Splitting trees with neutral mutations at birth

MATHIEU RICHARD

June 10, 2013

Abstract

We consider a population model where individuals behave independently from each other and whose genealogy is described by a chronological tree called splitting tree. The individuals have i.i.d. (non-exponential) lifetime durations and give birth at constant rate to clonal or mutant children in an infinitely many alleles model with neutral mutations.

First, to study the allelic partition of the population, we are interested in its frequency spectrum, which, at a fixed time, describes the number of alleles carried by a given number of individuals and with a given age. We compute the expected value of this spectrum and obtain some almost sure convergence results thanks to classical properties of Crump-Mode-Jagers (CMJ) processes counted by random characteristics.

Then, by using multitype CMJ-processes, we get asymptotic properties about the number of alleles that have undergone a fixed number of mutations with respect to the ancestral allele of the population.

Key words: Splitting tree; Crump-Mode-Jagers branching processes; neutral mutation; infinitely many alleles model; frequency spectrum; multitype branching processes; almost-sure limit theorem; epidemiology.

MSC 2010: Primary 60J80, secondary 60J28, 92D25, 60J85, 60J27, 92D10, 60G55.

1 Introduction

We consider a general branching model with neutral mutations occurring at birth. We suppose that individuals carry alleles, have i.i.d. (and not necessarily exponential) life lengths and give birth at constant rate $b$ during their lives to children who can be mutants with probability $p$ or clones of their parents with probability $1-p$. We are working with an infinite alleles model, that is, when a mutation occurs, the allele of the mutant child was never encountered before. Moreover, mutations are neutral because they do not imply advantages or disadvantages (all individuals have identically distributed dynamics).

Without mutation, the model is linked to a genealogical tree called splitting tree [15, 16, 24]. Moreover, if $\Xi(t)$ denotes the number of alive individuals at time $t$, the process $\Xi := (\Xi(t), t \geq 0)$ is a Crump-Mode-Jagers process (or general branching process) [20] which is binary (births occur singly) and homogeneous (constant birth rate).

We are first interested in the allelic partition of the population and more precisely in properties of the frequency spectrum $(M_t^n, i \geq 1)$ where $M_t^n$ is the number of distinct alleles

\[ M_t^n := \text{number of distinct alleles at time } t. \]

\footnote{CMAP, Ecole Polytechnique, Route de Saclay, 91128 Palaiseau Cedex, France. mathieu.richard@cmap.polytechnique.fr}
carried by exactly $i$ individuals at time $t$ and that appeared after time $t - a$ (or equivalently the alleles whose ages are less than $a$ at time $t$). Roughly speaking, this spectrum classifies the different alleles depending on their ages and on their sizes.

This kind of question was first studied by Ewens [13] who discovered the well known 'sampling formula' named after him and which describes the law of the allelic partition for a Wright-Fisher model with neutral mutations. In our model, we can not get a counterpart of the Ewens’ sampling formula in the sense that we can not compute the joint law of $(M^{i,a}_t, i \geq 1)$ for fixed $0 < a \leq t$. However, we obtain two kinds of results concerning the frequency spectrum $(M^{i,a}_t, i \geq 1)$. We first compute the expected value of the number of alleles carried by $i$ individuals at time $t$ and with age $a$. When the process $\Xi$ is supercritical, we also obtain the asymptotic behaviors of the frequency spectrum and of the relative abundances of alleles as $t \to \infty$ on the survival event of $\Xi$.

Similar models to ours have been studied in the literature. In [17], Griffiths and Pakes study the case of a Bienaymé-Galton-Watson (BGW) process where children can independently be mutants with a given probability: the authors obtained asymptotic results about the number of alleles and the frequency spectrum at generation $n$ as $n \to \infty$. In [28], Pakes gets analogous properties concerning continuous-time Markov branching processes. In particular, his formula of the expected frequency spectrum can be seen as a counterpart of ours, which is stated in Section 3.1.

In the two articles [2, 3], Bertoin considers an infinite alleles model with neutral mutations in a subcritical or critical BGW-process where individuals independently give birth to a random number of clonal and mutant children according to the same joint distribution. In [2], he defines a tree of alleles where all individuals of a common allele are gathered in clusters and specifies the law of the allelic partition of the total population by describing the joint law of the sizes of the clusters and the numbers of their mutant children. In [3], Bertoin obtains the convergence of the sizes of allelic families for a large initial population and a small mutation rate.

Recent results have been obtained about splitting trees with mutations. In [10], Delaporte studies sequences of splitting trees with general and neutral mutations occurring at birth and investigates scaling limits in a regime with large population size and rare mutations. In two related papers [5, 6], Champagnat and Lambert consider a model really close to ours: the authors also work with splitting trees but in their model, individuals independently experience mutations at Poisson times during their lives. In [5], explicit formulas about the expected frequency spectrum at time $t$ are stated. In [6], the authors are interested in large and old families; they obtain asymptotic results about the sizes of the largest families and about the ages of the oldest ones as $t \to \infty$. Their model with Poissonian mutations and our model with mutations occurring at birth are compared in [7] in the particular case of exponential lifelengths.

Finally, Taïb [33] considers CMJ-processes with a more general kind of mutations at birth (for example, the probability that a mutation occurs can depend on the age of the mother) and thanks to the theory of CMJ-processes counted with characteristics, he obtains several asymptotic results, especially about the frequency spectrum of the population. However, in our particular case, some of the non-explicit limits he obtained can be computed thanks to a recent paper of Lambert [24] giving for example the one-dimensional marginals of $\Xi$ and its asymptotic behavior.

On the other hand, we obtain properties about the number of mutations undergone by
alleles. More specifically, for $i \geq 0$ and $t \geq 0$, we study $L_i(t)$ the number of alleles at time $t$ that have been affected by $i$ more mutations than the ancestral allele and $K_i(t)$ the number of individuals that carry such alleles. Our model can represent the spread of an epidemic: individuals are infected hosts, deaths are recoveries or actual deaths and births are transmissions of the disease, which can mutate to new strains. Then, $L_i(t)$ is simply the number of strains of the disease that are present at $t$ and that are at a distance $i$ from the original strain of the disease. Moreover, the process $K_i$ describes the number of individuals infected by such strains of the disease.

We compute the expected values of $L_i(t)$ and $K_i(t)$ and obtain asymptotic results about $K_i(t)$ as $t \to \infty$ by considering a multitype CMJ-process where the type of an allele is the number of mutations it has undergone. Multitype branching processes are also used in carcinogenesis, that is, in the evolution of cancerous cells. In [9, 12], cancerous cells are modeled by a multitype branching process where a cell is of type $k$ if it has undergone $k$ mutations and where the more a cell has undergone mutations, the faster it grows. The object of study is the time $\tau_k$ of appearance of the first cell of type $k$. Branching processes and birth and death processes appear in other works dealing with the evolution of cancerous cells (see for example [27, 19, 32] and references therein). For instance, in [32], the authors study the arrival time of the first resistant cell in a model of cancerous cells undergoing a medical treatment and becoming resistant after having experienced a certain number of mutations.

The paper is organized as follows: in Section 2, we expose the model that we consider and give some of its properties that will be useful to get our main results. Section 3 is devoted to the study of the frequency spectrum and in Section 4, we are interested in properties of the number of mutations undergone by alleles.

## 2 Preliminaries

### 2.1 The model

In this paper, as a population model, we consider genealogical trees satisfying the branching property and called splitting trees [15, 16, 24]. They are random trees satisfying the following assumptions.

- At time $t = 0$, there is only one progenitor.
- All individuals have i.i.d. lifespans and reproduction behaviors.
- Conditional on her birth date $\alpha$ and her lifespan $\zeta$, each individual gives birth to children at constant rate $b \in (0, \infty)$ during $(\alpha, \alpha + \zeta]$.
- Births occur singly.

We denote the common lifespan distribution by $\Lambda(\cdot)/b$ where $\Lambda$ is a finite positive measure on $(0, +\infty]$ with total mass $b$ and called lifespan measure [24].

To this splitting tree, we had neutral mutations occurring at birth in the following way. We fix $p \in (0,1)$. When a birth event occurs, independently of others individuals, with probability $1 - p$, a child is a clone of her mother, that is carries the same allele and a mutant with probability $p$. Moreover, when a mutant appears in the population, its type was never
carried before by any other individual. Thus, we consider an infinitely many alleles model with neutral mutations because all individuals behave in the same way regardless of allele.

On Figure 1, one can find an illustration of this model.

Figure 1: An example of a splitting tree with mutations up to time $t$. Vertical axis is time and horizontal axis shows filiation (lengths of dashed lined are meaningless). Full circles represent mutations occurring at birth and thick lines, the clonal splitting tree of the ancestor up to time $t$. The different letters are the alleles of alive individuals at time $t$.

Without mutation, if $\Xi(t)$ is the number of extant individuals at time $t$, then the process $(\Xi(t), t \geq 0)$ is a Crump-Mode-Jagers (CMJ) process or general branching process (see [20, 21] and references therein). We use the formalism commonly employed in CMJ-processes. One associates with each individual $x$ in the population a non-negative r.v. $\lambda_x$ (its life length), and a point process $\xi_x$ called birth point process. The sequence $(\lambda_x, \xi_x)$ is assumed to be i.i.d. but $\lambda_x$ and $\xi_x$ are not necessarily independent.

In our particular case, the common distribution of lifespans is $\Lambda(\cdot)/b$ and conditional on her lifespan, the birth point process of an individual is distributed as a Poisson point process during its life. If we denote by $\xi$ the birth point process of the ancestor, its characteristic measure is then given by

$$\mu(dt) := \mathbb{E}[\xi(dt)] = dt\Lambda((t, +\infty)).$$

(1)

In the same way, if $\xi_m$ (resp $\xi_c$) is the birth point process of the mutant (resp. clonal) children of the ancestor, then by the thinning property of Poisson point processes, $\xi_m$ and $\xi_c$ are independent, $\xi = \xi_m + \xi_c$ and with obvious notation, $\mu_m(dt) = p\mu(dt)$ and $\mu_c(dt) = (1-p)\mu(dt)$.

2.2 Basic properties about the process $\Xi$

We consider here the model without mutation and recall some known facts about the CMJ-processes $\Xi$, which will be useful in the following.
We say that the process $\Xi$ is subcritical, critical or supercritical if

$$
m := \int_{(0, \infty]} r \Lambda(dr)
$$

is respectively less than, equal to or greater than 1. We set

$$
\psi(\lambda) := \lambda - \int_{(0, \infty]} \left(1 - e^{-\lambda r}\right) \Lambda(dr).
$$

(2)

Since this function is convex and satisfies $\psi(0^+) = 0$, we can define $\eta$ as its largest root. Moreover, since $\psi'(0^+) = 1 - m$, when the process $\Xi$ is subcritical or critical, we have $\eta = 0$ and when it is supercritical, $\eta$ is positive. In the last case, $\eta$ is called the Malthusian parameter of the population as explained in the forthcoming Proposition 2.1.

To obtain properties about the splitting trees, Lambert defined in [24] a contour process which characterized them. This contour process is a spectrally positive Lévy process with Laplace exponent $\psi$. Let $W$ be the scale function associated with it (see [1, Ch. VII]), defined as the unique increasing continuous function $(0, \infty) \to (0, \infty)$ satisfying

$$
\int_{0}^{\infty} W(x)e^{-\lambda x}dx = \frac{1}{\psi(\lambda)}, \quad \lambda > \eta.
$$

(3)

In the entire paper, by convenience, we assume that the following hypothesis holds.

The measure $\Lambda$ has no atom. (H)

According to [23, p. 234], under the hypothesis (H), the function $W$ is continuously differentiable and thanks to the Lemma 4.1 in [31], we have

$$
W'(t) = bW(t) - \int_{0}^{t} W(t-x)\Lambda(dx), \quad t \geq 0.
$$

(4)

In fact, if (H) is not satisfied, most of the results stated in this paper still hold by replacing $W'(t)$ by $bW(t) - \int_{0}^{t} W(t-x)\Lambda(dx)$. We mainly choose to assume (H) in order to simplify the results and their proofs.

Thanks to the scale function $W$, we deduce the one-dimensional marginals of $\Xi$ (see [31, p. 293])

$$
P(\Xi(t) = 0) = 1 - \frac{W'(t)}{bW(t)}
$$

(5)

and for $n \in \mathbb{N}^*$,

$$
P(\Xi(t) = n) = \left(1 - \frac{1}{W(t)}\right)^{n-1} \frac{W'(t)}{bW(t)^2}.
$$

(6)

In other words, conditional on being positive, $\Xi(t)$ is geometric with success probability $1/W(t)$. In particular, for $t \geq 0$,

$$
E[\Xi(t)] = \frac{W'(t)}{b}.
$$

(7)

Let $\text{Ext} := \left\{\Xi(t) \xrightarrow{t \to \infty} 0\right\}$ be the extinction event of the splitting tree. We denote by $P^* := P(\cdot|\text{Ext}^c)$ the law of the process $\Xi$ conditional on its survival event.
Proposition 2.1 (Lambert [24]). We have
\[ P(\text{Ext}) = 1 - \eta/b. \]
Moreover, if \( \Xi \) is supercritical \((m > 1)\), under \( P^* \),
\[ e^{-\eta t} \Xi(t) \xrightarrow{t \to \infty} \mathcal{E} \quad \text{a.s.} \]  
where \( \mathcal{E} \) is exponential with parameter \( \psi'(\eta) \).

In fact, Lambert proved the convergence in distribution in [24] and a.s. convergence holds according to [26] (see [31, p.285]). The convergence (8) justifies why we call \( \eta \) the Malthusian parameter of the population since \( \Xi(t) \) grows like \( e^{\eta t} \) on the non-extinction event.

Most of the results stated in Sections 3 and 4 deal with long-time behaviors of several processes. To obtain them, we need the asymptotic behaviors of the scale function \( W \) and of its derivative \( W' \). Different regimes appear depending on whether \( \Xi \) is subcritical, critical or supercritical. We record them in the following lemma.

Lemma 2.2. (i) If \( m > 1 \) (supercritical case), as \( t \to \infty \), we have
\[ W(t) \sim \frac{1}{\psi''(\eta)} e^{\eta t} \quad \text{and} \quad W'(t) \sim \frac{\eta}{\psi'(\eta)} e^{\eta t}. \]
(ii) If \( m = 1 \) (critical case) and \( \sigma^2 := \int_0^\infty r^2 \Lambda(dr) < \infty \), we have \( W(t) \sim \frac{2}{\sigma^2} t \). If we also suppose that
\[ \lim_{t \to \infty} t^2 \int_t^\infty (x-t) \Lambda(dx) = 0, \]  
we have \( \lim_{t \to \infty} W'(t) = \frac{2}{\sigma^2} \).
(iii) If \( m < 1 \) (subcritical case), \( \lim_{t \to \infty} W(t) = \frac{1}{1 - m} \). If we also assume that there is a negative \( \tilde{\eta} \) satisfying
\[ \psi(\tilde{\eta}) = 0 \quad \text{and} \quad \int_0^\infty re^{-\tilde{\eta}r} \Lambda(dr) < \infty, \]  
then \( W'(t) \sim \frac{\tilde{\eta}}{\psi'(\tilde{\eta})} e^{\tilde{\eta} t} \).

In the critical case, the condition (9) holds if \( \Lambda \) has a finite third moment. Concerning the subcritical case, it is possible to find a negative root of \( \psi \) because this function is convex, \( \psi(0) = 0 \) and \( \psi'(0) = 1 - m > 0 \). However, to have this root requires that the exponential moment \( \int_0^\infty e^{\lambda r} \Lambda(dr) \) is finite for \( \lambda \) large enough. The proof of Lemma 2.2 is postponed to the appendix.
2.3 The clonal process

In the sequel, an important role will be played by the clonal process \( \Xi_c(t), t \geq 0 \) (c for clonal) where \( \Xi_c(t) \) is the number of extant individuals at time \( t \), carrying the same allele as the ancestor (see Figure 1). Since mutations occur independently of \( (\Xi(t), t \geq 0) \), by the thinning property of Poisson processes, the process \( \Xi_c \) defines a splitting tree with lifespan measure

\[ \Lambda_c := (1 - p)\Lambda. \]

Moreover, conditional on its arrival time in the population, each allele evolves like \( \Xi_c \) and independently of other alleles.

According to the previous section, the process \( \Xi_c \) is subcritical, critical or supercritical if \((1 - p)m\) is respectively less than, equal to, or greater than 1. In particular, when \( \Xi \) is critical or supercritical, \( \Xi_c \) is obviously subcritical.

Moreover, we associate with \( \Xi_c \) the function \( \psi_c \), which satisfies

\[ \psi_c(\lambda) := \lambda - (1 - p) \int_{(0,\infty)} \left( 1 - e^{-\lambda r} \right) \Lambda(dr) = p\lambda + (1 - p)\psi(\lambda). \]  

(11)

Let \( \eta_c \) be the largest root of \( \psi_c \). When \( \Xi_c \) is subcritical or critical, \( \eta_c = 0 \) while in the supercritical case, \( \eta_c > 0 \). Furthermore, in the latter, by the definition of \( \eta_c \) and by using (11), we have

\[ \psi(\eta_c) = \frac{p\eta_c}{p - 1} < 0. \]

It implies that \( \eta_c < \eta \) since \( \psi \) is convex and its largest root is \( \eta \).

Finally, all the properties about \( \Xi \), stated in the paragraph 2.2, also hold for \( \Xi_c \). To obtain them, it suffices to respectively replace \( b, \psi, \eta \) and \( W \) by \( b(1 - p) \), \( \psi_c \), \( \eta_c \) and \( W_c \) where \( W_c \) is the scale function associated with \( \psi_c \), solution of

\[ \int_0^\infty W_c(x)e^{-\lambda x}dx = \frac{1}{\psi_c(\lambda)}, \quad \lambda > \eta_c. \]

3 Frequency spectrum

At a given time, for any allele, we call family the set of all individuals that share this allele. To study the allelic partition of the population, we study its associated frequency spectrum, which, roughly speaking, sort the different families according to their sizes. More precisely, for \( i \in \mathbb{N}^* \) and \( a > 0 \), let \( M_i^a \) be the number of alleles whose ages are less than \( a \) and carried by \( i \) individuals at \( t \). Then, for fixed \( 0 \leq a < t \), the sequence \((M_i^a, i \geq 1)\) is the frequency spectrum at time \( t \) of families with ages less than \( a \).

Notice that \( M_i^{t,a} \) is simply the number of alleles carried by \( i \) particles at time \( t \) (regardless of their ages) and that \( M_t := \sum_{i \geq 1} M_i^{t,i} \) is the number of different alleles at time \( t \).

In the example on Figure 1, taking no account of ages of alleles, the frequency spectrum \((M_i^{t,i}, i \geq 1)\) is \((3, 2, 1, 0, \ldots)\) because three alleles \((B, E, F)\) are carried by one individual, \( A \) and \( D \) are carried by two individuals and \( C \) is the only allele carried by three individuals. If we are only interested in alleles younger than \( a \) at time \( t \), we have \((M_i^{t,a}, i \geq 1)\) = \((3, 1, 0, \ldots)\)
since alleles $A$ and $D$ appeared in the population before $t - a$.

Although it is not possible to obtain the joint distribution of $(M_i^{t,a}, i \geq 1)$ for fixed $t$ and $a$ as in the 'Ewens sampling formula', we are able to get some properties of this frequency spectrum.

### 3.1 Expected frequency spectrum

We first give an exact expression of the expected frequency spectrum at any time $t$. For $0 < a < t$ and $i \geq 1$, we denote by $M_i^{t,a}$ the number of alleles carried by $i$ individuals at time $t$ and with ages in $[a - da, a]$. The following proposition yields its expected value.

**Proposition 3.1.** For $0 < a < t$ and $i \geq 1$, we have

$$
E \left[ M_i^{t,a} \right] = \frac{p}{b(1-p)} W''(t - a) \left(1 - \frac{1}{W_a(t)}\right)^{i-1} \frac{W'_a(t)}{W_a^2(t)} da.
$$

**Proof.** Conditional on $\Xi(t - a)$, thanks to the branching property and classical properties of Poisson point processes, $M_i^{t,a}$ is the sum of $\Xi(t - a)$ i.i.d. r.v. distributed as the number of atoms in the interval $[t - a, t - a + da]$ of a Poisson point process with parameter $bpP(\Xi_c(a) = i)$. Hence,

$$
E \left[ M_i^{t,a} \right] = E[\Xi(t - a)] \frac{1}{b} P(\Xi_c(a) = i) da.
$$

According to (6) and (7), $E[\Xi(t - a)] = W'(t - a)/b$ and

$$
P(\Xi_c(a) = i) = \left(1 - \frac{1}{W_a(t)}\right)^{i-1} \frac{W'_a(t)}{b(1-p)W_a^2(t)}
$$

and we obtain the desired result. 

As a consequence of this proposition, we deduce the expected value of the frequency spectrum. In particular, when $\Xi$ is supercritical, we see that $M_i^{t,a}$ grows as $e^{\eta t}$ when $t \to \infty$, that is, with the same growth rate as $\Xi(t)$.

**Corollary 3.2.** For $a \leq t$ and $i \geq 1$,

$$
E \left[ M_i^{t,a} \right] = \frac{p}{b(1-p)} \int_0^a W''(t - x) \left(1 - \frac{1}{W_a(x)}\right)^{i-1} \frac{W'_a(x)}{W_a^2(x)} dx
$$

$$
+ \frac{1}{b(1-p)} \left(1 - \frac{1}{W_a(t)}\right)^{i-1} \frac{W'_a(t)}{W_a^2(t)} 1_{(t=a)}. \tag{13}
$$

Moreover, if $m > 1$, for $a \geq 0$, as $t \to \infty$,

$$
e^{-\eta t}E \left[ M_i^{t,a} \right] \to \eta \frac{p}{b(1-p)} J_i^{1,a} \tag{14}
$$

where

$$
J_i^{1,a} := \int_0^a e^{-\eta u} \left(1 - \frac{1}{W_a(u)}\right)^{i-1} \frac{W'_a(u)}{W_a^2(u)} du.
$$
In [28], Pakes obtained a similar result: in Lemma 3.1.2, he computed the expected frequency spectrum for a Markov branching process where children can independently be mutants of their mothers with probability \( p \). In [30, Chap 3], the expected frequency spectrum formula (13) is used to obtain the orders of magnitude of the sizes (resp. of the ages) of the largest families (resp. of the oldest families) at time \( t \) when \( t \to \infty \) (see also [7]).

**Proof of Corollary 3.2.** We obtain (13) by integrating (12) on \((0, a)\). The second term of the r.h.s. corresponds to \( \mathbb{P}(\Xi_c(t) = i) \), that is, the probability that the progenitor has \( i \) alive clonal descendants at time \( t \).

We get the convergence result (14) by using the dominated convergence theorem. First, as \( t \to \infty \), using the Lemma 2.2, \( W'(t-x) \) is equivalent to \( \eta \psi'(\eta) e^{\eta(t-x)} \). Then, we obtain the convergence of the integral because \( 1 - \frac{1}{W_c(x)} \) is bounded by \( b \) for \( t \) large enough. \( \square \)

### 3.2 Convergence results

In this paragraph, we suppose that the process \( \Xi \) is supercritical \((m > 1)\) and are interested in improvements of the convergence result (14).

The following proposition yields the asymptotic behavior, as \( t \to \infty \) and under \( \mathbb{P}^* \), of the frequency spectrum \((M_{t}^{i,a}, i \geq 1)\). We also obtain the convergence of the relative abundances \( M_{t}^{i,a}/M_{t} \) of families of sizes \( i \) and ages less than \( a \). We recall that \( \mathbb{P}^* \) is the law of the process \( \Xi \) conditional on its survival event.

**Proposition 3.3.** Under \( \mathbb{P}^* \), with probability 1

\[
e^{-\eta t} M_t \to \frac{p}{1-p} J \mathcal{E} \quad (15)
\]

\[
e^{-\eta t} M_t^{i,a} \to \frac{p}{1-p} J_t^{i,a} \mathcal{E} \quad (16)
\]

\[
\frac{M_t^{i,a}}{M_t} \to \frac{J_t^{i,a}}{J} \quad (17)
\]

where

\[
J := \int_0^\infty e^{-\eta u} \frac{W'_c(u)}{W_c(u)} du
\]

and where \( \mathcal{E} \) is the exponential r.v. with parameter \( \psi'(\eta) \) defined by (8).

Notice that (16) is consistent with (14) since \( \mathbb{P}(\text{Ext}^c) = \eta/b \) and \( \mathbb{E}[\mathcal{E}] = 1/\psi'(\eta) \). Moreover, we point that the relative abundances \( M_t^{i,a}/M_t \) converges to a deterministic limit.

Similar convergence results were obtained by Z. Taïb in [33] who considered a similar model with more general birth point processes and mutations. However, his convergence theorems give non explicit limits. In our particular case, we can find the distribution of the limits because of the knowledges of the one-dimensional marginals and of the asymptotic behavior of \( \Xi \) (see above-mentioned Proposition 2.1 and formulas (5) and (6)).

**Proof of Proposition 3.3.** To prove the three convergence results, we use general properties about CMJ-processes counted by random characteristics developed by Jagers and Nerman (see [20, 21, 22]). We mainly follow ideas of Taïb and all the properties we use are recorded in the appendix of [33]. We first check that the general assumptions (C.1-4) of this appendix hold in our case: recalling that \( \mu \) is the measure defined by (1), we must have
(a) \( \mu(\mathbb{R}^+) > 1 \) (supercritical case).

(b) There is a number \( \lambda_0 > 0 \) such that \( \int_0^\infty e^{-\lambda_0 u} \mu(du) = 1 \).

(c) \( \beta := \int_0^\infty u e^{-\lambda_0 u} \mu(du) < \infty \).

(d) \( \mu \) is not supported by any lattice \( k\mathbb{Z} \) for \( k > 0 \).

First, by the definition of \( \mu \), (a) is satisfied since we have supposed that \( m > 1 \) and point (d) is straightforward. Moreover, the largest root \( \eta \) of \( \psi \) satisfies (b) seeing that

\[
\int_0^\infty e^{-\lambda_0 u} \mu(du) = \int_0^\infty e^{-\lambda_0 u} \Lambda((u, +\infty]) du = \int_{(0,\infty]} \Lambda(dr) \frac{1 - e^{-\eta r}}{\eta} = \frac{\eta - \psi(\eta)}{\eta} = 1. \tag{18}
\]

Finally, let us check that (c) holds. Since \( \Lambda \) is a finite measure with mass \( b \), we have

\[
\beta = \int_0^\infty u e^{-\lambda_0 u} \Lambda((u, +\infty]) du \leq b \int_0^\infty u e^{-\lambda_0 u} du < \infty.
\]

We are now able to prove (16). We follow the proof of Theorem 3.3 in [33] using a random characteristic. We denote by \( \tau_1 < \tau_2 < \cdots \) the successive birth times of the children of the ancestor, by \( \rho_1, \rho_2, \ldots \) the independent sequence of i.i.d. Bernoulli r.v. with parameter \( 1 - p \) saying if a child is a clone or a mutant and by \( \Xi_1^c, \Xi_2^c, \ldots \) the i.i.d. clonal population processes of the mutant children. Then, for \( u \geq a \), we set the characteristic

\[
\chi(u) := \sum_{j \geq 1} (1 - \rho_j) 1_{\{u - a < \tau_j < u \}} 1_{\{\Xi_j^c(u - \tau_j) = i \}},
\]

that is, \( \chi(u) \) is the number of mutant children of the ancestor born between \( u - a \) and \( u \) and whose alleles are carried by \( i \) individuals at \( u \). For \( u \leq a \), we set \( \chi(u) = 0 \).

In the same way, for any individual \( k \) and \( u > a \), denote by \( \chi_k(u) \) the number of mutant children of \( k \), born between \( u - a \) and \( u \) units of time after the birth of \( k \) and whose alleles are carried by \( i \) individuals \( u \) units of time after the birth of \( k \). Hence, \( (M_{i,a}^t, t \geq 0) \) can be counted by the characteristic \( \chi \), meaning that for \( t \geq 0 \),

\[
M_{i,a}^t = \sum_{k=1}^y \chi_k(t - \alpha_k) + 1_{\{t \leq a \}} 1_{\{\Xi(t) = i \}} \tag{19}
\]

where \( \alpha_k \) denotes the birth time of \( k \) and \( y_t \) is the number of individuals born before \( t \). The second term in the r.h.s. of (19) pertains to the ancestor which is not counted by the characteristic. However, this term vanishes until \( t > a \) and so disappears at the limit \( t \to \infty \).

According to Theorem 5 in the appendix of [33], if there exists \( \eta' < \eta \) such that

\[
\int_0^\infty e^{-\eta' t} \mu(dt) < \infty \tag{20}
\]

\[
\mathbb{E} \left[ \sup_{u \geq 0} e^{-\eta' u} \chi(u) \right] < \infty, \tag{21}
\]

then, under \( \mathbb{P}^* \),

\[
\lim_{t \to \infty} \frac{M_{i,a}^t}{y_t} = \int_0^\infty \eta e^{-\eta u} \mathbb{E} [\chi(u)] du \quad \text{a.s.}
\]
However, conditioning by the point process $\xi_m$ which is independent from the $\Xi^*_t$'s, we have

$$
\int_0^\infty \eta e^{-\eta u} \mathbb{E} \left[ \chi(u) \right] \, du = \int_0^\infty \eta e^{-\eta u} \, du \mathbb{E} \left[ \sum_{j \geq 1} (1 - \rho_j) \mathbf{1}_{\{u < \tau_j \}} \mathbf{1}_{\{\Xi^*_t(u) = j\}} \right]
$$

$$
= \int_0^\infty \eta e^{-\eta u} \, du \mathbb{E} \left[ \int_{u-a}^u \mathbb{P}(\Xi_c(u - t) = i) \, d\xi_m(dt) \right]
$$

$$
= \int_0^\infty \eta e^{-\eta u} \, du \int_{u-a}^u \mathbb{P}(\Xi_c(u - t) = i) \, d\mu_m(dt)
$$

$$
= \int_0^\infty \mu_m(dt) e^{-\eta t} \int_0^a \eta e^{-\eta u} \mathbb{P}(\Xi_c(u) = i) \, du
$$

using Fubini-Tonelli's theorem and a change of variables. Since $\mu_m = p\mu$ and $\eta$ satisfies (18), $\int_0^\infty e^{-\eta t} \mu_m(dt) = p$. Then, thanks to (6),

$$
\lim_{t \to \infty} \frac{M_t^{i,a}}{y_t} = \frac{pn}{b(1-p)} \int_0^{\eta} e^{-\eta u} \left(1 - \frac{1}{W_c(u)}\right)^{i-1} \frac{W'_c(u)}{W^2_c(u)} \, du \quad \text{a.s.} \quad (22)
$$

We still have to find $\eta' < \eta$ such that conditions (20) and (21) are satisfied. First, as in (18), for $\eta' \in (0, \eta)$, $\int_0^\infty e^{-\eta' u} \mu(dt) = 1 - \psi(\eta')/\eta' < \infty$. Second, the characteristic $\chi$ is stochastically dominated by a Poisson process with parameter $b$, say $(N_t, t \geq 0)$. Then, to prove (21), it is sufficient to show that $\mathbb{E} \left[ \sup_{t \geq 0} e^{-\eta't} N_t \right]$ is finite. At the end of the proof of Proposition 5.1 in [5, p.1028], the authors show that for $\kappa$ large enough, the process $A := \left(e^{-\eta't} N_t + \kappa\right)^2$ is a supermartingale, which implies that for $y \geq 0$,

$$
\mathbb{P} \left( \sup_{t \geq 0} e^{-\eta't} N_t \geq y \right) = \mathbb{P} \left( \sup_{t \geq 0} A_t \geq (y + \kappa)^2 \right) \leq \frac{\mathbb{E}[A_0]}{(y + \kappa)^2} = \frac{\kappa}{(y + \kappa)^2},
$$

where the last inequality holds thanks to [29, p. 58]. Then,

$$
\mathbb{E} \left[ \sup_{t \geq 0} e^{-\eta't} N_t \right] = \int_0^\infty dy \mathbb{P} \left( \sup_{t \geq 0} e^{-\eta't} N_t \geq y \right) < \infty.
$$

Furthermore, using again [33, Thm 5] with the characteristic $\chi'(u) = \mathbf{1}_{\{0 \leq u \leq \lambda\}}$, on $\mathbb{P}^*$

$$
\frac{\Xi_t}{yt} = \sum_{k=1}^{\lambda t} \frac{\chi'_k(t - \alpha_k)}{yt} \int_0^\infty \eta e^{-\eta x} \mathbb{E} \left[ \chi'(x) \right] \, dx = \int_0^\infty \eta e^{-\eta x} \, dx \int_{(x, \infty)} \frac{\Lambda(dx)}{b} = \frac{\eta}{b} \quad (23)
$$

because $\psi(\eta) = 0$. Notice that in that case, condition (21) is easily satisfied because $\chi'(u) \leq 1$ for $u \geq 0$. Finally, using together (8), (22) and (23) we get the convergence result (16).

We now prove (15). According to [33, Thm 3.3], on $\mathbb{P}^*$,

$$
\frac{M_t}{y_t} \to \int_0^\infty e^{-\eta x} \mu_m(dx) \left(1 - \eta \int_0^\infty e^{-\eta u} \mathbb{P}(\Xi_c(u) = 0) \, du\right) \, du = pn \int_0^\infty e^{-\eta u} \frac{W'_c(u)}{b(1-p)W_c(u)} \, du.
$$

By using the last display, (8) and (23), on $\mathbb{P}^*$, we have

$$
e^{-\eta t} M_t \to \mathcal{E} \frac{p}{1-p} \int_0^\infty e^{-\eta u} \frac{W'_c(u)}{W_c(u)} \, du.
$$

Finally, (17) is straightforward from (15) and (16).
The following result deals with the asymptotic behavior of $M_{i,t}^{t}$, the number of alleles carried by $i$ individuals at time $t$.

**Proposition 3.4.** Under $\mathbb{P}^*$, with probability 1 as $t \to \infty$,

$$
\frac{M_{i,t}^{t}}{M_t} \to \frac{1}{i} \int_0^\infty e^{-\eta u} \left(1 - \frac{1}{W_c(u)}\right)^i du.
$$

Notice that the limit can be considered as a kind of Fisher log-series distribution, which is commonly used to describe species abundances [14].

**Proof.** Since $M_{i,t}^{t}$ can be counted by the random characteristic

$$
\bar{\chi}(u) := \sum_{j \geq 1} (1 - \rho_j) 1_{\{\tau_j < u\}} 1_{\{\Xi_j < u\}},
$$

similar computations as in the proof of Proposition 3.3 lead to the convergence of $M_{i,t}^{t}/M_t$ to $J_i/\infty$ as $t \to \infty$. We then get the result by integrating by parts $J_i$ and $J_i^\infty$ by noticing that $W_c(0) = 1$ according to [31, Lem 4.1].

**Remark 3.5.** In most cases, that is not possible to compute the integrals $J$ and $J_{i,a}$ appearing in the limits in Propositions 3.3 and 3.4 since the scale functions $W$ and $W_c$ are generally unknown. However, even if $W_c$ is computable, it is not easier: when the life lengths are exponentially distributed with parameter $d$ (in that case, $\Xi$ is simply a Markovian birth and death process with birth rate $b$ and death rate $d$), we have from [24, p393]

$$
W_c(x) = \left\{ \begin{array}{ll}
\frac{d - (1 - p)b e^{(1 - p)b - d}x}{d - (1 - p)b} & \text{if } d \neq (1 - p)b \\
1 + dx & \text{if } d = (1 - p)b
\end{array} \right. \quad (24)
$$

and integrals are still uncomputable. It seems that we can only compute them in the pure death case: suppose that $\Lambda(\cdot)/b = \delta_\infty$. Hence, each individual lives an infinite time and gives birth at rate $b$: $\Xi$ is then a Markovian pure birth process and direct computations give $\psi(\lambda) = (\lambda - b) 1_{\{\lambda > 0\}}$, $\eta = b$ and from (24), $W_c(x) = e^{(1-p)bx}$. Obviously, in that case, $\mathbb{P}(\text{Ext}) = 0$ and $\mathbb{P}^* = \mathbb{P}$. Moreover, easy computations lead to simple expressions of $J$ and $J_{i,a}$ (for instance, the first one equals $1 - p$) and we obtain convergence results as

$$
\frac{M_{i,t}^{t}}{M_t} \to \int_0^\infty \frac{1}{i(1 - p)} \left(1 - \frac{1}{i(1 - p)}\right)^k \frac{1}{1 + (1 - p)k} \quad \text{as.}
$$

### 4 Number of undergone mutations

Let us say that the allele of the progenitor is of kind $0$. Then, recursively, for $k \geq 1$, we say that an allele is of kind $k$ if it is carried by a mutant child of an individual carrying an allele of kind $k - 1$. Equivalently, they are alleles that have been affected by $k$ more mutations than the ancestral allele.

For $t \geq 0$ and $i \geq 0$, we denote by $L_i(t)$ the number of alleles of kind $i$ at time $t$ and by $K_i(t)$ the number of individuals that carry such alleles at time $t$. 
In this section, we obtain the same kind of results as those of Section 3, that is, we compute the expected values $E[K_i(t)]$ and $E[L_i(t)]$ for fixed $i$ and $t$. We then study the asymptotic behavior of $K_i(t)$ as $t \to \infty$ when the clonal process is supercritical.

The main argument we use to get these results is to consider $(K_i(t), i \geq 0, t \geq 0)$ as a multitype CMJ-process where the types are the kinds of the different alleles. More precisely, regardless of its kind, every individual has a lifespan distributed as $\Lambda(\cdot)/b$ and an individual with kind $i$ gives birth to individuals of the $i$-th kind at rate $b(1-p)$ and to individuals of kind $i+1$ at rate $bp$. Moreover, the process $(K_i, i \geq 1)$ belongs to the class of reducible multitype processes since an individual of kind $i$ can not give birth to an individual of kind $j$ with $j < i$.

Then, if at some time, there is no more individuals of kind $i$, no such individual will reappear in the population.

Even though the reducible branching processes are less studied than the irreducible processes (which enables the use of Perron-Frobenius theory of positive matrices), they are useful in the model we consider. Indeed, if we are interested in the study of the process $K_i$ for a given $i$, since the kinds can not decrease, we can transform a problem with an infinite and countable number of types to a problem with a finite number of possible types by studying the multitype process $(K_j, j = 1, \ldots, i)$.

The reader interested in multitype CMJ-processes can refer to the third and fourth chapters of [25].

### 4.1 Expected number of mutations

For $f, g$ two continuous functions on $[0, +\infty)$, we denote by $f \ast g$ the standard convolution product

$$f \ast g(t) = \int_0^t f(t-x)g(x)dx, \quad t \geq 0.$$  

We use the following notation for the consecutive convolutions of a function. For a continuous function $f$, let $f^{*(1)} := f$ and for $i \geq 2$, $f^{*(i)} := f \ast f^{*(i-1)}$. In the following result, we give simple expressions of the mean values of $K_i(t)$ and $L_i(t)$ that are entirely determined by the functions $W_c$ and $W'_c$.

**Proposition 4.1.** We fix $t > 0$. We then have

$$E[K_i(t)] = \frac{1}{b(1-p)} \left( \frac{p}{1-p} \right)^i (W'_c)^{(i+1)}(t), \quad i \geq 0$$  

$$E[L_0(t)] = \frac{1}{b(1-p)} \frac{W'_c(t)}{W_c(t)}$$  

$$E[L_i(t)] = \frac{1}{b(1-p)} \left( \frac{p}{1-p} \right)^i (W'_c)^{(i)} \ast \frac{W'_c(t)}{W_c(t)}, \quad i \geq 1.$$  

In [25], C. Mode considers general multitype branching processes. In paragraph 4.4, he gives an equation satisfied by the Laplace transform of $E[K_i(\cdot)]$. In most cases, this equation can not be solved but it enables to obtain the asymptotic behavior of $E[K_i(t)]$ as $t \to \infty$. In our particular case, one can solve the equation of Mode to obtain the Laplace transform of $E[K_i(\cdot)]$ and compute $E[K_i(t)]$ thanks to an inverse Laplace transform. Nevertheless, in the following proof, we prefer to use a more direct approach which avoids long computations.
Proof of Proposition 4.1. We prove (25) by induction on \( i \geq 0 \). For \( i = 0 \), according to (7), \( \mathbb{E}[K_0(t)] = \mathbb{E}[\xi_c(t)] = \frac{W_c(t)}{b(1 - p)} \). We now suppose that the result is true at rank \( i - 1 \). Denote by \( K_i^{da}(t) \) the number of individuals of kind \( i \) that are alive at time \( t \) and whose alleles appear in the population between times \( t - a \) and \( t - a + da \). Then, in the same manner as in proof of Proposition 3.1, using the branching property, conditional on \( K_{i-1}(t - a) = n \), \( K_i^{da}(t) \) is the sum of \( n \) independent r.v. These r.v. are distributed as the number at time \( a \), of all the clonal descendants of mutants individuals born in a time interval \([0, da]\). Then,

\[
\mathbb{E}[K_i^{da}(t)] = \mathbb{E}[K_{i-1}(t - a)] \mathbb{E}[\xi_c(a)] bpd\alpha.
\]

Finally, using the induction hypothesis and integrating over \( a \in (0, t) \), we have

\[
\mathbb{E}[K_i^{da}(t)] = \frac{1}{b(1 - p)} \left( \frac{p}{1 - p} \right)^{i-1} \int_0^t (W_c^a)^{(i)}(t - a) \frac{W_c^a(a)}{b(1 - p)} bpd\alpha,
\]

which is what we wanted to prove.

Since \( L_0(t) = 1_{[\xi_c(t) > 0]} \), (26) is straightforward from (5). We prove (27) with similar techniques as we proved (25). The mean number of alleles of kind \( i \) with age \( a \) at time \( t \) is

\[
\mathbb{E}[L_i^{da}(t)] = \mathbb{E}[K_{i-1}(t - a)] \mathbb{E}[\xi_c(a) > 0] \mathbb{E}(\xi_c > 0) da.
\]

We then get the results thanks to (5), (25) and by integrating the last display on \([0, t]\). \( \square \)

From Proposition 4.1, we obtain the asymptotic behaviors of \( \mathbb{E}[K_i(t)] \) and \( \mathbb{E}[L_i(t)] \) as \( t \to \infty \). Different regimes appear depending on the class of criticality of the clonal process \( \xi_c \).

**Corollary 4.2.** We suppose that one of the following hypotheses holds

- \((1 - p)m > 1\),
- \((1 - p)m = 1\), \(\sigma^2 < \infty\) and (9) is satisfied,
- \((1 - p)m < 1\) and there is \( \tilde{\eta}_c \) satisfying (10).

Then, for \( i \geq 0 \), we have

\[
\lim_{t \to \infty} \mathbb{E}[K_i(t)] = \frac{1}{b(1 - p)i!} \left( \frac{p}{1 - p} \right)^i C_p^{i+1} t^i e^{\eta_p t}
\]

where

\[
\eta_p := \begin{cases} 
\eta_c & \text{if } (1 - p)m > 1 \\
0 & \text{if } (1 - p)m = 1 \\
\tilde{\eta}_c & \text{if } (1 - p)m < 1
\end{cases}
\quad \text{and } C_p := \begin{cases} 
\frac{\eta_c}{\psi_c(\eta_c)} & \text{if } (1 - p)m > 1 \\
\frac{1}{2} & \text{if } (1 - p)m = 1 \\
\frac{\eta_c}{\psi_c(\tilde{\eta}_c)} & \text{if } (1 - p)m < 1
\end{cases}.
\]

**Proof.** This result is a consequence of the following lemma thanks to Proposition 4.1 and Lemma 2.2 which gives the asymptotic behaviors of \( W_c^a(t) \) as \( t \to \infty \) in all cases. \( \square \)
Lemma 4.3. Let $f$ be a non-negative continuous function on $[0, +\infty)$ such that for some $a \in \mathbb{R}$ and $l \geq 0$,
\[ e^{at} f(t) \to l \quad (28) \]
Then, for $i \geq 1$
\[ \frac{e^{at}}{i-1} f^{(i)}(t) \to \frac{l^{i}}{(i-1)!} \quad (29) \]
Proof. We prove the result by induction on $i \geq 1$. It is obvious if $i = 1$ since we assume (28).
We now suppose that (29) holds for an integer $i \geq 1$. By a change of variables, we have
\[ f^{(i+1)}(t) = \int_{0}^{t} f(t-x) f^{(i)}(x) \, dx, \]
Hence,
\[ t^{-i} e^{at} f^{(i+1)}(t) = \int_{0}^{1} \left( e^{at(1-x)} f(t(1-x)) \right) \left( \frac{e^{ax}}{(tx)^i - t} f^{(i)}(tx) \right) x^{i-1} \, dx. \]
As $t \to \infty$, the r.h.s. of the last display converges to $l \frac{l^{i}}{(i-1)!} \int_{0}^{1} x^{i-1} \, dx = \frac{l^{i+1}}{i!}$ by using (28), the recurrence hypothesis and the dominated convergence theorem.

We end this paragraph by giving the asymptotic behavior of $E[L_i(t)]$ as $t \to \infty$.

Corollary 4.4. Let $i$ be a positive integer.
(i) If $(1-p)m > 1$, we have
\[ E[L_i(t)] \sim \frac{J_c}{b(1-p)(i-1)!} \left( \frac{p}{1-p} \right)^i \left( \frac{\eta_c}{\psi'(\eta_c)} \right)^i t^{i-1} e^{\eta_c t} \]
where $J_c = \int_{0}^{\infty} e^{-\eta_c u} W'_c(u) W_c(u) \, du$.
(ii) If $(1-p)m = 1$, if $\sigma^2 < \infty$ and if (9) holds,
\[ E[L_i(t)] \sim C_i t^i \ln t \]
where $C_i = \frac{1}{b(1-p)(i-1)!} \left( \frac{2p}{\sigma^2(1-p)} \right)^i$.
(iii) If $(1-p)m < 1$ and if there is $\tilde{\eta}_c$ satisfying (10),
\[ E[L_i(t)] \sim \frac{1 - (1-p)m}{b(1-p)i!} \left( \frac{p}{1-p} \right)^i \left( \frac{\tilde{\eta}_c}{\psi'(\tilde{\eta}_c)} \right)^i t^{i+1} e^{\tilde{\eta}_c t} \]
Proof. We begin by proving the subcritical case. According to Lemma 2.2(iii), $W'_c(t) W_c(t)$ behaves as $(1-(1-p)m) \frac{\tilde{\eta}_c}{\psi'(\tilde{\eta}_c)} e^{\tilde{\eta}_c t}$ as $t \to \infty$. Then point (iii) stems from a slightly modification of Lemma 4.3.
Concerning the critical case, according to Lemma 4.3, for any \( \varepsilon > 0 \), there is \( A > 0 \) such that, if \( t - x \geq A \), then

\[
\left| \frac{(W_c')^{(i)}(t - x)}{t^i} - \frac{1}{(i-1)!} \left( \frac{2}{(1-p)\sigma^2} \right) t^i \right| :\! := A_i
\]

Then, for \( \varepsilon > 0 \) and \( t \geq A \), we have

\[
t^{-i}(W_c')^{(i)} \star \frac{W_c'}{W_c}(t) = I_1(t) + I_2(t)
\]

where

\[
I_1(t) := \int_0^{t-A} t^{-i}(W_c')^{(i)}(t-x) \frac{W_c'(x)}{W_c(x)} dx \in \left[ (A_i - \varepsilon) \int_0^{t-A} \frac{W_c'(x)}{W_c(x)} dx, (A_i + \varepsilon) \int_0^{t-A} \frac{W_c'(x)}{W_c(x)} dx \right]
\]

and

\[
I_2(t) := \int_{t-A}^{t} t^{-i}(W_c')^{(i)}(t-x) \frac{W_c'(t-x)}{W_c(t-x)} dx \leq t^{-i}b(1-p) \int_0^{t} (W_c')^{(i)}(x) dx
\]

where we have used that \( \frac{W_c'(x)}{W_c(x)} = b(1-p)\mathbb{P}(\Xi_c(t) > 0) \). Hence, as \( t \to \infty \), \( I_2(t) \) vanishes and \( I_1(t) \) is equivalent to \( A_i \ln(t) \) since we know from Lemma 2.2(ii) that \( W_c'(x)/W_c(x) \sim 1/x \) as \( x \to \infty \). Thus,

\[
\mathbb{E}[L_i(t)] = \frac{1}{b(1-p)} \left( \frac{p}{1-p} \right)^i (W_c')^{(i)} \star \frac{W_c'}{W_c}(t) \sim \frac{A_i}{b(1-p)} \left( \frac{p}{1-p} \right)^i t^i \ln(t).
\]

Finally, we get the asymptotic behavior in the supercritical case by a direct application of the following Lemma 4.5. Indeed, according to the proof of Lemma 4.4, \( (W_c')^{(i)}(t) \) grows as \( t^i e^{\eta_c t} \) as \( t \to \infty \) and from Lemma 2.2 (i), we have \( \frac{W_c(t)}{W_c(t)} \to \eta_c \) as \( t \to \infty \). \( \square \)

**Lemma 4.5.** Let \( f, g \) be two continuous and positive functions such that \( f(t) \) converges as \( t \to \infty \) and such that for some \( a > 0 \),

\[
\lim_{t \to \infty} \frac{e^{-at}}{t^i} g(t) = l'.
\]

Then, as \( t \to \infty \),

\[
f \ast g(t) \sim l' e^{\eta_c t} \int_0^\infty e^{-ax} f(x) dx.
\]

**Proof.** We have

\[
e^{-at}/t^i f \ast g(t) = \int_0^\infty e^{-ax} f(x) h(t, x) dx
\]

where \( h(t, x) := \frac{e^{-a(t-x)}g(t-x)}{p} 1_{0 \leq x \leq t} \). Moreover, \( h(t, x) \to l' 1_{x \geq 0} \) as \( t \to \infty \) and according to (30)

\[
h(t, x) \leq \frac{e^{-a(t-x)}g(t-x)}{(t-x)^i} 1_{x \leq t} \leq M
\]

where \( M \) is a constant that does not depend on \( t \) and \( x \). Finally, since \( f(t) \) converges as \( t \to \infty \), \( \int_0^\infty f(x)e^{-ax} dx \) is finite and the result stems from the dominated convergence theorem. \( \square \)
4.2 Asymptotic behavior of \( K_i(t) \) when \( \Xi_c \) is supercritical

In Corollary 4.2, we proved that \( \mathbb{E}[K_i(t)] \) grows as \( t^i e^{\eta c t} \) as \( t \to \infty \) when the clonal process is supercritical. We now improve that convergence result by studying the asymptotic behavior of \( K_i(t) \). Notice that in the subcritical and critical cases, the process \( K_i \) goes extincted a.s.

**Theorem 4.6.** We suppose that \((1-p)m > 1\) and that \( \sigma^2 := \int_0^\infty \Lambda(dz)z^2 < \infty \). Then, for \( i \geq 0 \), almost surely and in quadratic mean, we have

\[
e^{-\eta c t} t^i K_i(t) \xrightarrow{t \to \infty} \kappa_i
\]

where \( \mathbb{P}(\kappa_i = 0) = 1 - \frac{\eta}{b(1-p)} \) and conditional on being non zero, \( \kappa_i \) is exponential with mean

\[
\frac{1}{i!} \left( \frac{p}{1-p} \right)^i \frac{\eta_i}{\psi_c'(\eta_c)^{i+1}}.
\]

**Proof.** The case \( i = 0 \) is straightforward since \( K_0 = \Xi_c \) and the result holds according to Proposition 2.1. Then, from now on, we suppose that \( i \geq 1 \).

The proof is divided in four steps. In the first one, we prove that the a.s. and \( L^2 \) convergences hold. We then identify the law of the limit: we find an equation satisfied by its Laplace transform (step 2), exhibit one solution of this equation (step 3) and show that it is the only one (step 4).

**Step 1: Proof of the convergence**

To prove the a.s. and quadratic mean convergences toward \( \kappa_i \) as \( t \to \infty \), we use Theorem 4.3 (iv) in [25, p.173]. Actually, in this book, C. Mode only proved that \( t^{-1} e^{-\eta c t} K_1(t) \) converges as \( t \to \infty \) but the same result holds about \( K_i \) for \( i \geq 2 \) by using similar techniques.

Let us check the hypotheses of Theorem 4.3 of [25]. First, we have to show that \( N \), the number of children of the ancestor, is square integrable. Conditional on having a life length \( z \), \( N \) is a Poisson r.v. with parameter \( bz \). Then,

\[
\mathbb{E}[N^2] = b^{-1} \int_{0}^{\infty} \Lambda(dz)((bz)^2 + bz) = b\sigma^2 + m < \infty.
\]

Second, for \( i, j \geq 0 \), let \( \mu_{i,j}(t) \) be the mean number of children of kind \( j \) that has an immortal individual of kind \( i \) before he reaches age \( t \). We have to prove that \( \max_{i, j} \mu_{i,j}(t) \) is null when \( t = 0 \) and finite when \( t > 0 \). In our case, it is obvious since \( \mu_{i,i}(t) = b(1-p)t \) if \( j = i \), \( \mu_{i,j}(t) = bpt \) if \( j = i+1 \) and is null in other cases.

Finally, if \( \zeta \) is the life length of an individual, we have to check that \( \int_0^\infty dt \mathbb{P}(\zeta \geq t)^p \) is finite for \( p = 1 \) and also for some \( p > 1 \). We have

\[
\int_0^\infty dt \mathbb{P}(\zeta \geq t)^p \leq \int_0^\infty dt \mathbb{P}(\zeta \geq t) = \mathbb{E}[\zeta] = \frac{m}{b} < \infty.
\]

**Step 2: An equation satisfied by the Laplace transform of \( \kappa_i \)**

The aim of this step is to prove that for \( i \geq 0 \) and \( a > 0 \), we have

\[
\varphi_i(a) = \int_0^\infty \frac{\Lambda(dz)}{b} \exp \left\{ b(1-p) \left( \int_0^z \varphi_i(ae^{-\eta c u})du - z \right) \right\}.
\]
where for $i \geq 0$, $\varphi_i$ denotes the Laplace transform of $\kappa_i$

$$\varphi_i(a) := \mathbb{E}\left[e^{-a\kappa_i}\right], \quad a > 0.$$  

To show that (31) holds, we look at the children of the ancestor. More precisely, conditional on having a lifespan $z$, before time $t$, the ancestor has a number of mutant (resp. clonal) children distributed as a Poisson r.v. with parameter $bp(z \wedge t)$ (resp. $b(1-p)(z \wedge t)$). Moreover, these two r.v. are independent.

For $z > 0$, $t \geq 0$ and $k, l \in \mathbb{N}$, by the branching property and classical properties about Poisson processes, conditional on the event “during its lifespan $z$ and before age $t$, the ancestor has $k$ mutant and $l$ clonal children”, we have

$$K_i(t) \overset{d}{=} \sum_{q=1}^{k} \tilde{K}_i^q(t-U_q) + \sum_{r=1}^{l} \tilde{K}_i^r(t-V_r)$$

where

- the processes $\tilde{K}_i^1, \ldots, \tilde{K}_i^k$ are i.i.d. and distributed as $K_{i-1}$, 
- the processes $\tilde{K}_i^1, \ldots, \tilde{K}_i^k$ are i.i.d. and distributed as $K_i$, 
- the r.v. $U_1, \ldots, U_p, V_1, \ldots, V_q$ are i.i.d. and uniform in $[0, z \wedge t]$.

Moreover, all these quantities are mutually independent. Hence, for $s \in [0, 1]$, we have

$$\mathbb{E}\left[s^{K_i(t)}\right] = \int_0^\infty \frac{\Lambda(dz)}{b} \sum_{k,l \geq 0} e^{-bz\Lambda} \frac{(bp(z \wedge t))^k}{k!} \frac{(b(1-p)(z \wedge t))^l}{l!} \mathbb{E}\left[s^{K_{i-1}(t-U_1)}\right]^k \mathbb{E}\left[s^{K_i(t-V_1)}\right]^l$$

$$= \int_0^\infty \frac{\Lambda(dz)}{b} \exp\left\{ -bz\wedge t + pb(z \wedge t) \mathbb{E}\left[s^{K_{i-1}(t-U_1)}\right] + (1-p)b(z \wedge t)\mathbb{E}\left[s^{K_i(t-V_1)}\right] \right\}. $$

Thus, if we denote by $G_{i,t}$ the generating function of $K_i(t)$, we have for $s \in [0, 1]$

$$G_{i,t}(s) = \int_0^\infty \frac{\Lambda(dz)}{b} \exp\left\{ -bz\wedge t + pb \int_0^s g_{i-1,t-u}(s)du + (1-p)b \int_0^s g_{i,t-v}(s)dv \right\}. $$

(32)

According to the first step of the proof, as $t \to \infty$, $t^{-i}e^{-\eta_i t}K_i(t)$ converges a.s. and then in distribution to $\kappa_i$. Then,

$$G_{i,t}\left(e^{-at^{-i}e^{-\eta_i t}}\right) = \mathbb{E}\left[e^{-aK_i(t)t^{-i}e^{-\eta_i t}}\right] \underset{t \to \infty}{\longrightarrow} \varphi_i(a),$$

$$G_{i-1,t-u}\left(e^{-at^{-i}e^{-\eta_i t}}\right) \underset{t \to \infty}{\longrightarrow} 1, \quad (33)$$

$$G_{i,t-v}\left(e^{-at^{-i}e^{-\eta_i t}}\right) \underset{t \to \infty}{\longrightarrow} \varphi_i(ae^{-\eta_i v}). \quad (34)$$

We set

$$C(t, z) := -b(z \wedge t) + pb \int_0^s g_{i-1,t-u}(e^{-at^{-i}e^{-\eta_i t}})du + (1-p)b \int_0^s g_{i,t-v}(e^{-at^{-i}e^{-\eta_i t}})dv.$$

18
Then, since for all $t \geq 0$ and $i \in \mathbb{N}$, $G_{i,t}$ and $G_{i-1,t}$ are less than 1, using the dominated convergence theorem and equations (33) and (34), we get

$$\lim_{t \to \infty} C(t, z) = -b(1-p)z + (1-p)b \int_0^z \varphi_i(ae^{-\eta_c v}) dv$$

Finally, if we apply (32) with $s = e^{-at-i\eta_c t}$ and let $t$ go to $\infty$, we obtain (31). Indeed, we can again use the dominated convergence theorem since $C(t, z) \leq 0$ for $t \geq 0$ and $z > 0$ and since $\Lambda$ is a finite measure.

**Step 3: Existence of a solution of (31)**

If $P := \frac{\eta_c}{b(1-p)}$, let $X$ be a non-negative r.v. such that $P(X = 0) = 1 - P$ and conditional on being positive, $X$ is exponential with parameter $c$. We prove in this paragraph that the Laplace transform of $X$, that is

$$\varphi_X : a \mapsto 1 - P + P \frac{e^c}{a + c},$$

is a solution of (31). We have

$$\int_0^z \varphi_X(ae^{-\eta_c u}) du = z(1-P) + P \int_0^z \frac{ce^{\eta_c u}}{a + ce^{\eta_c u}} du$$

$$= z(1-P) + \frac{P}{\eta_c} (\ln(a + ce^{\eta_c z}) - \ln(a+c))$$

Then,

$$\int_0^\infty \frac{\Lambda(dz)}{b} \exp\left\{ b(1-p) \left( \int_0^z \varphi_X(ae^{-\eta_c u}) du - z \right) \right\} = \int_0^\infty \frac{\Lambda(dz)}{b} e^{-\eta_c z} \frac{a + ce^{\eta_c z}}{a + c}$$

$$= \frac{c}{a + c} + \frac{a}{a + c} \int_0^\infty \frac{\Lambda(dz)}{b} e^{-\eta_c z}.$$

Since $\eta_c$ is a root of the function $\psi_c$ defined by (11), we have $(1-p) \int_0^\infty \Lambda(dz)e^{-\eta_c z} = b(1-p) - \eta_c$. Then,

$$\int_0^\infty \frac{\Lambda(dz)}{b} \exp\left\{ b(1-p) \left( \int_0^z \varphi_X(ae^{-\eta_c u}) du - z \right) \right\} = \frac{c}{a + c} + \frac{a}{a + c} (1-P) = \varphi_X(a).$$

**Step 4: Unicity of the equation (31)**

We prove here that if a non-negative r.v. has a positive and finite mean $\mu := \mathbb{E}[Y]$ and if its Laplace transform satisfies (31), then

$$\mathbb{P}(Y > 0) = \frac{\eta_c}{b(1-p)},$$

and conditional on being positive, $Y$ is exponential with mean

$$\mu \frac{b(1-p)}{\eta_c}.$$
If we show this, the proof of Theorem 4.6 is complete. Indeed, thanks to the Step 1, the convergence holds in quadratic mean, which implies that

$$\mathbb{E}[\kappa_i] = \frac{1}{b(1-p)} \int \frac{1}{i!} \left( \frac{p}{1-p} \right)^i \left( \frac{\eta_c}{\psi_c(\eta_c)} \right)^{i+1}$$

according to Corollary 4.2.

Let $Y$ be a r.v. with mean $\mu$ and satisfying

$$\varphi_Y(a) = \int_0^\infty \frac{\Lambda(dz)}{b} \exp \left\{ b(1-p) \left( \int_0^z \varphi_Y(ae^{-\eta_c u})du - z \right) \right\}. \quad (35)$$

We want to prove that the successive moments of $Y$ are characterized by this last equation. To avoid complicated computations, instead of giving their exact expressions, we prove that they are uniquely determined by the value of $\mu$.

To do this, we compute the derivatives of $\varphi_Y$ at 0 and need the following simple result: let $F$ be a function of class $C^\infty$. By induction on $n \geq 1$, one can prove that there exists a polynomial $P_n$ with $n - 1$ variables such that the $n$-th derivative of $e^F$ satisfies

$$(e^F)^{(n)} = e^F \left( F^{(n)} + P_n \left( F', F'', \ldots, F^{(n-1)} \right) \right).$$

Using this result, for $n \geq 2$, if we differentiate (35) $n$ times with respect to $a$ and take $a = 0$, we get

$$\varphi_Y^{(n)}(0) = \int_0^\infty \frac{\Lambda(dz)}{b} \left\{ (1-p)b \left( \int_0^z e^{-n\eta_c u}du \right) \varphi_Y^{(n)}(0) + P_n \left( D_z^{(1)}, D_z^{(2)}, \ldots, D_z^{(n-1)} \right) \right\} \quad (36)$$

where for $1 \leq k \leq n - 1$, $D_z^{(k)} = (1-p)b \varphi_Y^{(k)}(0) \int_0^z e^{-kn\eta_c u}du$. Since

$$(1-p) \int_0^\infty \frac{\Lambda(dz)}{b} \left( 1 - e^{-n\eta_c z} \right) = 1 - \frac{\psi_c(n\eta_c)}{\eta_c},$$

(36) is equivalent to

$$\varphi_Y^{(n)}(0) \frac{\psi_c(n\eta_c)}{n\eta_c} = \int_0^\infty \frac{\Lambda(dz)}{b} P_n \left( D_z^{(1)}, \ldots, D_z^{(n-1)} \right).$$

Moreover, we have $\varphi_Y^{(n)}(0) = (-1)^n \mathbb{E}[Y^n]$ and $D_z^{(k)} = b(1-p)(-1)^k \mathbb{E}[Y^k] \int_0^z e^{-kn\eta_c u}du$. Then, for $n \geq 2$, there exists a polynomial $P_n$ with $n - 1$ variables such that

$$\mathbb{E}[Y^n] \frac{\psi_c(n\eta_c)}{n\eta_c} = \tilde{P}_n \left( \mathbb{E}[Y], \mathbb{E}[Y^2], \ldots, \mathbb{E}[Y^{n-1}] \right). \quad (37)$$

If $n > 1$, $\psi_c(n\eta_c)$ is positive since by definition, $\eta_c$ is the largest root of $\psi_c$. Then, using (37), we can prove by induction that the moments $\mathbb{E}[Y^2], \mathbb{E}[Y^3], \ldots$ can be written as functions of $\mu = \mathbb{E}[Y]$. It implies that $Y$ has exactly the same moments as those of any r.v. with mean
μ and satisfying (35). In particular, Y has the same moments as the “pseudo-exponential” r.v. defined in the step 3 with c = P/μ, that is,

\[ \mathbb{E}[Y^k] = k! \frac{\mu^k}{pk-1}, \quad k \geq 0. \]

These moments satisfy the Carleman’s condition, which ensures that there is at most one r.v. with such moments (for instance, see [11, p.110-111]), that is, Y = X and the proof is complete.

**Remark 4.7.** We can also see \((L_i, i \geq 1)\) as a multitype CMJ-process: an allele of kind \(i\) has a lifespan distributed as the extinction time of \(\Xi_c\) and gives birth to alleles of kind \(i + 1\) according to the jump times of \(Y_{\int_0^t \Xi(s)ds, t \geq 0}\) where \(Y\) is a homogeneous Poisson process with rate \(bp\), independent of \(\Xi_c\).

In the supercritical case, it seems not possible to study the asymptotic behavior of \(L_i(t)\) with the techniques we used to obtain that of \(K_i(t)\). Indeed, some conditions of [25, Thm 4.3] are not fulfilled. As an example, the mean number of “children-alleles” of an allele is infinite since each allele has a positive probability \(\eta b(1−p)\) to survive forever and then to give birth to an infinite number of children.

### A Asymptotic behaviors of W and W′

#### A.1 Proof of Lemma 2.2(i)

We suppose here that \(m > 1\) which implies that \(\eta > 0\). To obtain the asymptotic behavior of \(W\), we use a Tauberian theorem about Laplace transform. Indeed, we have

\[ \int_0^\infty e^{-\lambda x} \left(e^{-\eta x}W(t)\right) dx = \frac{1}{\psi(\eta + \lambda)} \sim \frac{1}{\lambda \psi(\eta)}. \]

Then, since \(t \mapsto e^{-\eta t}W(t)\) is non-decreasing according to equation (4) in [8] (applied with \(q = 0\)), a Tauberian theorem entails the desired result (see for instance [1, p.10]).

To get the behavior of \(W'\), since \(t \mapsto e^{-\eta t}W'(t)\) is not necessarily non-decreasing, we can not use the same method. However, according to (4),

\[ e^{-\eta t}W'(t) = be^{-\eta t}W(t) - \int_0^t e^{-\eta(t-x)}W(t - x)e^{-\eta x} \Lambda(dx). \]

Since \(t \mapsto e^{-\eta t}W(t)\) is non-decreasing and converges to \(1/\psi'(\eta)\), the monotone convergence theorem implies that

\[ \lim_{t \to \infty} e^{-\eta t}W'(t) = \frac{b}{\psi'(\eta)} - \frac{1}{\psi'(\eta)} \int_0^\infty e^{-\eta x} \Lambda(dx). \]

The proof is then complete since from (2), we have \(0 = \psi(\eta) = \eta - b + \int_0^\infty e^{-\eta x} \Lambda(dx)\).
A.2 Proof of Lemma 2.2(ii)

We assume here that \( m = 1 \) and \( \sigma^2 = \int_0^\infty z^2 \Lambda(dz) < \infty \). First, notice that in that case \( \psi'(0) = 1 - m = 0 \) and that \( \psi''(0) = \sigma^2 \). Then, we have

\[
\int_0^\infty e^{-\lambda x} W(x)dx = \frac{1}{\psi(\lambda)} \lambda \sim \frac{2}{\sigma^2 \lambda^2}.
\]

Hence, a Tauberian theorem yields the asymptotic behavior of \( W(t) \) as \( t \to \infty \).

As in the supercritical case, it is not possible to obtain the asymptotic behavior of \( W' \) with the help of a Tauberian theorem because this function is not monotone. We use known results about the extinction time of a critical CMJ-process. According to the main result of [18], recalling that the measure \( \mu \) is defined by (1) and that \( \zeta \) is the lifespan of the ancestor, if

\[
\lim_{t \to \infty} t^2(1 - \mu([0, t])) = 0
\]

then

\[
\lim_{t \to \infty} t^2 \mathbb{P}(\zeta > t) = 0,
\]

where \( V \) is the variance of the number of children of the ancestor. Conditional on “the ancestor has a lifespan \( z \)”, this number is Poisson with parameter \( bz \). Then, we have \( V = b \int_0^\infty r^2 \Lambda(dr) \).

Moreover, according to (1),

\[
\int_0^\infty r \mu(dr) = \int_0^\infty rdr \int_r^\infty \Lambda(dx) = \sigma^2/2.
\]

Thus,

\[
\mathbb{P}(\Xi(t) > 0) \sim \frac{1}{tb}.
\]

It remains to prove that the conditions (38) and (39) hold. By using (1), the first one is equivalent to the condition (9) that we have assumes here. Moreover, \( \mathbb{P}(\zeta > t) = b^{-1} \Lambda((t, +\infty]) \), which goes to 0 faster than \( t^{-2} \) as \( t \to \infty \) since we the second moment of \( \Lambda \) is finite.

On the other hand, according to (5), we know that \( \mathbb{P}(\Xi(t) > 0) = \frac{W(t)}{W'(t)} \). Hence, \( W'(t) \) converges to \( 2/\sigma^2 \) as a consequence of (40) and of the fact that \( W(t) \) behaves as \( 2/(\sigma^2 t) \) as \( t \to \infty \).

A.3 Proof of Lemma 2.2(iii)

As in the critical case, the asymptotic of \( W \) can be obtained via a Tauberian theorem by using that \( \psi(\lambda) \sim (1 - m) \lambda \) as \( \lambda \to 0 \).

To study \( W' \) in that case, we use again known results about the time of extinction of CMJ-processes. The hypotheses (10) that we make about \( \tilde{\eta}_c \) enables us to use Theorem 6.7.10 in [20], that is,

\[
\tilde{C} := \lim_{t \to \infty} e^{-\tilde{\eta}t} \mathbb{P}(\Xi(t) > 0)
\]

exists and is positive if and only if

\[
\mathbb{E} \left[ \int_0^\infty e^{-\tilde{\eta}t} \xi(dt) \log \{ \xi((0, \infty)) \} \right] < \infty
\]
where we recall that $\xi$ is the birth point process of the ancestor. By conditioning by the lifespan of the ancestor and by using classical properties about Poisson point processes, we have

$$E \left[ \int_0^\infty e^{-\tilde{\eta}t} \xi(dt) \log\{\xi((0, \infty))\} \right] \leq \int_{(0,\infty)} \Lambda(dz) \frac{1-e^{-\tilde{\eta}z}}{\tilde{\eta}} ((1-p) + b(1-p)^2 z)$$

which is finite thanks to (10). Then, we have proved that there exists $\tilde{C} > 0$ such that

$$\mathbb{P}(\Xi(t) > 0) \sim t \to \infty \tilde{C} e^{\tilde{\eta}t}. \quad (41)$$

It remains to compute $\tilde{C}$ (which is unknown in most of CMJ-processes). Since $W(t)$ converges to $(1-m)^{-1}$ as $t \to \infty$ and using (41),

$$W'(t) \sim t \to \infty \frac{b}{1-m} \tilde{C} e^{\tilde{\eta}t}. \quad (42)$$

Moreover, by integrating (3) by parts, since $W(0) = 1$ according to [31, Lem 4.1], we have for $\lambda > \tilde{\eta}$

$$\int_0^\infty W'(u) e^{-\lambda u} du = \frac{\lambda}{\psi(\lambda)} - 1$$

with the convention $0/\psi(0) = 1/(1-m) < \infty$. Then,

$$\int_0^\infty (W'(u)e^{-\tilde{\eta}u}) e^{-\lambda u} du = \frac{\lambda + \tilde{\eta}}{\psi(\lambda + \tilde{\eta})} \sim \frac{\tilde{\eta}}{\lambda \psi'(\tilde{\eta})}.$$

On the other hand, using [4, Thm. 1.7.6] with (42), we get

$$\int_0^\infty (W'(u)e^{-\tilde{\eta}u}) e^{-\lambda u} du \sim \frac{b\tilde{C}}{1-m} \frac{1}{\lambda^2},$$

which implies that $\tilde{C} = \frac{\tilde{\eta}}{\psi(\tilde{\eta})}$.

Acknowledgments

This work was supported by project MANEGE ANR-09-BLAN-0215 (French national research agency). The author want to thank Amaury Lambert for his very helpful remarks and comments and Guillaume Achaz who raised a question about the number of mutations undergone by an allele during a talk given at Collège de France.

References

[1] J. Bertoin. *Lévy processes*, volume 121 of *Cambridge Tracts in Mathematics*. Cambridge University Press, Cambridge, 1996.

[2] J. Bertoin. The structure of the allelic partition of the total population for Galton-Watson processes with neutral mutations. *Ann. Probab.*, 37(4):1502–1523, 2009.
[3] J. Bertoin. A limit theorem for trees of alleles in branching processes with rare neutral mutations. *Stochastic Process. Appl.*, 120(5):678–697, 2010.

[4] N. H. Bingham, C. M. Goldie, and J. L. Teugels. *Regular variation*, volume 27 of *Encyclopedia of Mathematics and its Applications*. Cambridge University Press, Cambridge, 1989.

[5] N. Champagnat and A. Lambert. Splitting trees with neutral Poissonian mutations I: Small families. *Stochastic Process. Appl.*, 122(3):1003–1033, 2012.

[6] N. Champagnat and A. Lambert. Splitting trees with neutral poissonian mutations II: Largest and oldest families. *Stochastic Processes and their Applications*, 123(4):1368 – 1414, 2013.

[7] N. Champagnat, A. Lambert, and M. Richard. Birth and death processes with neutral mutations. *International Journal of Stochastic Analysis*, 2012 Article ID 569081, 2012.

[8] T. Chan, A. E. Kyprianou, and M. Savov. Smoothness of scale functions for spectrally negative Lévy processes. *Probab. Theory Related Fields*, 150(3-4):691–708, 2011.

[9] K. Danesh, R. Durrett, L. J. Havrilesky, and E. Myers. A branching process model of ovarian cancer. *Journal of Theoretical Biology*, 314(0):10 – 15, 2012.

[10] C. Delaporte. Lévy processes with marked jumps II : Application to a population model with mutations at birth. Preprint available at http://fr.arxiv.org/abs/1305.6491, 2013.

[11] R. Durrett. *Probability: theory and examples*. Duxbury Press, Belmont, CA, second edition, 1996.

[12] R. Durrett and S. Moseley. Evolution of resistance and progression to disease during clonal expansion of cancer. *Theoretical Population Biology*, 77(1):42 – 48, 2010.

[13] W. J. Ewens. The sampling theory of selectively neutral alleles. *Theoret. Population Biology*, 3:87–112; erratum, ibid. 3 (1972), 240; erratum, ibid. 3 (1972), 376, 1972.

[14] R. A. Fisher, A. S. Corbet, and C. B. Williams. The relation between the number of species and the number of individuals in a random sample of an animal population. *Journal of Animal Ecology*, 12:42–58, 1943.

[15] J. Geiger. Size-biased and conditioned random splitting trees. *Stochastic Process. Appl.*, 65(2):187–207, 1996.

[16] J. Geiger and G. Kersting. Depth-first search of random trees, and Poisson point processes. In *Classical and modern branching processes (Minneapolis, MN, 1994)*, volume 84 of *IMA Vol. Math. Appl.*, pages 111–126. Springer, New York, 1997.

[17] R. C. Griffiths and A. G. Pakes. An infinite-alleles version of the simple branching process. *Adv. in Appl. Probab.*, 20(3):489–524, 1988.

[18] J. M. Holte. Extinction probability for a critical general branching process. *Stochastic Processes and their Applications*, 2(3):303 – 309, 1974.
[19] Y. Iwasa, M. A. Nowak, and F. Michor. Evolution of resistance during clonal expansion. *Genetics*, 172(4):2557–2566, April 2006.

[20] P. Jagers. *Branching processes with biological applications*. Wiley-Interscience, London, 1975. Wiley Series in Probability and Mathematical Statistics—Applied Probability and Statistics.

[21] P. Jagers and O. Nerman. The growth and composition of branching populations. *Adv. in Appl. Probab.*, 16(2):221–259, 1984.

[22] P. Jagers and O. Nerman. Limit theorems for sums determined by branching and other exponentially growing processes. *Stochastic Process. Appl.*, 17(1):47–71, 1984.

[23] A. E. Kyprianou. *Introductory lectures on fluctuations of Lévy processes with applications*. Universitext. Springer-Verlag, Berlin, 2006.

[24] A. Lambert. The contour of splitting trees is a Lévy process. *Ann. Probab.*, 38(1):348–395, 2010.

[25] C. J. Mode. *Multitype branching processes. Theory and applications*. Modern Analytic and Computational Methods in Science and Mathematics, No. 34. American Elsevier Publishing Co., Inc., New York, 1971.

[26] O. Nerman. On the convergence of supercritical general (C-M-J) branching processes. *Z. Wahrsch. Verw. Gebiete*, 57(3):365–395, 1981.

[27] M. A. Nowak, F. Michor, and Y. Iwasa. The linear process of somatic evolution. *Proceedings of the National Academy of Sciences*, 100(25):14966–14969, 2003.

[28] A. G. Pakes. An infinite alleles version of the Markov branching process. *J. Austral. Math. Soc. Ser. A*, 46(1):146–169, 1989.

[29] D. Revuz and M. Yor. *Continuous martingales and Brownian motion*, volume 293 of *Grundlehren der Mathematischen Wissenschaften [Fundamental Principles of Mathematical Sciences]*. Springer-Verlag, Berlin, third edition, 1999.

[30] M. Richard. *Arbres, Processus de branchement non markoviens et Processus de Lévy*. PhD thesis, UPMC, Paris 6, 2011. Available at http://tel.archives-ouvertes.fr/tel-00649235/fr/.

[31] M. Richard. Limit theorems for supercritical age-dependent branching processes with neutral immigration. *Advances in Appl. Probability*, 43-1:276–300, 2011.

[32] S. Sagitov and M. C. Serra. Multitype Bienaymé-Galton-Watson processes escaping extinction. *Adv. in Appl. Probab.*, 41(1):225–246, 2009.

[33] Z. Taïb. *Branching processes and neutral evolution*. Lecture Notes in Biomathematics. 93. Berlin: Springer-Verlag. viii, 112 p., 1992.