{t \to \infty}{\longrightarrow} 1 \qquad (i \in \Lambda),$$

which may happen even if $R \ge 0$.

Recall that

$$\begin{split} \mathcal{D}(\texttt{bra}_{ij}) &= \{j\} & \mathcal{R}_j(\texttt{bra}_{ij}) = \{i, j\}, \\ \mathcal{D}(\texttt{death}_i) &= \{i\} & \mathcal{R}_i(\texttt{death}_i) = \emptyset. \end{split}$$

For the nearest neighbor contact process on \mathbb{Z}^d , this yields

$$R=2d\lambda-1,$$

which is < 0 iff $\lambda < 1/2d$. Since $\mathbf{X}_{s,t}(\underline{0}) = \underline{0}$, we conclude that for $\lambda < 1/2d$, the measure δ_0 is the unique invariant law.

Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 0.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 1.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 2.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 3.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 4.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 5.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 6.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 7.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 8.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 9.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 10.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 11.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 12.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 14.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 15.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 16.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 17.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 18.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 19.

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Contact process with infection rate $\lambda = 2$ and death rate d = 1. Time t = 20.

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A stochastic Ising model

Let (Λ, E) be an undirected graph and let $S = \{-1, +1\}$. For any $x \in S^{\Lambda}$, we call

$$M_i(x) := \sum_{j: \{i,j\} \in E} x(j)$$

the *local magnetization* around $i \in \Lambda$. Let $\kappa_i(x, \cdot)$ denote the law of a random variable X such that

$$\mathbb{P}[X(i)=\pm 1]=rac{e^{eta\pm M_i(x)}}{e^{eta\pm M_i(x)}+e^{eta\mp M_i(x)}},$$

and X(j) = x(j) a.s. for all $j \neq i$. Then κ_i is a local probability kernel and

$$Gf(x) = \sum_{i \in \Lambda} \left(\int \kappa_i(x, \mathrm{d}y) f(y) - f(x) \right)$$

defines the generator of a *stochastic Ising model* with *Glauber dynamics*.

When the parameter β is large, nearby spins like have the same sign.

We start the process in product measure for different values of β and see what happens.



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Instead of allowing only two states -1, +1, we can more generally allow $q \ge 2$ states $1, \ldots, q$.

Each person i chooses a new state at times of a Poisson process with rate 1.

The probability that the newly chosen state is $k \in \{1, \dots, q\}$ equals

$$\frac{\mathrm{e}^{\beta M_i(k)}}{\sum_{m=1}^{q} \mathrm{e}^{\beta M_i(m)}},$$

where $M_i(k)$ denotes the number of neighbors of *i* that are in the state *k*.

Setting q = 2 and replacing β by 2β yields the Ising model.



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 $\beta = 1.2$, time t = 8.

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 $\beta = 1.2$, time t = 16.

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 $\beta = 1.2$, time t = 64.

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The voter model

In the *voter model*, each site $i \in \Lambda$ is occupied by an organism with genetic type $x(i) \in S$.

The organism at site i dies at times of a Poisson process with rate 1 and is replaced by the offspring of a randomly chosen neighbor.

Using the *voter map*

$$ext{vot}_{ji}(x)(k) := \left\{egin{array}{c} x(j) & ext{if } k=i, \ x(k) & ext{otherwise,} \end{array}
ight.$$

we can give the following random mapping representation of the generator:

$$Gf(x) = \sum_{(i,j)\in E} \frac{1}{N_i} \{f(\operatorname{vot}_{ji} x) - f(x)\},$$

where $N_i := |\{j : (j, i) \in E\}|$ is the number of neighbors of *i*.



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Time t = 0.25.

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The voter model



Time t = 0.5.

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The voter model



Time t = 1.

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The voter model



Time t = 2.



Time t = 4.



Time t = 8.



Time t = 16.



Time t = 31.25.

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The voter model



Time t = 62.5.



Time t = 125.



Time t = 250.

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Time t = 500.

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The behavior of the voter model strongly depends on the dimension.

Clustering in dimensions d = 1, 2.

Stable behavior in dimensions $d \ge 3$.

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Cut of 3-dimensional model, time t = 2.

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Cut of 3-dimensional model, time t = 4.

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Cut of 3-dimensional model, time t = 8.



Cut of 3-dimensional model, time t = 16.

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Cut of 3-dimensional model, time t = 32.

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Cut of 3-dimensional model, time t = 64.

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Cut of 3-dimensional model, time t = 125.

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Cut of 3-dimensional model, time t = 250.

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A one-dimensional voter model



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A one-dimensional Potts model



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In the *biased* voter model with two states $\{0, 1\}$, each organism *i* changes its type $X_t(i)$ with the rates

- $0\mapsto 1$ with rate $(1+s)\cdot$ fraction of type 1 neighbors,
- $1\mapsto 0$ with rate $1\cdot$ fraction of type 0 neighbors,

where s > 0 gives type 1 a (small) advantage.

Contrary to the voter model, even if we start with just a single organism of type 1, there is a positive probability that type 1 never dies out.

Models spread of advantageous mutation.

Biased voter model with s = 0.2. Time t = 0.

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Biased voter model with s = 0.2. Time t = 10.

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Biased voter model with s = 0.2. Time t = 20.

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Biased voter model with s = 0.2. Time t = 30.

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Biased voter model with s = 0.2. Time t = 40.

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Biased voter model with s = 0.2. Time t = 50.

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Biased voter model with s = 0.2. Time t = 60.



Biased voter model with s = 0.2. Time t = 70.



Biased voter model with s = 0.2. Time t = 80.



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Biased voter model with s = 0.2. Time t = 110.


Biased voter model with s = 0.2. Time t = 120.

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Biased voter model with s = 0.2. Time t = 130.

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Biased voter model with s = 0.2. Time t = 140.

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Biased voter model with s = 0.2. Time t = 150.

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Biased voter model with s = 0.2. Time t = 160.

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We can extend the biased voter model by also allowing spontaneous jumps from 1 to 0.

$0\mapsto 1$	with rate $(1+s)\cdot$ fraction of type 1 neighbors,
$1\mapsto 0$	with rate $1\cdot fraction$ of type 0 neighbors
	+ d,

where s > 0 gives type 1 an advantage and $d \ge 0$ is a *death rate*. This models the fact that genes may become disfunctional due to deleterious mutations.

Whether 1's have a positive probability to survive now depends in a nontrivial way on s and d.

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A rebellious voter model

The rebellious voter map is defined as

$$\operatorname{rvot}_{kji}(x)(l) := \begin{cases} 1-x(i) & \text{if } l=i \text{ and } x(k) \neq x(j), \\ x(l) & \text{otherwise.} \end{cases}$$

The *rebellious voter model* is the one-dimensional model with generator

$$Gf(x) := \alpha \sum_{i} \{f(\operatorname{vot}_{i,i+1}(x)) - f(x)\} \\ + \alpha \sum_{i} \{f(\operatorname{vot}_{i,i-1}(x)) - f(x)\} \\ + (1 - \alpha) \sum_{i} \{f(\operatorname{rvot}_{i-1,i,i+1}(x)) - f(x)\} \\ + (1 - \alpha) \sum_{i} \{f(\operatorname{rvot}_{i+1,i,i-1}(x)) - f(x)\}.$$

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A rebellious voter model



A rebellious voter model



Reaction diffusion models

Another rich class of models are reaction diffusion models.

These are systems of particles that perform independent random walks and interact when they are near to each other.

Let $X_t(i) = 1$ (resp. 0) signify the presence (resp. absence) of a particle and consider the maps $rw_{ij} : \{0,1\}^{\mathbb{Z}} \to \{0,1\}^{\mathbb{Z}}$

$$\mathbf{rw}_{i,j} \mathbf{x}(k) := \begin{cases} 0 & \text{if } k = i, \\ \mathbf{x}(i) \lor \mathbf{x}(j) & \text{if } k = j, \\ \mathbf{x}(k) & \text{otherwise.} \end{cases}$$

The process with generator

$$G = \frac{1}{2} \sum_{i \in \mathbb{Z}} \left\{ f(\texttt{rw}_{i,i+1}x) - f(x) \right\} + \frac{1}{2} \sum_{i \in \mathbb{Z}} \left\{ f(\texttt{rw}_{i,i-1}x) - f(x) \right\}$$

describes coalescing random walks.

Coalescing random walks



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In analogy with the branching map

$$ext{bra}_{ij}x(k) := \left\{ egin{array}{cc} x(i) \lor x(j) & ext{if } k=j, \ x(k) & ext{otherwise,} \end{array}
ight.$$

we can also define a cooperative branching map

$$\operatorname{coop}_{ii'j} x(k) := \begin{cases} (x(i) \wedge x(i')) \lor x(j) & \text{if } k = j, \\ x(k) & \text{otherwise.} \end{cases}$$

Branching and coalescing random walks



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Cooperative branching and coalescence



Cooperative branching



Two more maps of interest are the annihilating random walk map

$$\operatorname{arw}_{ij} x(k) := \begin{cases} 0 & \text{if } k = i, \\ x(i) + x(j) \mod(2) & \text{if } k = j, \\ x(k) & \text{otherwise,} \end{cases}$$

and the annihilating branching map

$$abra_{ij}x(k) := \begin{cases} x(i) + x(j) \mod(2) & \text{if } k = j, \\ x(k) & \text{otherwise,} \end{cases}$$

A cancellative system



Jan M. Swart Spatial Models in Population Biology

A cancellative system



Define a killing map as

$$ext{kill}_{ij}x(k) := \left\{ egin{array}{cc} (1-x(i)) \wedge x(j) & ext{if } k=j, \ x(k) & ext{otherwise}, \end{array}
ight.$$

which says that the particle at i, if present, kills any particle at j.

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Branching and killing



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