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The Contour Process of Crump-Mode-Jagers Branching Processes

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June 24, 2015

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Crump-Mode-Jagers trees

- Crump–Mode–Jagers (CMJ) branching processes generalize Galton–Watson process.
- Individuals live for a random duration and give birth at random times during their life-time.
- Each individual is characterized by a random pair $(\mathcal{P}, V) \in \mathcal{M} \times \mathbb{R}^+$
 - V is the life-length of the individual

$$V \ge \tau^1 \ge \tau^2 \cdots \ge \tau^{|\mathcal{P}|}$$

$$\mathcal{P} = \sum_{i=1}^{|\mathcal{P}|} \delta_{\tau^k}$$

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Results

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Crump-Mode-Jagers forests

- A CMJ forest is constructed from a sequence of i.i.d. life-descriptors {(P_n, V_n)}_{n≥0} ∈ M × ℝ⁺
- The root's life length is equal to V_0 . We graft new individuals at the atoms of \mathcal{P}_0 .
- The process is repeated until the population gets extinct, which happens almost surely in the (sub)critical case $\mathbb{E}(|\mathcal{P}|) \leq 1.$
- At extinction time, generate a new tree according to the same procedure.



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Chronological and genealogical structure

- Genealogy: the Galton Watson forest generated from the sequence {|P_n|}_{n≥0}. This GW forest encodes the genealogical structure of the CMJ.
- Chronology: the CMJ itself.





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Population dynamics

• The population size of a finite (but) large CMJ process is not Markovian.

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- Ex: Bellman-Harris processes. individuals beget their children at death, independently of their life-time.
- Need to keep track of the population size and the age structure in the population. This might be represented by a measure-valued process (Sagitov ('97)).
- General case: need to keep track of the age structure and birth events occurring in the past.





- Different point of view: contour processes.
- The contour process of a CMJ process encodes not only the population dynamics, but also the geometry of the underlying trees (or forests).
- In the particular case of Poissonian birth events, Lambert ('10) showed that the contour process is a Lévy process.
- Using this approach, we will be able to show that CMJ forests fall into three main categories:
 - If edges are short enough, CMJ forests are obtained by a deterministic stretching of their underlying genealogy.
 - If the offspring distribution has a finite second finite, a CMJ forest looks asymptotically like a Bellman-Harris forest.
 - For long edges and large number of offspring, our approach provides an educated guess for a natural scaling limit.

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Coding discrete planar trees



Figure : pictures taken from Le Gall's lecture notes

- Contour process: distance from the origin of a particle exploring the tree from left to right by travelling at unit speed along the edges.
- Height process: distance from the origin of the *n*th vertex visited in lexicographical order.

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From tree to contour and vice-versa.

- To every rooted planar tree corresponds a contour process.
- The genealogy can be recovered from the sub-excursions nested in the contour process.
- Real tree: tree constructed from an arbitrary excursion.









- Let f be a non-negative continuous function made of finite excursions with f(0) = 0.
- $x \sim y$ iff $\inf_{[x \wedge y, x \vee y]} f = f(x) = f(y)$

•
$$\mathcal{T}_f = \mathbb{R}^+ / \sim$$
.

• $d_f(x,y) = f(x) + f(y) - 2 \inf_{[x \land y, x \lor y]} f.$

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| Real tre | es | | | |

- $(\mathcal{T}_f, \mathit{d}_f)$ defines a real forest in the sense that
 - (i) (Unique geodesics.) There is a unique isometric map ψ^{a,b} from [0, d_f(a, b)] into T_f such that ψ^{a,b}(0) = a and ψ^{a,b}(d_f(a, b)) = b.
 - (ii) **(Loop free.)** If q is a continuous injective map from [0, 1] into \mathcal{T}_f , such that q(0) = a and q(1) = b, we have $q([0,1]) = \psi^{a,b}([0, d_f(a, b)]).$
 - Continuous Random Forest (Aldous ('91)): Real forest encoded by a reflected Brownian motion.

 \bullet Let ${\mathcal F}$ be a Galton Watson forest such that

$$\sum_{k=0}^{\infty} kp(k) = 1$$
 (critical case) and $0 < \sum_{k=0}^{\infty} k^2 p(k) - 1 = \sigma^2 < \infty$

and $\ensuremath{\mathcal{C}}$ its contour process.

- $\frac{1}{\sqrt{n}}(\mathcal{H}(n\cdot),\mathcal{C}(n\cdot))$ converges in the weak topology to $\frac{2}{\sigma}(|w|(\cdot),|w|(\cdot/2))$, where w is a standard Brownian motion.
- This indicates that the (rescaled) Galton Watson forest converges to the real forest encoded by a reflected Brownian motion.

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 Scaling limit of Galton Watson genealogies: infinite variance case (Le Gall Le Jan ('98), Duquesne Le Gall ('02))
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 $\bullet\,$ Let ${\mathcal F}$ be a Galton Watson forest such that

$$\sum_{k=0}^{\infty} kp(k) = 1 \quad \text{(critical case)}$$

- The offspring distribution {p(k)} is in the domain of attraction of the stable law with exponent α ∈ (1,2).
- The limiting process is not Markovian but can be expressed as a functional of a spectrally positive Lévy process



- In this talk, we will consider the height process and the contour process of a CMJ forest.
- Let $\mathbb H$ (resp., $\mathbb C)$ be the height (resp., contour) process of the CMJ.
- Let \mathcal{H} (resp., \mathcal{C}) be the height (resp., contour) process of the underlying Galton Watson forest.



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Back to the original question

- Question 1: limiting behavior of $\mathbb C$ at large time.
- Question 2: joint convergence of (\mathbb{C}, C) and $(\mathbb{H}, \mathcal{H})$, i.e., relation between the CMJ forest and its underlying genealogy.

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• Define y to be the random variable such that for every test function f:

$$\mathbb{E}[f(y)] = \sum_{k=1}^{\infty} \left(\frac{1}{k} \sum_{r=1}^{k} \mathbb{E}[f(A(\mathcal{P}, r)) \mid |\mathcal{P}| = k]\right) k \mathbb{P}(|\mathcal{P}| = k)$$

where $A(\mathcal{P}, r)$ is the position of the r^{th} atom on \mathcal{P}

- Assume that $\mathbb{E}(y) < \infty$
- Assume that there exists $\epsilon_n \rightarrow 0$ and \mathcal{H}_{∞} such that

$$\epsilon_n \mathcal{H}(n \cdot) \implies \mathcal{H}_{\infty}.$$

Then

$$\epsilon_n(\mathcal{H}(n\cdot), \mathbb{H}(n\cdot)) \rightarrow (\mathcal{H}_\infty, \mathbb{E}(y)\mathcal{H}_\infty).$$

in the sense of f.d.d.

- Assume that $\mathbb{E}(V) < \infty$.
- Assume that there exists $\epsilon_n \rightarrow 0$

$$\epsilon_n \mathcal{H}([n \cdot]) \Longrightarrow \mathcal{H}_{\infty}(\cdot), \ \epsilon_n \mathcal{C}([n \cdot]) \Longrightarrow \mathcal{C}_{\infty}(\cdot)$$

• Assume that there exists $\bar{\epsilon}_n \to 0$ such that $\{\bar{\epsilon}_n \mathbb{H}(nt)\}_n$ is tight (for every determinstic t) then

$$\overline{\epsilon}_n\left(\mathbb{H}\left(\frac{[nt]}{2\mathbb{E}(V)}\right)-\mathbb{C}([nt])\right)\implies 0$$

in the sense of f.d.d..

 Generalization of a result by Duquesne & Le Gall in the discrete setting. Non trivial due to the absence of tightness.



- X in the domain of attraction of an α -stable law with $\alpha \in (1, 2)$.
- $\mathcal{P} = \delta_1(X-1) + \delta_X$ in such a way that $|\mathcal{P}|$ is distributed as X.
- $\mathbb{E}(y) < \infty$.
- $\frac{1}{n^{1-\frac{1}{\alpha}}}\mathcal{H}([n\cdot]) \implies \mathcal{H}_{\infty}.$
- $\frac{1}{n^{1-\frac{1}{\alpha}}}\mathbb{H}([n\cdot]) \implies \mathbb{E}(y)\mathcal{H}_{\infty}.$
- $\max_{0,\dots,n} \mathbb{H} \geq \max_{0,\dots,n} X_i \sim n^{1/\alpha}$
- Taking $\alpha < \frac{1}{2}(1+\sqrt{5})$, we have $n^{1/lpha} >> n^{1-\frac{1}{lpha}}$
- As n goes to ∞, one can find infinitely many edges which do not scale as the height process.

- $\mathbb{E}(V) < \infty, \mathbb{E}(y) < \infty$
- Combining (R1) and (R2)

 $\epsilon_n(\mathcal{C}([n\cdot],\mathbb{C}([n\cdot]) \to (\mathcal{C}_\infty, \mathbb{E}(y)\mathcal{C}_\infty(\cdot/2\mathbb{E}(V)))$

• Sagitov ('95)

$$\epsilon_n Z([nt]) o rac{\mathbb{E}(V)}{\mathbb{E}(y)} \mathcal{Z}(t/2\mathbb{E}(y))$$

where Z is a CSBP starting at 1 and Z the discrete CMJ branching processes BP starting with $[1/\epsilon_n]$ individuals.

Question: What if E(y) = ∞, E(V) = ∞? We distinguish between two cases (1) E(|P|²) < ∞, and (2) E(|P|²) = ∞.

finite variance case

- $\mathbb{E}(|\mathcal{P}|^2) < \infty$ but y is the domain of attraction of an α -stable law with $\alpha \in (0, 1)$.
- There exists $\epsilon_n \rightarrow 0$ such that the joint distribution of

$$\left(\frac{1}{n^{1/2}}\mathcal{H}(n\cdot), \ \epsilon_n\mathbb{H}(n\cdot)\right)$$

can be asymptotically described in terms of the Poisson snake.



- Continuum analog of Bellman-Harris forests.
- Start from the genealogical structure.
- Mark every edge with an independent random number according to life-length distribution V.
- The BH forest can be recovered by a simple (random) stretching of the underlying genealogy.
- Informelly, our results indicates that if |P| has a finite second moment, then CMJ branching processes behave as a Bellman Harris forest.

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Poisson snake (Warren ('02), Abraham Delmas ('02) etc.)

- Let $(\mathcal{F}_{\frac{2}{\sigma}|w|}, d_{\frac{2}{\sigma}|w|})$ be the real forest induced by a reflected Brownian motion.
- Let $\lambda_{\frac{2}{\sigma}|w|}$ be the branch length measure, i.e. $\forall a, b$ $\lambda_{\frac{2}{\sigma}|w|}([a, b]) = d_{\frac{2}{\sigma}|w|}(a, b).$
- Conditioned on $(\mathcal{F}_{\frac{2}{\sigma}|w|}, d_{\frac{2}{\sigma}|w|})$, mark the forest with a Poisson Point process on $\mathcal{F}_{\frac{2}{\sigma}|w|} \times \mathbb{R}^+$ with intensity measure

$$\lambda_{rac{2}{\sigma}|w|} imes rac{dl}{l^{lpha+1}}$$



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• $\mathbb{H}_{\infty}(t)$ is the sum of all the atoms along the branch [
ho,t].

$$egin{aligned} &\left(rac{1}{\sqrt{n}}\mathcal{H}([nt]), \ \epsilon_{n}\mathbb{H}([nt]); \ t\geq 0
ight) \ &
ightarrow \ \left(\ rac{2}{\sigma}|w|(t), \ \mathbb{H}_{\infty}(t); \ t\geq 0
ight) \end{aligned}$$

in the sense of f.d.d.

• The chronological ancestral line is obtained by random dilatation of the genealogical ancestral line.



Result 4: Contour process for offspring distribution with finite variance case

- E(|P|²) < ∞. V is in the domain of attraction of a β-stable law with β ∈ (0,1).
- Assume that there exists $\epsilon_n, \overline{\epsilon}_n \to 0$

$$\epsilon_n \mathcal{H}([n\cdot]) \Longrightarrow \mathcal{H}_{\infty}(\cdot), \overline{\epsilon}_n \mathbb{H}([n\cdot]) \Longrightarrow \mathbb{H}_{\infty}(\cdot),$$

There exists $\tilde{\epsilon}_n
ightarrow 0$,

$$(\epsilon_n \mathbb{H}([n \cdot]), \epsilon_n \mathbb{C}([n \cdot])) \rightarrow (\mathbb{H}_{\infty}, \mathbb{C}_{\infty}),$$

in the sense of f.d.d. such that \mathbb{H}_{∞} and \mathbb{C}_{∞} are independent. • Let $\overline{\mathbb{C}}(t)$ be the date of birth (height) of the individual visited at time t. Then

$$(\epsilon_n \mathbb{H}, \tilde{\epsilon}_n \mathbb{\bar{C}}) \rightarrow (\mathbb{H}_{\infty}, \mathbb{H}_{\infty} \circ \Gamma^{-1}),$$

(again in the sense of f.d.d.) where Γ is a β -stable subordinator independent of \mathbb{H}_{∞} .

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- E(V), E(y) < ∞: the chronology is recovered from the genealogy by a deterministic space-time change.
- E(V), E(y) = ∞ by E(|P|²) < ∞: the chronology is obtained by a random dilation of the genealogy and a random time change.

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- $\mathcal{H}(n)$: number of ancestors of n
- $y(k) = age of the k^{th}$ ancestor when it begets the $(k 1)^{th}$ ancestor.
- The number of ancestors and the length can be computed from the Lukasiewicz path. S(0) = 0 and

$$S(n+1) - S(n) = |\mathcal{P}_n| - 1.$$

Random walk with $\mathbb{E}(S(n+1) - S(n)) = 0$ and negative increments of size 1.

Fundamental decomposition of the spine

Results

Contour process

• Define {*T*(*k*)} the sequence of (weak) ascending ladder times:

Spine decomposition

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$$T(0) = 0; \text{ and for } k \ge 1$$

 $T(k+1) = \inf\{k > T(k) : S(k) = \sup_{\{0, \dots, k\}} S\}.$

 Define R(k) as the undershoot upon reaching the running maximum at time T(k) incremented by 1 unit

for
$$k \ge 1$$
, $R(k) = S(T(k-1)) - S(T(k)-1) + 1$





The dual walk and the genealogical height process.

•
$$\omega = (\mathcal{P}_k, V_k)_{k \in \mathbb{Z}}$$

• $\vartheta^n(\omega) = (\mathcal{P}_{n-1-k}, V_{n-1-k})_{k \in \mathbb{Z}}$

Dual walk at n:

$$S \circ \vartheta^n = (S(n) - S(n-k); k \ge 0),$$

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• $\mathcal{A}(n)$ indices of the ancestors of n.

$$\mathcal{A}(n) = \{n - T(k) \circ \vartheta^n : T(k) \circ \vartheta^n \le n\}.$$

• $\mathcal{H}(n) = |\{k \le n : T(k) \circ \vartheta^n \le n\}|$

Contour process Results

The dual walk and the chronological height process

- $A(\mathcal{P}, k)$: location of the *r*th atom on \mathcal{P} .
- y(k) = A(P_{T(k)-1}, R(k)) random functional of the ladder height process.
- $y(k) \circ \vartheta^n$ is the contribution of the k^{th} ancestor: age of the k^{th} when it begets the $(k-1)^{th}$ ancestor.
- The pair $(\mathcal{H}(n) \ , \ \mathbb{H}(n))$ is equal to

$$\left(\left| \{k \leq n : T(k) \leq n\} \right|, \sum_{k \leq n: T(k) \leq n} y(k) \right) \circ \vartheta^n.$$



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• From standard excursion theory, the process

$$\left(\left(T(k) \ , \ \sum_{i=1}^{k} y(i)\right); \ k \ge 0\right)$$

defines a bivariate renewal process.

- In general, the difficulty stems from the fact that those two renewal processes are correlated in a non-trivial way.
- $\mathbb{E}(y) < \infty$: $\sum_{k=1}^{n} y(k) \sim n\mathbb{E}(y)$
- $\mathbb{E}(|\mathcal{P}^2|) < \infty$: the two subordinators become independent.

Long edges and large number of offsprings

Results

Contour process

Start from a P.P.P. {(t_i, P_{ti})}_{ti∈l} on ℝ × M with intensity measure dt × λ(dP).

Spine decomposition

- Let π be the push-forward measure of λ by the map $(t, \mathcal{P}) \rightarrow (t, |\mathcal{P}|)$ and assume that $\int_0^\infty x \wedge x^2 \pi(dx) < \infty$.
- {(t_i, |P_{ti}|}_{ti∈I} is a Poisson Point Process on ℝ × ℝ⁺ with intensity measure dt × π(dx).
- From the sequence {(t_i, |P_{ti}|)}_{ti∈I}, one can construct a Lévy process X_t with Laplace exponent

$$\alpha\lambda + \int_0^\infty (e^{-\lambda x} - 1 + \lambda x)\pi(dx)$$

• Choose α, π such that X is of infinite variation $(\int_0^\infty x \pi(dx) = \infty)$ and does not drift to $+\infty$.

3th universality class

- $S_t = \max_{[0,t]} X$
- Let *L* be the local at 0 of S X.
- Construct a discrete sequence from the ascending ladder height process corresponding to X and analogously to the discrete case, we consider

$$\left(L_{u_i}^{-1}, S_{L_{u_i}^{-1}}^{-} - X_{L_{u_i}^{-1}}^{-}, \mathcal{P}_{L_{u_i}^{-1}}^{-}\right)_{t_i \in I}$$

where the t_i 's correspond to the jump times of S.

• This defines a P.P.P. with intensity

$$\mu(dt \ dr \ d\mathcal{P}) = 1_{r \in [0, |\mathcal{P}|]} \lambda(d\mathcal{P}) \mu_r(dt)$$

where μ_r is the law of the $\inf\{u: X_u = -r\}$

• By mimicking the spine decomposition described above, one can construct a continuum analog of the height process.

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• Start P.P.P.
$$\{(t_i, \mathcal{P}_{t_i})\}_{i \in I}$$
.

•
$$X \circ \vartheta^t = X(t) - X(t - \cdot)$$
: dual Lévy process

•
$$A(\mathcal{P}, x) = \sup\{u : \mathcal{P}([u, \infty)) \geq x\}.$$

• At jump times of S,

$$y(u) = A(\mathcal{P}_u, S_u^- - X_u^-)$$

• $\mathbb{H}(t) = \left(\sum_{u \le t : \Delta S_u > 0} y(u)\right) \circ \vartheta^t$. (Random transformation of the Lévy process $X \circ \vartheta^t$.)

•
$$\mathcal{H}(t) = \left(\sum_{u \leq t : \Delta S_u > 0} 1\right) \circ \vartheta^t$$
 (Duquesne Le Gall)

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- Convergence in the long edges and large number of offspring case.
- Ray-Knight type of theorems.
 - Local time for generalized height processes.
 - Can we identify those local times with an underlying continuum branching processes (measure valued process).

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Thank you !