

ON COMPOUND POISSON POPULATION MODELS

Martin Möhle, University of Tübingen

(joint work with Thierry Huillet, Université de Cergy-Pontoise)

Workshop on Probability, Population Genetics and Evolution

Centre International de Rencontres Mathématiques (CIRM)

Marseille-Luminy, France

June 11, 2012

Exchangeable population models (Cannings)

Non-overlapping generations $r \in \mathbb{Z} = \{\dots, -1, 0, 1, \dots\}$

Population size N , i.e. N individuals (genes, particles) in each generation

$\nu_i^{(r)}$:= number of offspring of individual i of generation r , $\nu_1^{(r)} + \dots + \nu_N^{(r)} = N$

Additional assumptions on the offspring

Exchangeability: $(\nu_{\pi 1}^{(r)}, \dots, \nu_{\pi N}^{(r)}) \stackrel{d}{=} (\nu_1^{(r)}, \dots, \nu_N^{(r)}) \quad \forall \pi$

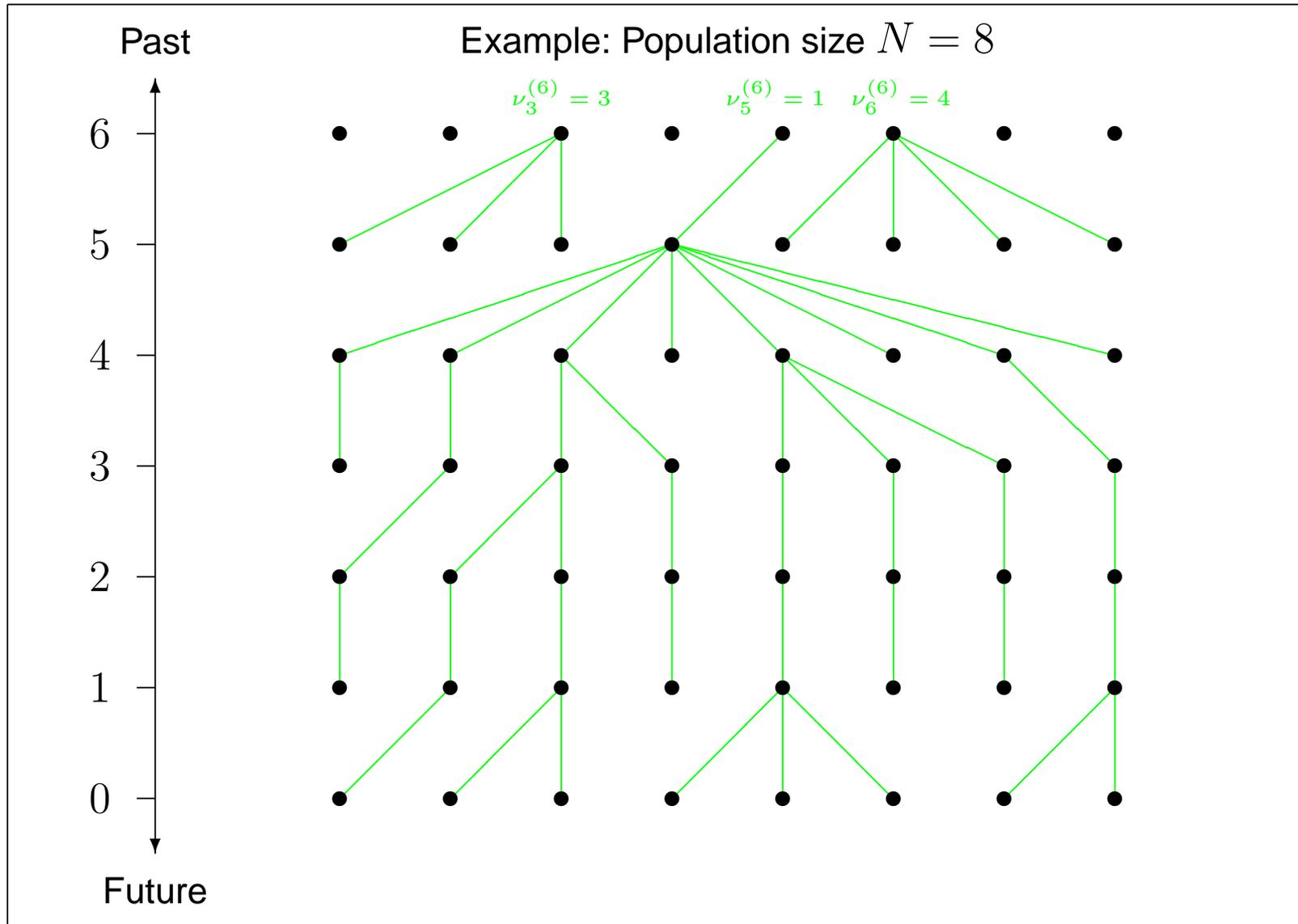
Homogeneity: $(\nu_1^{(r)}, \dots, \nu_N^{(r)})_r$ i.i.d.

Avoid the trivial model $(\nu_i^{(r)} \equiv 1)$. Define $(\nu_1, \dots, \nu_N) := (\nu_1^{(0)}, \dots, \nu_N^{(0)})$.

Examples. **Wright-Fisher:** $(\nu_1, \dots, \nu_N) \stackrel{d}{=} \text{Multinomial}(N, \frac{1}{N}, \dots, \frac{1}{N})$

Moran: $(\nu_1, \dots, \nu_N) = \text{random permutation of } (2, \underbrace{1, \dots, 1}_{N-2}, 0)$

Exchangeable population models (graphical representation)



Conditional branching population models

Let ξ_1, ξ_2, \dots be independent random variables taking values in $\mathbb{N}_0 := \{0, 1, 2, \dots\}$.

Assume that $\mathbb{P}(\xi_1 + \dots + \xi_N = N) > 0$ for all $N \in \mathbb{N} := \{1, 2, \dots\}$.

Perform the following two steps:

1. Conditioning: Let μ_1, \dots, μ_N be random variables such that the distribution of them coincides with that of ξ_1, \dots, ξ_N conditioned on the event that $\xi_1 + \dots + \xi_N = N$.

2. Shuffling: Let ν_1, \dots, ν_N be the μ_1, \dots, μ_N randomly permuted.

Conditional branching process models are particular Cannings models with offspring variables ν_1, \dots, ν_N constructed as above (Karlin and McGregor, 1964).

Compound Poisson population models

Compound Poisson population models are particular conditional branching process models for which ξ_n has p.g.f.

$$\mathbb{E}(x^{\xi_n}) = \exp(-\theta_n(\phi(z) - \phi(zx))), \quad |x| \leq 1, n \in \mathbb{N},$$

with parameters $0 < \theta_n < \infty$ and with a power series ϕ of the form $\phi(z) = \sum_{m=1}^{\infty} \phi_m \frac{z^m}{m!}$, $|z| < r$, with positive radius r of convergence and $\phi_m \geq 0$, $\phi_1 > 0$.

Distribution and factorial moments of μ

Notation: Taylor expansion $\exp(\theta\phi(z)) = \sum_{k=0}^{\infty} \frac{\sigma_k(\theta)}{k!} z^k, |z| < r.$

(The coefficients $\sigma_k(\theta)$ can be computed recursively.)

Distribution of μ :
$$\mathbb{P}(\mu_1 = j_1, \dots, \mu_N = j_N) = \frac{N!}{j_1! \cdots j_N!} \frac{\sigma_{j_1}(\theta_1) \cdots \sigma_{j_N}(\theta_N)}{\sigma_N(\sum_{n=1}^N \theta_n)}$$

$$(j_1, \dots, j_N \in \mathbb{N}_0 \text{ with } j_1 + \cdots + j_N = N)$$

Factorial moments of μ :

$$\mathbb{E}((\mu_1)_{k_1} \cdots (\mu_N)_{k_N}) = \frac{N!}{\sigma_N(\sum_{n=1}^N \theta_n)} \sum_{\substack{j_1 \geq k_1, \dots, j_N \geq k_N \\ j_1 + \cdots + j_N = N}} \frac{\sigma_{j_1}(\theta_1) \cdots \sigma_{j_N}(\theta_N)}{(j_1 - k_1)! \cdots (j_N - k_N)!}$$

$$(k_1, \dots, k_N \in \mathbb{N}_0)$$

A subclass of compound Poisson models

We focus on the subclass of compound Poisson models satisfying

$$\frac{\sigma_{k+1}(\theta)}{\sigma_k(\theta)} + \frac{\sigma_{k'+1}(\theta')}{\sigma_{k'}(\theta')} = \frac{\sigma_{k+k'+1}(\theta + \theta')}{\sigma_{k+k'}(\theta + \theta')} \quad k, k' \in \mathbb{N}_0, \theta, \theta' \in (0, \infty). \quad (*)$$

Lemma. A compound Poisson model satisfies (*) if and only if $\phi_m = (m-1)! \phi_1 (\phi_2 / \phi_1)^{m-1}$ for all $m \in \mathbb{N}$. (These are essentially Wright–Fisher models and Dirichlet models.) If (*) holds then μ has factorial moments

$$\mathbb{E}((\mu_1)_{k_1} \cdots (\mu_N)_{k_N}) = (N)_{k_1 + \cdots + k_N} \frac{\sigma_{k_1}(\theta_1) \cdots \sigma_{k_N}(\theta_N)}{\sigma_k(\theta_1 + \cdots + \theta_N)}$$

$$(k_1, \dots, k_N \in \mathbb{N}_0)$$

Assumption. In the following it is always assumed that (*) holds.

Ancestral process

Take a sample of $n (\leq N)$ individuals from some generation and consider their ancestors.

$(i, j) \in \mathcal{R}_t = \mathcal{R}_t^{(n)} : \iff$ individuals i and j have a common parent t generations backwards in time.

The **ancestral process** $\mathcal{R} := (\mathcal{R}_t)_{t=0,1,\dots}$ is **Markovian** with state space \mathcal{E}_n (set of equivalence relations on $\{1, \dots, n\}$).

Transition probabilities: $\mathbb{P}(\mathcal{R}_{t+1} = \eta \mid \mathcal{R}_t = \xi) = \Phi_j^{(N)}(k_1, \dots, k_j)$, $\xi, \eta \in \mathcal{E}_n$ with $\xi \subseteq \eta$ with

$$\Phi_j^{(N)}(k_1, \dots, k_j) := \frac{1}{\sigma_{k_1 + \dots + k_j}(\sum_{n=1}^N \theta_n)} \sum_{\substack{n_1, \dots, n_j=1 \\ \text{all distinct}}}^N \sigma_{k_1}(\theta_{n_1}) \cdots \sigma_{k_j}(\theta_{n_j})$$

where $j := |\eta|$ = number of blocks of η ,

$k_1, \dots, k_j :=$ group sizes of merging classes of ξ . ($\Rightarrow k_1 + \dots + k_j = |\xi|$)

Two basic transition probabilities

Notation: For $N, k \in \mathbb{N}$ define $\Theta_k(N) := \sum_{n=1}^N \theta_n^k$.

$$\begin{aligned} c_N &:= \text{'coalescence probability'} := \mathbb{P}(2 \text{ individuals have same parent}) \\ &= \Phi_1^{(N)}(2) = \frac{1}{\sigma_2(\Theta_1(N))} \sum_{n=1}^N \sigma_2(\theta_n) = \frac{\phi_2 \Theta_1(N) + \phi_1^2 \Theta_2(N)}{\phi_2 \Theta_1(N) + \phi_1^2 (\Theta_1(N))^2}. \end{aligned}$$

We also need

$$\begin{aligned} d_N &:= \mathbb{P}(3 \text{ individuals have same parent}) \\ &= \Phi_1^{(N)}(3) = \frac{\phi_3 \Theta_1(N) + 3\phi_1 \phi_2 \Theta_2(N) + \phi_1^3 \Theta_3(N)}{\phi_3 \Theta_1(N) + 3\phi_1 \phi_2 (\Theta_1(N))^2 + \phi_1^3 (\Theta_1(N))^3}. \end{aligned}$$

Exchangeable coalescent processes

- **Exchangeable coalescents** are discrete-time or continuous-time Markov processes $\Pi = (\Pi_t)_t$ with state space \mathcal{E} , the set of equivalence relations (partitions) on $\mathbb{N} := \{1, 2, \dots\}$.
- During each transition, equivalence classes (blocks) merge together. **Simultaneous multiple collisions** of blocks are allowed.
- Schweinsberg (2000) characterizes these processes via a **finite measure** Ξ on the infinite simplex

$$\Delta := \left\{ x = (x_1, x_2, \dots) : x_1 \geq x_2 \geq \dots \geq 0, |x| := \sum_{i=1}^{\infty} x_i \leq 1 \right\}.$$

- These processes are therefore also called **Ξ -coalescents**.

Domain of attraction

For $n \in \mathbb{N}$ let $\varrho_n : \mathcal{E} \rightarrow \mathcal{E}_n$ denote the restriction of \mathcal{E} to \mathcal{E}_n , the set of equivalence relations on $\{1, \dots, n\}$.

Definition.

- We say that the considered population model is in the **domain of attraction of a continuous-time coalescent** $\Pi = (\Pi_t)_{t \geq 0}$, if, for each sample size $n \in \mathbb{N}$, the time-scaled ancestral process $(\mathcal{R}_{\lfloor t/c_N \rfloor}^{(n)})_{t \geq 0}$ weakly converges to $(\varrho_n \circ \Pi_t)_{t \geq 0}$ as $N \rightarrow \infty$.
- We say that the considered population model is in the **domain of attraction of a discrete-time coalescent** $\Pi = (\Pi_t)_{t=0,1,\dots}$, if, for each sample size $n \in \mathbb{N}$, the ancestral process $(\mathcal{R}_t^{(n)})_{t=0,1,\dots}$ weakly converges to $(\varrho_n \circ \Pi_t)_{t=0,1,\dots}$ as $N \rightarrow \infty$.

Results (Regime 1)

Theorem 1. Suppose that (*) holds. If $\sum_{n=1}^{\infty} \theta_n < \infty$, then the compound Poisson population model is in the domain of attraction of a **discrete-time Ξ -coalescent**.

Characterization of Ξ . There exists a consistent sequence $(Q_j)_{j \in \mathbb{N}}$ of probability measures Q_j on the j -simplex $\Delta_j := \{(x_1, \dots, x_j) \in [0, 1]^j : x_1 + \dots + x_j \leq 1\}$ uniquely determined via their moments

$$\int_{\Delta_j} x_1^{k_1} \cdots x_j^{k_j} Q_j(dx_1, \dots, dx_j) = \frac{\sigma_{k_1}(\theta_1) \cdots \sigma_{k_j}(\theta_j)}{\sigma_{k_1 + \dots + k_j}(\sum_{n=1}^{\infty} \theta_n)}, \quad k_1, \dots, k_j \in \mathbb{N}_0.$$

Let Q denote the projective limit of $(Q_j)_{j \in \mathbb{N}}$, let X_1, X_2, \dots be random variables with joint distribution Q , and let ν be the joint distribution of the ordered variables $X_{(1)} \geq X_{(2)} \cdots$. Then, Ξ has density $x \mapsto (x, x) := \sum_{n=1}^{\infty} x_n^2$ with respect to ν . The measure Ξ is concentrated on the subset Δ^* of points $x = (x_1, x_2, \dots) \in \Delta$ satisfying $|x| := \sum_{n=1}^{\infty} x_n = 1$.

Regime 1 (continued)

Remark. The proof of Theorem 1 is based on [general convergence theorems for ancestral processes of Cannings models](#) (M. and Sagitov 2001) and on the [moment problem for the \$j\$ -dimensional simplex](#) (Gupta).

Examples. Suppose that $\theta := \sum_{n=1}^{\infty} \theta_n < \infty$.

Wright-Fisher models. If $\phi(z) = \phi_1 z$, then $\sigma_k(\theta) = \theta^k \phi_1^k$. In this case ν is the Dirac measure at $p = (\theta_1/\theta, \theta_2/\theta, \dots) \in \Delta^*$. The measure Ξ assigns its total mass $\Xi(\Delta) = (p, p) = (\sum_{n=1}^{\infty} \theta_n^2)/\theta^2$ to the single point p .

Dirichlet models. If $\phi(x) = -\log(1 - x)$, then $\phi_m = (m - 1)!$, $m \in \mathbb{N}$, and $\sigma_k(\theta) = [\theta]_k := \theta(\theta + 1) \cdots (\theta + k - 1)$, $k \in \mathbb{N}_0$. The limiting coalescent is the discrete-time Dirichlet-Kingman coalescent with parameter $(\theta_n)_{n \in \mathbb{N}}$.

Results (Regime 2)

Theorem 2. Suppose that (*) holds. If $\sum_{n=1}^{\infty} \theta_n = \infty$ and if $\sum_{n=1}^{\infty} \theta_n^2 < \infty$, then the compound Poisson population model is in the domain of attraction of the **Kingman coalescent**.

Remarks.

1. Time-scaling satisfies $c_N = \Theta_2(N)/(\Theta_1(N))^2$ if $\phi_2 = 0$ and $c_N \sim \phi_2/(\phi_1^2 \Theta_1(N))$ if $\phi_2 > 0$, where $\Theta_k(N) := \sum_{n=1}^N \theta_n^k$ for $k \in \mathbb{N}$.
2. In contrast to the situation in Theorem 1, the limiting coalescent in Theorem 2 does not depend on the function ϕ of the compound Poisson model. Theorem 2 is for example applicable if $\theta_n = n^{-\alpha}$ with $\alpha \in (\frac{1}{2}, 1]$.

Sketch of proof

The proof of Theorem 2 is based on the following technical lemma.

Lemma. If (*) holds, then the following five conditions are equivalent.

$$(i) \lim_{N \rightarrow \infty} \frac{\Theta_2(N)}{(\Theta_1(N))^2} = 0.$$

$$(ii) \lim_{N \rightarrow \infty} \frac{\Theta_3(N)}{\Theta_1(N) \Theta_2(N)} = 0.$$

$$(iii) \lim_{N \rightarrow \infty} c_N = 0.$$

$$(iv) \lim_{N \rightarrow \infty} \frac{d_N}{c_N} = 0.$$

(v) The compound Poisson model is in the domain of attraction of the [Kingman coalescent](#).

Remark. The proof that (i) - (iii) are equivalent is technical but elementary. The equivalence of (iv) and (v) and the implication '(iv) \Rightarrow (iii)' hold even for arbitrary Cannings models (M., 2000). [The interesting point is that, for compound Poisson models, \(iii\) implies \(iv\).](#) Note that this implication does not hold for arbitrary Cannings models.

Results (Regime 3)

Theorem 3. Suppose that (*) holds, that $\sum_{n=1}^{\infty} \theta_n = \infty$ and that $\sum_{n=1}^{\infty} \theta_n^2 = \infty$. Then the compound Poisson model is in the domain of attraction of the **Kingman coalescent** if and only if $\Theta_2(N)/(\Theta_1(N))^2 \rightarrow 0$ as $N \rightarrow \infty$. In this case the time-scaling c_N satisfies $c_N \sim \phi_2/(\phi_1^2\Theta_1(N)) + \Theta_2(N)/(\Theta_1(N))^2$.

Corollary. (unbiased case, Huillet and M., 2010) If (*) holds and if $\theta_n = \theta$ does not depend on n , then the compound Poisson model is in the domain of attraction of the **Kingman coalescent**. The time-scaling c_N satisfies $c_N \sim (1 + \phi_2/(\phi_1^2\theta))/N$.

Results (Regime 3, continued)

Theorem 4. Suppose that (*) holds and that all the limits

$$p_1(k) := \lim_{N \rightarrow \infty} \frac{\Theta_k(N)}{(\Theta_1(N))^k}, \quad k \in \mathbb{N},$$

exist. Then all the limits

$$p_j(k_1, \dots, k_j) := \lim_{N \rightarrow \infty} \frac{1}{(\Theta_1(N))^{k_1 + \dots + k_j}} \sum_{\substack{n_1, \dots, n_j = 1 \\ \text{all distinct}}}^N \theta_{n_1}^{k_1} \dots \theta_{n_j}^{k_j},$$

$k_1, \dots, k_j \in \mathbb{N}$, exist. Suppose now in addition that $\sum_{n=1}^{\infty} \theta_n = \infty$ and that $p_1(2) > 0$. Then, the compound Poisson model is in the domain of attraction of a **discrete-time Ξ -coalescent Π** . The characterizing measure $\nu(dx) := \Xi(dx)/(x, x)$ of Π is the Dirac-measure at $x = (x_1, x_2, \dots) \in \Delta$, where $x_1 := \lim_{k \rightarrow \infty} (p_1(k))^{1/k}$ and $x_{n+1} := \lim_{k \rightarrow \infty} (p_1(k) - (x_1^k + \dots + x_n^k))^{1/k}$, $n \in \mathbb{N}$.

Regime 3 (continued)

Remarks.

- Let Z_N be a random variable taking the value $\theta_n/\Theta_1(N)$ with probability $\theta_n/\Theta_1(N)$, $n \in \{1, \dots, N\}$. The existence of all the limits $p_1(k)$, $k \in \mathbb{N}$, is equivalent to the convergence $Z_N \rightarrow Z$ in distribution, where Z has characteristic function $t \mapsto \sum_{k=0}^{\infty} (t^k/k!)p_1(k+1)$, $t \in \mathbb{R}$.
- In contrast to the situation in Theorem 1, the limiting discrete-time Ξ -coalescent in Theorem 4 does not depend on the function ϕ of the compound Poisson model.

Example. Fix $\lambda > 1$ and suppose that $\theta_n = \lambda^n$, $n \in \mathbb{N}$. Then, $p_1(k) = (\lambda - 1)^k / (\lambda^k - 1) > 0$, $k \in \mathbb{N}$. In this case the measure Ξ of the limiting Ξ -coalescent assigns its total mass $\Xi(\Delta) = p_1(2) = (\lambda - 1)/(\lambda + 1)$ to the single point $x = (x_1, x_2, \dots) \in \Delta^*$ defined via $x_n := (\lambda - 1)/\lambda^n = (1 - 1/\lambda)(1/\lambda)^{n-1}$, $n \in \mathbb{N}$.

Generalization: Assume that (*) does not hold

Theorem. (Huillet, M. 2011)

Fix $\theta \in (0, \infty)$ and suppose that the equation $\theta z \phi'(z) = 1$ has a real solution $z(\theta) \in (0, r)$. Then $\mu_1 \rightarrow X$ in distribution as $N \rightarrow \infty$, where X has distribution

$$\mathbb{P}(X = k) = \sigma_k(\theta) \frac{(z(\theta))^k}{k!} e^{-\theta \phi(z(\theta))} \quad k \in \{0, 1, 2, \dots\}.$$

The associated **symmetric compound Poisson model** is in the domain of attraction of the **Kingman coalescent**. The **effective population size** $N_e := 1/c_N$ satisfies $N_e \sim \varrho N$ as $N \rightarrow \infty$, where $\varrho := 1/\mathbb{E}((X)_2) = 1/(1 + \theta(z(\theta))^2 \phi''(z(\theta))) \in (0, 1]$.

Remark. Proof uses the **saddle point method** to establish the asymptotics of $\sigma_N(N\theta)$.

Open cases.

- a) Symmetric models without a solution $z(\theta)$, condensation (work in progress).
- b) Non symmetric models.

Conclusions

- Asymptotics of the ancestry for some compound Poisson population models analyzed
- **Results essentially based on convergence theorems** (M. 2000 and M. and Sagitov 2001) for ancestral processes of **exchangeable Cannings population models**
- Convergence to the **Kingman coalescent** if and only if $\Theta_2(N)/(\Theta_1(N))^2 \rightarrow 0$
- Compound Poisson models satisfying (*) are **never in the domain of attraction of a continuous-time coalescent different from Kingman's coalescent**; discrete-time Ξ -coalescents (with simultaneous multiple collisions) arise if the parameters θ_n are 'unbalanced'
- **Three regimes** depending on the behavior of the series $\sum_n \theta_n$ and $\sum_n \theta_n^2$; **complete convergence results for the first two regimes**; **partial convergence results for the third regime** (when both series diverge)
- **Convergence to the Kingman coalescent** holds even for more general symmetric compound Poisson models, which do not necessarily satisfy the restriction (*)

References

- CANNINGS, C. (1974) The latent roots of certain Markov chains arising in genetics: a new approach, I. Haploid models. *Adv. Appl. Probab.* **6**, 260–290.
- GUPTA, J. C. (1999) The moment problem for the standard k -dimensional simplex. *Sankhyā A* **61**, 286–291.
- HUILLET, T. AND MÖHLE, M. (2011) Population genetics models with skewed fertilities: a forward and backward analysis. *Stochastic Models* **27**, 521–554.
- HUILLET, T. AND MÖHLE, M. (2012) Correction on ‘Population genetics models with skewed fertilities: a forward and backward analysis’. *Stochastic Models*, to appear.
- KARLIN, S. AND MCGREGOR, J. (1964) Direct product branching processes and related Markov chains. *Proc. Nat. Acad. Sci. U.S.A.* **51**, 598–602.

References (continued)

KINGMAN, J. F. C. (1982) The coalescent. *Stochastic Process. Appl.* **13**, 235–248.

MÖHLE, M. (2000) Total variation distances and rates of convergence for ancestral coalescent processes in exchangeable population models. *Adv. Appl. Probab.* **32**, 983–993.

MÖHLE, M. (2011) Coalescent processes derived from some compound Poisson population models. *Electron. Comm. Probab.* **16**, 567–582.

MÖHLE, M. AND SAGITOV, S. (2001) A classification of coalescent processes for haploid exchangeable population models. *Ann. Probab.* **29**, 1547–1562.

SCHWEINSBERG, J. (2000) Coalescents with simultaneous multiple collisions. *Electron. J. Probab.* **5**, 1–50.