

Let Markov chains evolve along genealogies

Vincent Bansaye,
based on a joint work Chunmao Huang

Ecole Polytechnique

10 december. Angers, SMEEG.



Discrete time genealogy and Markov chains

We consider a population of individuals with a trait described by

- a **genealogical tree** \mathbb{T} in discrete time (fixed or random)
- a **trait for each individual**, whose dynamic is given by (Markovian) **transition kernels**, with possible dependence on the number of offsprings and on the generation (time inhomogeneity).

Limit theorems for large populations. We assume *here* that the number of individuals in generation n goes to ∞ as $n \rightarrow \infty$.

Two particular classes studied here : emergence of deterministic proportions under neutrality assumption or branching property.

Discrete time genealogy and Markov chains

We consider a population of individuals with a trait described by

- a **genealogical tree** \mathbb{T} in discrete time (fixed or random)
- a **trait for each individual**, whose dynamic is given by (Markovian) **transition kernels**, with possible dependence on the number of offsprings and on the generation (time inhomogeneity).

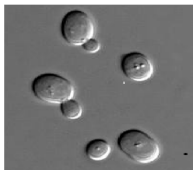
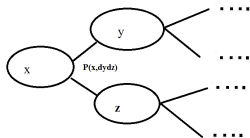
Limit theorems for large populations. We assume *here* that the number of individuals in generation n goes to ∞ as $n \rightarrow \infty$.

Two particular classes studied here : emergence of deterministic proportions under neutrality assumption or branching property.

Motivation 1 : random transmission in cell division

One generation = cell division time. $k \in \{0, 1, 2\}$.

Trait = number of parasites, plasmids, mitochondrias, external DNA ... ;
age, growth rate, damages ... of the cell.



Strong asymmetry may be observed.

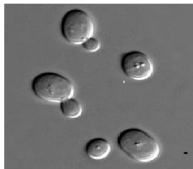
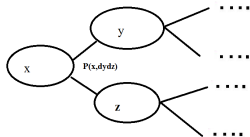
Two examples of models :

- Kimmel's branching models. (Binomial repartition of plasmids or parasites in the two daughter cells)
- Bifurcating autoregressive process for cellular aging
($X_{n+1} = a_n X_n + b_n$).

Motivation 1 : random transmission in cell division

One generation = cell division time. $k \in \{0, 1, 2\}$.

Trait = number of parasites, plasmids, mitochondrias, external DNA ... ;
age, growth rate, damages ... of the cell.



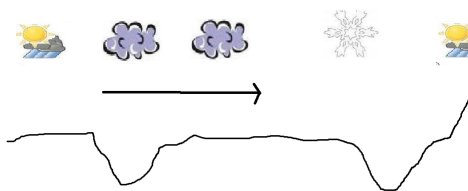
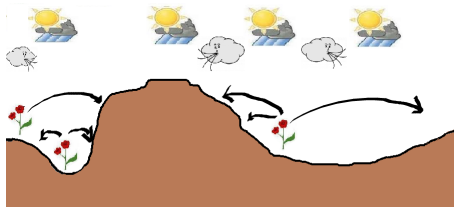
Strong asymmetry may be observed.

Two examples of models :

- Kimmel's branching models. (Binomial repartition of plasmids or parasites in the two daughter cells)
- Bifurcating autoregressive process for cellular aging
($X_{n+1} = a_n X_n + b_n$).

Motivation 2 : Reproduction-dispersion models

1 generation = 1 year ; Trait = location.

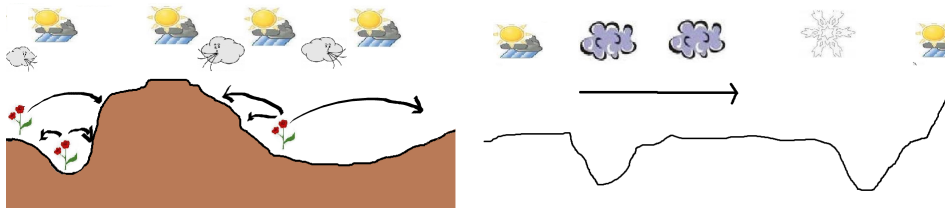


Shape of the repartition of the population with global limitation of resources on compact set ;
Invasion dynamics and effect of time and space non-homogeneity.

Evolution with time non-homogeneity.

Motivation 2 : Reproduction-dispersion models

1 generation = 1 year ; Trait = location.



Shape of the repartition of the population with global limitation of resources on compact set ;
 Invasion dynamics and effect of time and space non-homogeneity.

Evolution with time non-homogeneity.

- Markov chain **along binary or Galton-Watson tree** (Athreya Khang 98, Guyon 05, Delmas Marsalle and al ...)
- **Multitype** branching processes (well known with finite number of types ; pioneering works of Moy, Seneta, Vere Jones, Kesten for infinite number of types).
- **Branching random walks** when P is additive (Biggins), with possibly random environment (see e.g. works of Comets, Gantert Müller, Yoshida.)

Here the novelties lie in *non branching trees*, *time non homogeneity* (with non additive P) and/or *infinite numbers of types*.

- Markov chain **along binary or Galton-Watson tree** (Athreya Khang 98, Guyon 05, Delmas Marsalle and al ...)
- **Multitype** branching processes (well known with finite number of types ; pioneering works of Moy, Seneta, Vere Jones, Kesten for infinite number of types).
- **Branching random walks** when P is additive (Biggins), with possibly random environment (see e.g. works of Comets, Gantert Müller, Yoshida.)

Here the novelties lie in *non branching trees*, *time non homogeneity* (with non additive P) and/or *infinite numbers of types*.

Markov chain along genealogies

The model is specified by $P_n^{(k)}(x, dx_1 \cdots dx_k)$: if $u \in \mathbb{T}$ belongs to the generation n and has k offsprings, then

$$\begin{aligned} \mathbb{P} \left(X(u_1) \in dx_1 \cdots X(u_k) \in dx_k \mid (X(v) : |v| \leq n) \right) \\ = P_n^{(k)}(X(u), dx_1 \cdots dx_k). \end{aligned}$$

Question :

What is the **proportion of individuals** with some given trait, i.e. the asymptotic behavior of $X_n(A)/X_n(\mathcal{X})$?

Early separation of the genealogies

Proposition

Let $A \in \mathcal{B}_{\mathcal{X}}$. We assume that

- (i) $N_n \rightarrow \infty$ as $n \rightarrow \infty$;
- (ii) $\limsup_{n \rightarrow \infty} \mathbb{P}(|U_n \wedge V_n| \geq K) \rightarrow 0$ as $K \rightarrow \infty$, where U_n, V_n are uniformly and independently chosen in generation n ;
- (iii) there exists $\mu_n(A)$ such that for all $u \in \mathbb{T}$ and $x \in \mathcal{X}$,

$$\lim_{n \rightarrow \infty} \mathbb{P} \left(X(U_n^{(u)}) \in A \mid X(u) = x \right) - \mu_n(A) = 0,$$

where $U_n^{(u)}$ is uniformly chosen in generation n .

Then $\frac{X_n(A)}{X_n(\mathcal{X})} - \mu_n(A) \xrightarrow{n \rightarrow \infty} 0$ in L^2 .

Examples for (i-ii) : supercritical branching processes.

Non late separation of the genealogies

Proposition

Let $A \in \mathcal{B}_{\mathcal{X}}$. We assume that

- (i) $N_n \rightarrow \infty$ as $n \rightarrow \infty$;
- (ii) $\limsup_{n \rightarrow \infty} \mathbb{P}(|U_n \wedge V_n| \geq n - K) \rightarrow 0$ as $K \rightarrow \infty$, where U_n, V_n are uniformly and independently chosen in generation n ;
- (iii) there exists $\mu_n(A)$ such that

$$\lim_{n \rightarrow \infty} \sup_{u \in \mathbb{T}, x \in \mathcal{X}} \left| \mathbb{P} \left(X(U_n^{(u)}) \in A \mid X(u) = x \right) - \mu_n(A) \right| = 0,$$

where $U_n^{(u)}$ is an individual uniformly chosen in generation n .

Then

$$\frac{X_n(A)}{X_n(\mathcal{X})} - \mu_n(A) \rightarrow 0 \quad \text{in } L^2.$$

Example for (ii) : bounded number of offsprings. 

Two examples where (iii)[ergodic assumptions] can be obtained easily as the ergodicity of an auxiliary Markov chain.

- when the transitions $P^{(k)}$ do not depend on the number of offsprings ;
- when the genealogical tree \mathbb{T} is a branching process.

What about branching Markov chains ? (non neutral framework, multitype branching processes, with possibly infinite number of types).

The individuals reproduce independently and each individual with trait $x \in \mathcal{X}$ in environment \mathbf{e}

- the reproduction law is $N(x, \mathbf{e})$
- the traits of the k offsprings are given by

$$P^{(k)}(x, \mathbf{e}, dx_1 \dots dx_k) \quad (k \geq 1)$$

- the offsprings live in environment $T\mathbf{e}$.

The auxiliary chain

We consider the transition kernel

$$Q_k(x, \mathbf{e}, dy) = \mathbb{E}_{\delta_x, \mathbf{e}}(Z_1(dy)) \frac{\mathbb{E}_{\delta_y, T\mathbf{e}}(Z_{k-1}(\mathcal{X}))}{\mathbb{E}_{\delta_x, \mathbf{e}}(Z_k(\mathcal{X}))}$$

and the trait $X_i(u)$ of the ancestor of u in generation i .

Lemma

For all $n \in \mathbb{N}$, $x \in \mathcal{X}$ and $F \in \mathcal{B}(\mathcal{X}^{n+1})$ non-negative, we have

$$\mathbb{E}_{\mathbf{e}, \delta_x} \left(\sum_{|u|=n} F(X_0(u), \dots, X_n(u)) \right) = \mathbb{E}_{\mathbf{e}, \delta_x}(Z_n(\mathcal{X})) \mathbb{E}_{\mathbf{e}, x}(F(Y_0^{(n)}, \dots, Y_n^{(n)})),$$

where $(Y_i^{(n)} : i = 0, \dots, n)$ is a non-homogeneous Markov chain with kernels $(Q_{n-i}(\cdot, T^i \mathbf{e}, \cdot) : i = 0, \dots, n-1)$.

In particular $\mathbb{E}_{\mathbf{e}, \delta_x} \left(\sum_{|u|=n} f(Z(u)) \right) = \mathbb{E}_{\mathbf{e}, \delta_x}(Z_n(\mathcal{X})) \mathbb{E}_{\mathbf{e}, x}(f(Y_n^{(n)})).$

Law of large numbers (I)

Theorem

We assume that there exists a measure ν with finite first moment such that for all $x \in \mathcal{X}, k, l \geq 0$,

$$\mathbb{P}(N(x, T^k \mathbf{e}) \geq l) \leq \nu[l, \infty).$$

Assume also that there exists a sequence of probability measure μ_n such that

$$\sup_{\lambda \in \mathcal{M}_1(\mathcal{X})} |Q_{i,n}(\lambda, T^i \mathbf{e}, f \circ f_n) - \mu_n(f)| \rightarrow 0,$$

uniformly for $n - i \rightarrow \infty$. Then,

$$\frac{f_n \cdot Z_n(f)}{Z_n(\mathcal{X})} - \mu_n(f) \xrightarrow{n \rightarrow \infty} 0$$

$\mathbb{P}_{\mathbf{e}}$ a.s. on the event $\left\{ \forall n, Z_n(\mathcal{X}) > 0; \liminf_{n \rightarrow \infty} Z_{n+1}(\mathcal{X})/Z_n(\mathcal{X}) > 1 \right\}$.

Law of large numbers (II)

Theorem

Let $\mathbf{e} \in E$, $x \in \mathcal{X}$ and $f \in \mathcal{B}(\mathcal{X})$ bounded.

Under some technical assumptions (bounded second moment and control on $(x, k) \rightarrow \mathbb{E}_{T^k \mathbf{e}, \delta_x}(Z_n(\mathcal{X}))$) and assuming that there exists a sequence of probability measures μ_n on \mathcal{X} such that

$$\sup_{i \in \mathbb{N}} \sum_{n \geq i} \sup_{\lambda \in \mathcal{M}_1} |Q_{i,n}(\lambda, T^i \mathbf{e}, f \circ f_n) - \mu_n(f)|^2 < \infty.$$

Then, $Z_n(\mathcal{X})/\mathbb{E}_{\mathbf{e}}(Z_n(\mathcal{X}))$ is bounded in $L_{\mathbf{e}}^2$ and

$$\frac{f_n \cdot Z_n(f) - \mu_n(f) Z_n(\mathcal{X})}{\mathbb{E}_{\mathbf{e}}(Z_n(\mathcal{X}))} \xrightarrow{n \rightarrow \infty} 0 \quad \mathbb{P}_{\mathbf{e}} \text{ a.s.}$$

Applications

- Multitype branching process in varying / random environment
- Deriving the shape of the population from limit theorems for Random Walk in Random Environment
- A sufficient condition is given by [Doebelin type assumptions]

$$\mathbb{E}_{\mathbf{e}, \delta_x}(Z_1(A)) \leq M(\mathbf{e})\mathbb{E}_{\mathbf{e}, \delta_y}(Z_1(A)) \quad (x, y \in \mathcal{X}).$$

+control on $\mathbf{e} \rightarrow M(\mathbf{e})$.

- Using Lyapounov assumptions : *WIP*.

Applications

- Multitype branching process in varying / random environment
- Deriving the shape of the population from limit theorems for Random Walk in Random Environment
- A sufficient condition is given by [Doebelin type assumptions]

$$\mathbb{E}_{\mathbf{e}, \delta_x}(Z_1(A)) \leq M(\mathbf{e})\mathbb{E}_{\mathbf{e}, \delta_y}(Z_1(A)) \quad (x, y \in \mathcal{X}).$$

+control on $\mathbf{e} \rightarrow M(\mathbf{e})$.

- Using Lyapounov assumptions : *WIP*.

Two more questions :

- How many individuals have some given (non common) trait ? (local densities and extremal traits)
- What is the growth rate of the population ? When does the population survives ?

Local densities and extremal individuals

Question : can we get

$$Z_n(A_n) \asymp \mathbb{E}_{\mathbf{e}}(Z_n(\mathcal{X}))\mathbb{P}(Y_n^{(n)} \in A_n)$$

from the many-to-one formula

$$\mathbb{E}_{\mathbf{e}}(Z_n(A_n)) = \mathbb{E}_{\mathbf{e}}(Z_n(\mathcal{X}))\mathbb{P}(Y_n^{(n)} \in A_n)$$

and the trajectory of $Y^{(n)}$ associated to the (large deviation event) A_n .

Problem : lower bound.

Solution : coupling by a branching process in varying environment in the first steps and LLN I for the rest of the time.

Application : under monotonicity and neutrality assumption, with a control of the trajectory of the auxiliary process.

Let X be the Markov chain with transition inherited from $P^{(k)}$.

Theorem

Assume that X satisfies a LDP with good rate function $I_{\mathbf{e}}$ in environment \mathbf{e} and $\log m : \mathcal{X} \times E \rightarrow (-\infty, \infty)$ is continuous and bounded. Then, for every $x \in \mathcal{X}$,

$$\lim_{n \rightarrow \infty} \frac{1}{n} \log \mathbb{E}_{\mathbf{e}, \delta_x}(Z_n(\mathcal{X})) = \sup_{\mu \in \mathcal{M}_1(\mathcal{X} \times E)} \left\{ \int_{\mathcal{X} \times E} \log(m(x, e)) \mu(dxde) - I_{\mathbf{e}}(\mu) \right\}$$

and

$$M_{\mathbf{e}} := \left\{ \mu \in \mathcal{M}_1(\mathcal{X} \times E) : \int \log(m(x, e)) \mu(dxde) - I_{\mathbf{e}}(\mu) = \varrho_{\mathbf{e}} \right\}$$

is compact and non empty.

Applications in fixed environment (via Sanov's theorem) and stationary ergodic random environment (via Seppäläinen LDP 95).