

**A disintegration theorem for  
infinite variance superprocesses**

*Marcella Capaldo*

*joint work with*

*Jochen Blath and Alison Etheridge*

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**Superprocesses** as diffusion approximation of spatial population models:

- Dawson-Watanabe (DW) superprocess

Branching Brownian motion

Population size evolves randomly

- Fleming-Viot (FV) superprocess

“Spatial” Moran model

Population size is fixed

*Feller rescaling:*

Mass of each individual  $\sim 1/n$

Branching/Sampling rate speeded by  $n$

## **Perkins' Disintegration Theorem.**

If we normalize a DW superprocess  $X_t$  by its total mass  $|X_t|$  (stopped before extinction) we recover a FV superprocess with time varying sampling rate  $\sim 1/|X_t|$ .

“Skew-Product” representation for DW superprocess :

$$X_t = |X_t|Z_t$$

$|X_t|$ : Total mass process, given by a Feller diffusion.

$Z_t$ : Time-changed FV superprocess.

$\Rightarrow$  Genealogy of the DW superprocess can be linked via a time change to that of the FV superprocess.

## **Donnelly-Kurtz construction.**

Simultaneous representation of measure-valued population models and their genealogy.

Countable representation.

Relabelling according to the longest line of descent.

Infinite variance setting:

[Etheridge, Williams]

[Birkner, Möhle, Wakolbinger]

Offspring generating function:

$$\Phi(s) = \frac{1}{1 + \beta} (1 - s)^{1 + \beta} + s, \quad 0 < \beta \leq 1$$

$$\text{Tails} \sim n^{-1 - \beta}.$$

$\Rightarrow$  Stable branching mechanism of index  $1 + \beta$ .

Population size described by a CSBP with unbounded variation.

## Exchangeable $\Lambda$ -coalescents.

[Pitman97], [Sagitov99]

Multiple mergers of ancestral lineages occur

If there are  $b$  lineages, then a  $k$ -tuple of lineages merges with rate :

$$\lambda_{b,k} = \int_0^1 x^{k-2} (1-x)^{b-k} \Lambda(dx)$$

where  $\Lambda$  is a finite measure on  $[0, 1]$ .

- No *simultaneous* multiple mergers are allowed.
- $\Lambda = \delta_0 \rightarrow$  Kingman coalescent.

## Generalized Fleming-Viot processes.

[Bertoin, LeGall]

$(\rho_t, t \geq 0) \in \mathcal{M}_1(\mathbb{R}^d)$

At a branching event:

$$\rho_{t-} \rightarrow (1 - X)\rho_{t-} + X\delta_Y$$

$X \sim \nu(\cdot)/\nu(]0, 1])$ ,  $\nu$  finite measure on  $]0, 1]$ .

$Y \sim \rho_{t-}$ .

Generalized Fleming-Viot (GFV) processes can be interpreted as diffusion approximation of Cannings' models.

Generator for  $\rho_t$ :

$$\mathcal{L}G(\rho_t) = \int \nu(dx) \int \rho_t(da) G((1-x)\rho_t + x\delta_a) - G(\rho_t)$$

The GFV process appears as *measure-valued dual* of the  $\Lambda$ -coalescents.

- $\nu(dx) = x^{-2}\Lambda(dx)$

$X_t$  : Infinite variance superprocess

We study the normalized process (stopped before extinction):

$$Y_t = X_t/|X_t| \in \mathcal{M}_1(\mathbb{R}^d)$$

$Y_t$  is a *time-changed GFV process*, with coalescent measure  $\Lambda = \text{Beta}(1 - \beta, 1 + \beta)$ .

Generator for  $Y_t$  (ignoring spatial component):

$$\mathcal{G}f(Y_t) = \frac{C}{|X_t|^\beta} \mathcal{L}f(Y_t)$$

The jumping rate is function of the total mass.

## A Disintegration Theorem for infinite variance superprocesses.

“Skew-product” representation :

$$X_t = |X_t|Y_t$$

$|X_t|$ : Total mass process, given by a CSBP.

$Y_t$ : Time-changed GFV process.

$\Rightarrow$  Genealogy of a sample from an infinite variance superprocess can be represented in terms of a  $\Lambda$ -coalescent, with rates rescaled by its total mass.

For more general branching mechanisms, the jump measure  $\nu$  will depend on  $|X_t|$  in a complicated way.

$\Rightarrow$  “Skew-product” representation *only in the stable case*.