# Tree valued spatial $\Lambda$ -Cannings and $\Lambda$ -Fleming-Viot dynamics

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#### A resampling model

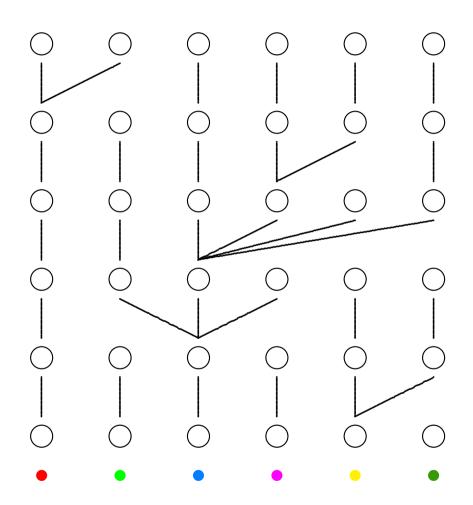
#### Type space K, compact

We consider a multi-type asexual population of fixed size N.

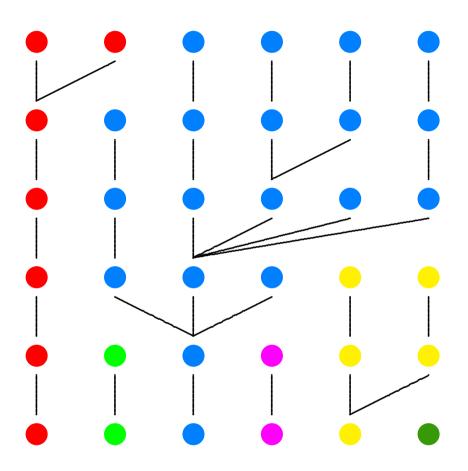
For each  $k \in \{2, ..., N\}$  at rate  $\lambda_{N,k}$ ,

- a k-tuple  $\{i_1,...,i_k\}$  of individuals is killed, and
- replaced by k copies of the individual  $i_{\ell}$  chosen at random among  $\{i_1, ..., i_k\}$ . That is, the offspring inherits the type from  $i_{\ell}$ .

# $\Lambda\text{-Cannings dynamics}$



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#### **Consequences of consistency**

**Consistency.** (= same dynamics is observed in any sample)

$$\lambda_{N,k} = \lambda_{N+1,k} + \lambda_{N+1,k+1}.$$

#### Pitman 1999, Sagitov 1999

There exists a finite measure  $\Lambda$  on [0,1] with

$$\lambda_{N,k} := \int_0^1 \Lambda(\mathrm{d}x) \, x^{k-2} (1-x)^{n-k}.$$

#### **Examples of** $\Lambda$ **-Cannings.**

 $\Lambda = \delta_0$  (Kingman coalescent);  $\Lambda = \delta_1$  (star-shaped)

#### From particle model to diffusion limits

#### Interesting functional.

 $X_t^{N,\Lambda}:=$  empirical type distribution at time t

Bertoin & Le Gall (2003)

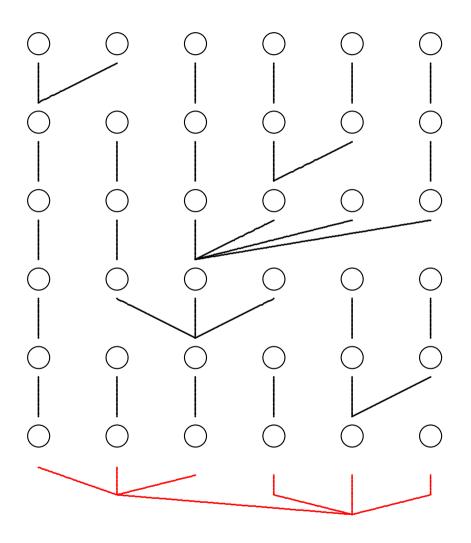
Measure-valued process  $(N \to \infty)$ .  $X^{\Lambda}$  is a strong Markov process with values in  $\mathcal{M}_1(K)$  whose generator acts on functions of the form

$$\mu \mapsto \prod_{i=1}^{n} \langle \mu, \psi_i \rangle = \langle \mu^{\otimes n}, \prod_{i=1}^{n} \psi_i \rangle$$

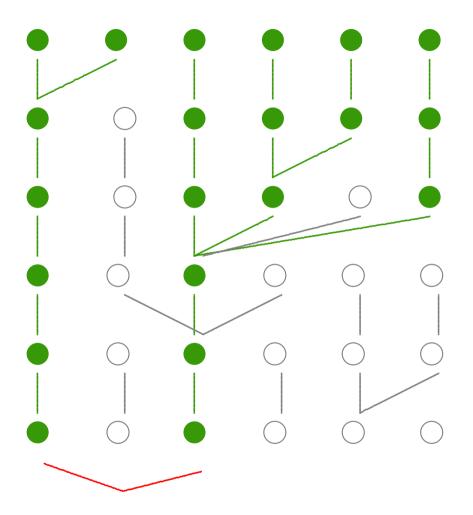
as follows:

$$\Omega_{\Lambda-\text{FV}} \prod_{i=1}^{n} \langle \cdot, \psi_i \rangle (\mu) 
= \sum_{\substack{J \subseteq \{1, 2, ..., n\} \\ \#J > 2}} \lambda_{n, \#J} (\langle \cdot, \prod_{j \in J} \psi_j \rangle - \prod_{j \in J} \langle \cdot, \psi_j \rangle) \cdot \prod_{i \in \{1, 2, ..., n\} \setminus J} \langle \cdot, \psi_i \rangle (\mu).$$

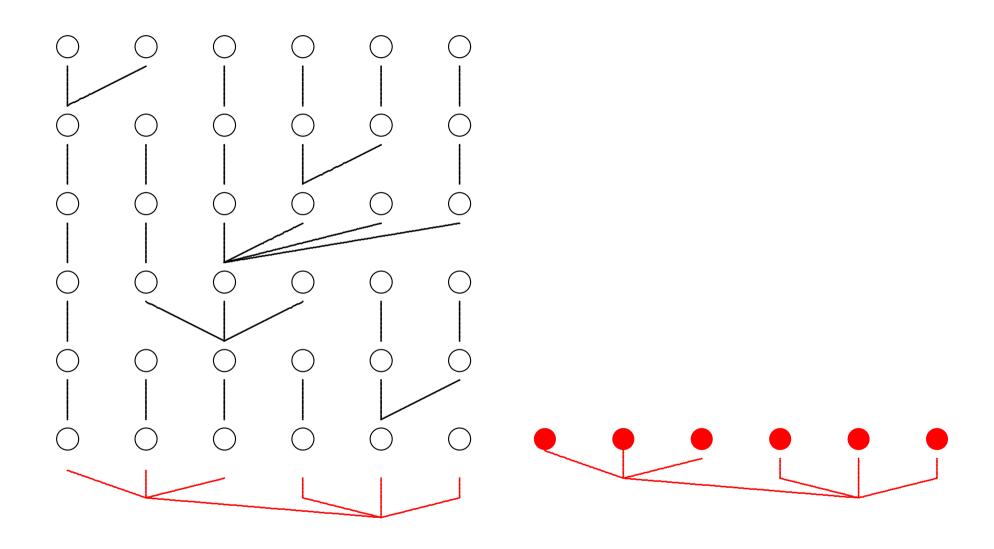
### **Tracing back ancestry**

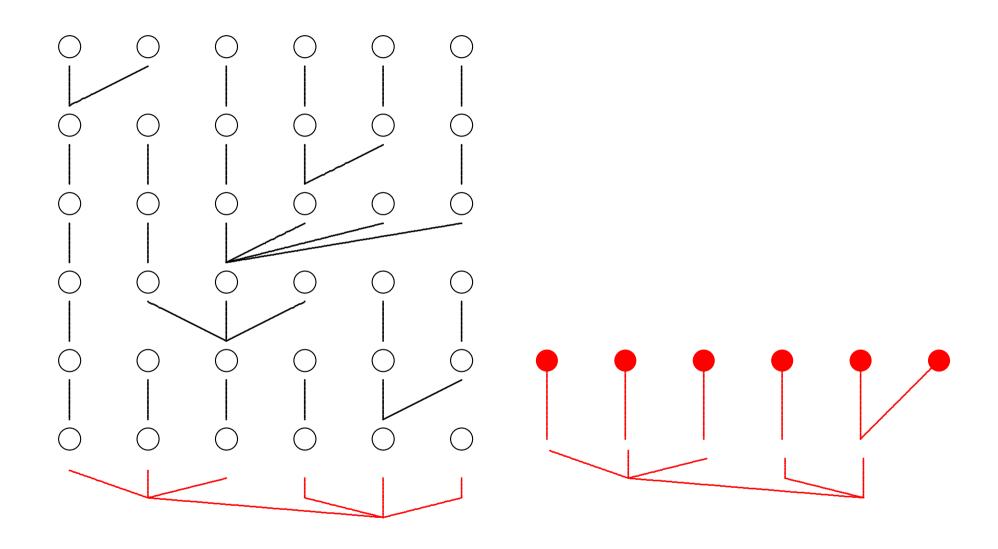


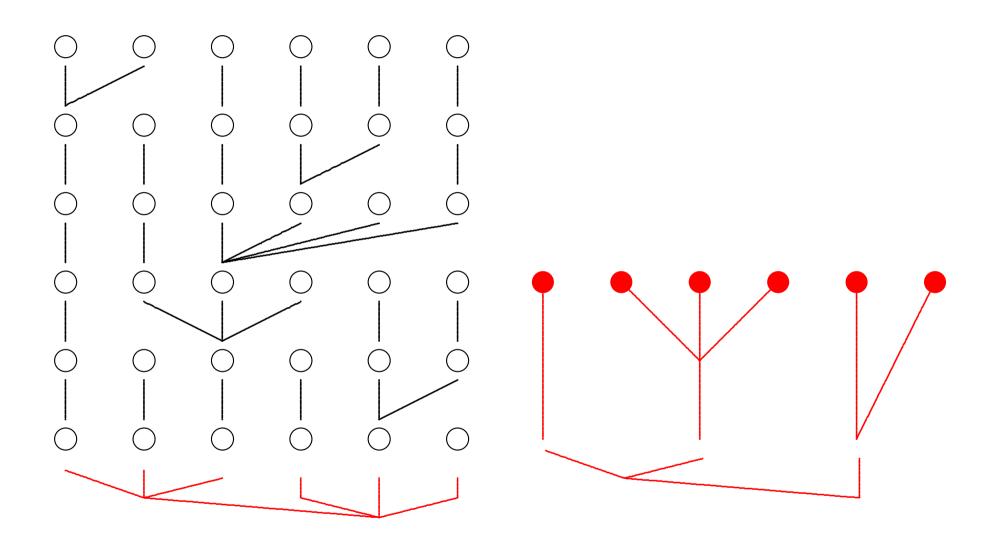
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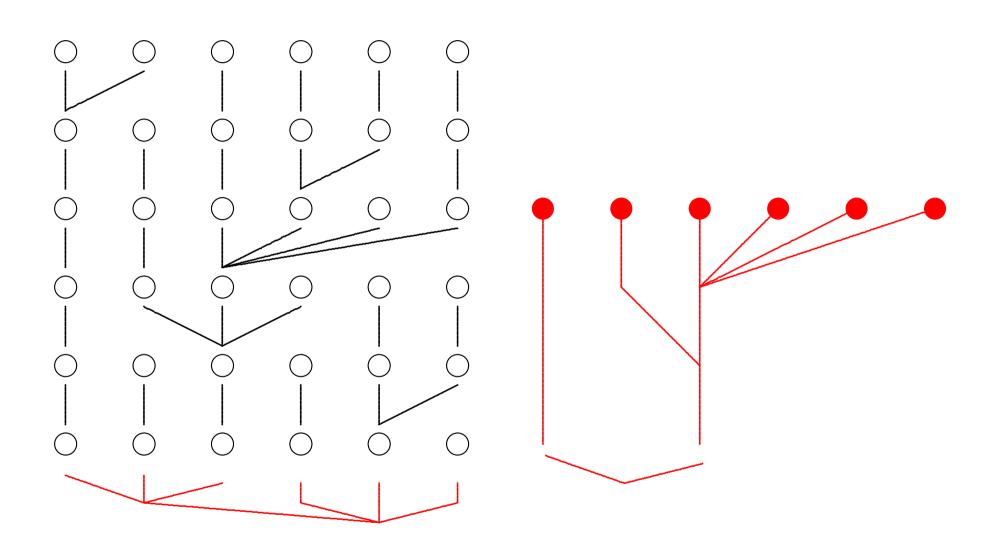


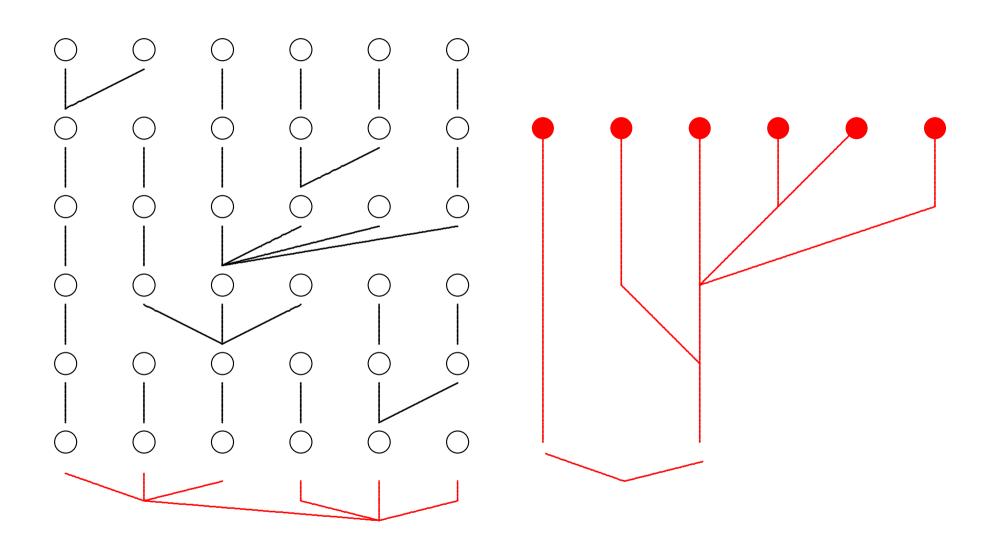
 $\Lambda$ -coalescent (in backward picture) given n ancestral lines, k of them merge at rate  $\lambda_{n,k}$ 

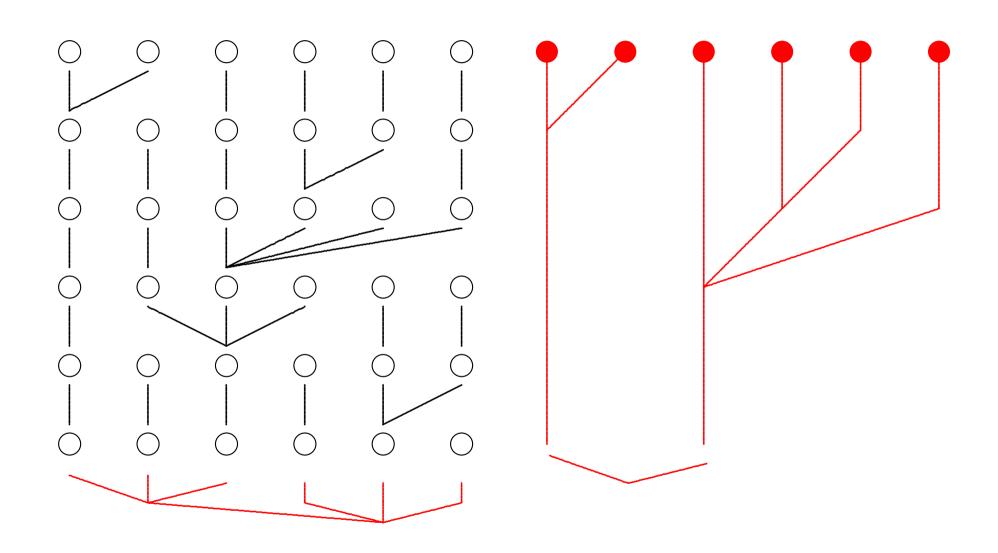












#### **Encoding genealogies ...**

We aim to describe the **genealogical tree** of the **whole population** while making ancestral lines of **all possible samples explicit**.

We encode our genealogies by 
$$(U, r, \mu)$$

set of individuals

genealogical distances

genealogical distances

appling measure

and evaluate samples via test functions of the form

$$\Phi^{n,\phi}(U,r,\mu) := \int_{U^n} \mu^{\otimes n}(\underline{\mathrm{d}}\underline{u}) \,\phi\big((r(u_i,u_j))_{1 \le i < j \le n}\big).$$

Such test functions are referred to as polynomials.

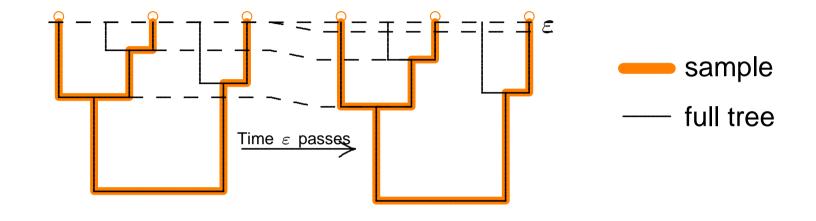
#### The state space: more formal

 $\mathbb{U} := \{\text{isometry classes of ultra metric probability spaces}\}.$ 

Gromov (2000); Greven, Pfaffelhuber & Winter (2009)

We equip U with the **Gromov-weak topology** which means convergence in the sense of **convergence of all polynomials** (with continuous bounded test functions).

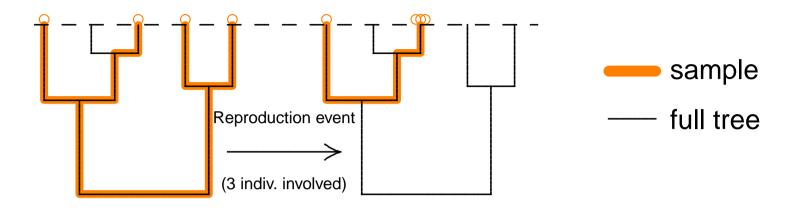
#### Tree growth



$$\Omega_{\text{growth}}^{N,\Lambda} \Phi(U, r, \mu) 
= 2 \int_{U^n} \mu^{\otimes n} (\underline{d}\underline{u}) \sum_{1 \leq i < j \leq n} \frac{\partial \phi}{\partial r_{i,j}} ((r(u_i, u_j))_{1 \leq i < j \leq n}) + \mathcal{O}(\frac{1}{N}),$$

where the error term comes from multiples in a sample.

#### Reproduction



$$\begin{split} &\Omega_{\text{repro}}^{N,\Lambda} \Phi(U,r,\mu) \\ &= \sum_{J \subset \{1,2,...,n\}, \#J > 2} \lambda_{n,\#J} \int_{U^n} \mu^{\otimes n} (\mathrm{d}u) \, \frac{1}{\#J} \sum_{j_0 \in J} \big\{ R_J^{j_0} \phi - \phi \big\} \big( (r(u_i,u_j))_{1 \le i < j \le n} \big) + \mathcal{O} \big( \frac{1}{N} \big) \end{split}$$

with the replacement operator

$$R_J^{j_0} \phi \left( (r_{i,j})_{1 \le i < j \le n} \right) := \phi \left( (r_{\tilde{i}, \tilde{j}})_{1 \le i < j \le n} \right)$$

where for all  $1 \leq i \leq n$ ,

$$ilde{i} := \left\{ egin{array}{ll} j_0, & ext{if } i \in J, \ i, & ext{if } i 
ot\in J. \end{array} 
ight.$$

#### The tree-valued generalized $\Lambda$ -FV

Consider the limiting operator

$$\begin{split} \Omega^{\Lambda} \Phi \big( U, r, \mu \big) &:= \Omega^{\Lambda}_{\text{repro}} \Phi (U, r, \mu) + \Omega^{\Lambda}_{\text{growth}} \Phi (U, r, \mu) \\ &= \sum_{J \subseteq \{1, 2, ..., n\}, \#J \ge 2} \lambda_{n, \#J} \int_{U^n} \mu^{\otimes n} (\mathrm{d}u) \, \frac{1}{\#J} \sum_{j_0 \in J} \big\{ R_J^{j_0} \phi - \phi \big\} \big( (r(u_i, u_j))_{1 \le i < j \le n} \big) \\ &+ 2 \int_{U^n} \mu^{\otimes n} (\mathrm{d}\underline{u}) \sum_{1 < i < j < n} \frac{\partial \phi}{\partial r_{i,j}} \big( (r(u_i, u_j))_{1 \le i < j \le n} \big) \end{split}$$

acting on the set

 $\Pi^1 :=$  polynomials with differentiable, bounded test functions.

Theorem 1. (Greven, Klimovsky & W.) Let  $\mathbf{P_0}$  be a probability measure on  $\mathbb{U}$ . The  $(\mathbf{P_0}, \Omega^{\Lambda}, \Pi^1)$ -martingale problem is well-posed provided that the "dust-free" property holds, i.e.,  $\int_0^1 \Lambda(\mathrm{d}x) \frac{1}{x} = \infty$ . The solution  $\mathcal{U}^{\Lambda}$  is a strong Markov process with the Feller property.

#### **Existence: Particle Approximation**

Theorem 2. (Greven, Klimovsky & W.) Let  $\mathcal{U}^{N,\Lambda}$  the tree-valued  $\Lambda$ -Cannings dynamics with population size N. Assume that the initial conditions convergence in  $\mathcal{U}_0 \in \mathbb{U}$ . Then

$$(\mathcal{U}_t^{N,\Lambda})_{t\geq 0} \underset{N\to\infty}{\Longrightarrow} (\mathcal{U}_t^{\Lambda})_{t\geq 0}.$$

**Keys in proof.** Convergence of generators + Compact containment

For population models with "ancestor-descendant" relationships whose type proportions evolve as a martingale, **compact containment follows** already **from convergence of 1-dimensional marginals**.

#### The $\Lambda$ -coalescent measure tree

- run a Λ-coalescent
- ullet equip  $\mathbb N$  with genealogical distance,  $r_{\mathrm{gen}}$  (and complete as a metric space)

For each  $n \in \mathbb{N}$  put, consider

$$\mathcal{U}_n^{\Lambda,\downarrow} := \left( (\bar{\mathbb{N}}, r_{\mathrm{gen}}), \frac{1}{n} \sum_{i=1}^n \delta_i \right)$$

Gromov (2000); Greven, Pfaffelhuber & W. (2009)

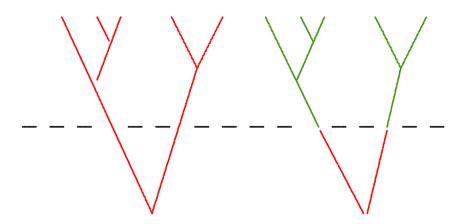
**Fact.** Provided the dust-free property holds there exists  $\mathcal{U}_{\infty}^{\Lambda,\downarrow} \in \mathbb{U}$  such that

$$\mathcal{U}_n^{\Lambda,\downarrow} \underset{n\to\infty}{\Longrightarrow} \mathcal{U}_{\infty}^{\Lambda,\downarrow}.$$

 $\mathcal{U}_{\infty}^{\Lambda,\downarrow}$  is called  $\Lambda$ -coalescent measure tree.

#### **Uniqueness of MP = Tree-valued duality**

generalized  $\Lambda$ -FV  $(U, r^{\uparrow}, \mu)_t$  dual to  $\Lambda$ -coalescent  $(K, r^{\downarrow})_t$ 



Greven, Pfaffelhuber, W. (2012)

Theorem 3. [Greven, Klimovsky, W.]

$$\mathbf{E} \Big[ \int \mu_t^{\otimes n} (\mathrm{d}\underline{u}) \phi \big( (\mathbf{r}_t^{\uparrow}(\mathbf{u}_i, \mathbf{u}_j))_{1 \leq i < j \leq n} \big) \Big]$$

$$= \mathbf{E} \Big[ \int \prod_{\varpi \in K_t} \mu_0 (\mathrm{d}v_{\varpi}) \phi \big( (\mathbf{r}_t^{\downarrow}(i, j) + \mathbf{r}_0^{\uparrow}(\mathbf{v}_{\varpi(i)}, \mathbf{v}_{\varpi(j)}))_{1 \leq i < j \leq n} \big) \Big]$$

#### The infinitely old population

Theorem 4. (Greven, Klimovsky & W.) Assume that  $\Lambda$  satisfies the dust-free property. Then

$$\mathcal{U}_t^{\Lambda} \underset{t \to \infty}{\Longrightarrow} \mathcal{U}_{\infty}^{\Lambda,\downarrow}.$$

**Proof.** It is enough to show that

$$\mathbb{E}\left[\Phi\left(\mathcal{U}_{t}^{\Lambda}\right)\right]\underset{t\to\infty}{\longrightarrow}\mathbb{E}\left[\Phi\left(\mathcal{U}_{\infty}^{\Lambda,\downarrow}\right)\right],$$

for all polynomials  $\Phi \in \Pi^1$ .

This, however, follows by duality.

# Adding mutation

#### A resampling model with mutation

Type space K, compact

We consider a multi-type asexual population of fixed size N.

For each  $k \in \{2, ..., N\}$  at rate  $\lambda_{N,k}$ ,

- k-individuals  $\{i_1, ..., i_k\}$  are killed, and
- are replaced by k copies of the individual  $i_{\ell}$  chosen at random among  $\{i_1, ..., i_k\}$ . That is, the offspring inherits the type from  $i_{\ell}$ .

For each individual of type x, at rate m

• the type **mutates** from x to y with probability M(x, dy).

#### Enriching the state space with types ...

We aim to describe the **genealogical tree** of the **whole population** while making ancestral lines and types of all possible samples explicit.

We encode our genealogies by 
$$(U, r, \mu, \kappa)$$

set of individuals set of individuals sampling measure genealogical distances sampling measure samples via test functions of the form

and evaluate samples via test functions of the form

$$\Phi^{n,\phi,f}(X,r,\mu,\kappa) := \int_{U^n} \mu^{\otimes n}(\underline{\mathrm{d}}\underline{u}) \big( (\phi \circ \underline{\underline{r}}) \cdot (f \circ \kappa) \big) (\underline{u}).$$

with

$$\underline{\underline{r}} : \underline{u} \mapsto (r(u_i, u_j))_{1 \le i < j \le n}$$

#### The state space including types: more formal

 $\mathbb{U}^K := \{ \text{mark function invariant isometry classes of ultra metric probability spaces} \}.$ 

Depperschmidt, Greven & Pfaffelhuber (2011)

We equip  $\mathbb{U}^K$  with the marked Gromov-weak topology which means convergence in the sense of convergence of all polynomials (with continuous bounded test functions).

#### Well-posed martingale problem

#### Consider the operator

$$\Omega^{\Lambda,M}\Phi\big(U,r,\mu,\kappa\big):=\Omega_{\mathrm{repro}}^{\Lambda,M}\Phi(T,r,\mu,\kappa)+\Omega_{\mathrm{growth}}^{\Lambda,M}\Phi(T,r,\mu,\kappa)+\Omega_{\mathrm{mutation}}^{\Lambda,M}\Phi(T,r,\mu,\kappa)$$

acting on the set

 $\Pi^1 := \text{polynomials with "smooth" bounded test functions } \phi \text{ and } f$ ,

where

$$\begin{split} &\Omega_{\mathrm{mut}}^{\Lambda,M} \Phi(T,r,\mu,\kappa) \\ &:= m \cdot \int \mu^{\otimes n} (\mathrm{d}u) \, \phi \circ \underline{\underline{r}}(\underline{u}) \cdot \int_{K} M(\kappa(u_{i}),\mathrm{d}y_{i}) \big\{ f(\kappa(u_{1}),...,y_{i},...,\kappa(u_{n})) - f(\kappa(u_{1}),...,\kappa(u_{i}),...,\kappa(u_{n})) \big\}. \end{split}$$

Theorem 5. (Greven, Klimovsky & W.) Let  $\mathbf{P_0}$  be a probability measure on  $\mathbb{U}^K$ . The  $(\mathbf{P_0}, \Omega^{\Lambda,M}, (\Pi^K)^1)$ -martingale problem is well-posed provided that the "dust-free" property holds. The solution  $\mathcal{U}^{\Lambda,M}$  is a strong Markov process with the Feller property.

#### The equilibrium with mutation

Assume that  $M(\cdot, \cdot)$  is **ergodic**, i.e., there is a probability measure  $\pi \in \mathcal{M}_1(K)$  with  $\pi M = \pi$  and  $M^{(n)}(x, \cdot) \Longrightarrow_{n \to \infty} \pi$ , for al  $x \in K$ .

Theorem 6. Greven, Klimovsky & W. Under these assumptions, the sequence

$$\left(\mathcal{U}_t^{\Lambda,M}\right)_{t\geq 0}$$

converges for all initial  $\mathcal{U}_0^{\Lambda,M}$  , as  $t \to \infty$  .

The equilibrium  $\mathcal{U}_{\infty}^{\Lambda,M,\downarrow}$  can be represented by first realizing the the  $\Lambda$ -coalescent tree  $\mathcal{U}_{\infty}^{\Lambda}$ , and then given the latter, realizing a tree-indexed mutation random walk in equilibrium.

**Note.** The rate of convergence will dependent on the measure  $\Lambda$ .

Question. Do the tree-length decrease stochastically in  $\Lambda$ ?

# Adding migration

#### The spatial $\Lambda$ -Cannings model

#### Geographic space. G, discrete

We consider a multi-type asexual population of fixed size N which individuals placed at a site  $x \in G$ 

- Migration. The individuals perform independently rate 1 random walks with transition kernel a(x,y)
- Reproduction. At each site  $x \in G$ , for each  $k \in \{2, ..., N\}$  at rate  $\lambda_{N,k}$ ,
  - k-individuals  $\{i_1, ..., i_k\}$  currently situated in G are killed, and
  - replaced by k copies of the individual  $i_{\ell}$  chosen at random among  $\{i_1, ..., i_k\}$ . That is, the offspring inherits the type from  $i_{\ell}$ .
- Mutation. For each individual of type x, at rate m, the type mutates from x to y with probability M(x, dy).

#### ... and its dual spatial $\Lambda$ -coalescent

**Spatial**  $\Lambda$ -coalescent is a strong Markov process which takes values in the set of partitions of all individuals where each partition element is assigned a site in G such that any "locally finite" subpopulation/-partition behaves as follows:

- Migration. Partition elements change their position according to a rate 1 random walk with transition probabilities  $\bar{a}(x,y) := a(y,x)$ .
- $\Lambda$ -coalescence. Each local partition performs a  $\Lambda$ -coalescent.

#### Constructions of the $\Lambda$ -coalescent.

- Limic & Sturm (2006) for finite G
- via Donnelly & Kurtz (1990ies)'s look-down

#### **Observing genealogies**

For observing the genealogies as marked metric measure spaces, we have different choices:

- Global point of view (*G* finite necassary). One sampling measure for the whole population.
  - Start with locally finite populations on a finite G.
  - Take the uniform distribution  $\mu$  on all individuals.
  - Let the local intensity tend to infinity.
- Local point of view. One sampling measure for each local population.
  - Start with locally finite populations on possible infinite G.
  - Take in each site  $x \in G$  the uniform distribution  $\mu_x$  on all individuals placed at site x.
  - Let the local intensity tend to infinity.

# Global point of view

#### Well-posed martingale problem; G finite

#### Consider the operator

$$\Omega^{\Lambda,M,a} = \Omega^{\Lambda,M} + \Omega^a_{\text{migration}}$$

with  $\kappa:U\to K\times G$  and

$$\Omega_{\text{migration}}^{a} \Phi^{n,\phi,f} (U, r, \mu, \kappa) := \int \mu^{\otimes n} (du) \, \phi \circ \underline{\underline{r}}(\underline{u}) \cdot \sum_{i=1}^{n} A^{(i)} f \circ \kappa(\underline{u})$$

and  $A^{(i)}$  being the generator of a single individual random walk acting on the  $n^{\text{th}}$  individual in the sample.

Theorem 7. (Greven, Klimovsky & W.) For each initial tree in  $\mathbb{U}^{K\times G}$ , the  $(\Omega^{\Lambda,M,a},(\Pi^{K\times G})^1)$ -martingale problem is well-posed.

Call its solution  $\mathcal{U}^{\Lambda,M,a}$  the spatial tree-valued  $\Lambda$ -Fleming-Viot.

#### "Wrapping around torus"; $d \ge 3$

$$G_N := [-N, N]^d \cap \mathbb{Z}^d, \ a_N(x, y) := \sum_{z: z=y \bmod G_N} a(0, z)$$

 $\widehat{\mathcal{U}}^{\Lambda,a_N} := \text{rescaled tree-valued spatial } \Lambda\text{-Fleming-Viot dynamics:}$ 

- speed up time by a factor  $(2N+1)^d$
- scale down distances by a factor of  $(2N+1)^{-d}$

$$\kappa := 2 \cdot \left(\rho + \frac{2}{\lambda_{2,2}}\right)^{-1}, \qquad \rho := \text{escape probability on } \mathbb{Z}^d$$

Greven, Limic & W. (2005), Limic & Sturm (2006)

For all 
$$t > 0$$
,  $\Phi \in \Pi^1$ ,  $\mathbb{E}\left[\Phi\left(\widehat{\mathcal{U}}_t^{\Lambda, a_N}\right)\right] \xrightarrow[N \to \infty]{} \mathbb{E}\left[\Phi\left(\mathcal{U}_t^{\kappa \delta_0}\right)\right]$ .

Theorem 8. (Greven, Klimovsky and W.) If the initial states converges in  $\mathbb{U}$  and  $\sum_{x \in \mathbb{Z}^d} \hat{a}(0,x)|x|^{2+d} < \infty$ , then

$$(\widehat{\mathcal{U}}_t^{\Lambda,a_N})_{t\geq 0} \underset{N\to\infty}{\Longrightarrow} (\mathcal{U}_t^{\kappa\delta_0})_{t\geq 0}.$$

proof uses techniques from Dawson, Greven & Vaillancourt (1995)

#### Diffusive clustering of genealogies in d=2

$$G_N := [-N, N]^2 \cap \mathbb{Z}^2, \ a_N(x, y) := \sum_{z: z=y \bmod G_N} a(0, z), \ \alpha \in (0, 1]$$

 $\widehat{\mathcal{U}}^{\Lambda,a_N} := \text{rescaled tree-valued spatial } \Lambda\text{-Fleming-Viot dynamics:}$ 

- observe at times  $N^{\frac{2}{\alpha}}$
- stretch distances (non-linearly) via  $\tau^{\alpha}(s) := \frac{\log(1+s)}{2\log N}$

Greven, Limic & W., Heuer & Sturm

For all 
$$\alpha \in (0,1]$$
,  $\Phi \in \Pi^1$ ,  $\mathbb{E}\left[\Phi(\widehat{\mathcal{U}}_{\alpha}^{\Lambda,a_N})\right] \xrightarrow[N \to \infty]{} \mathbb{E}\left[\Phi(\mathcal{U}_{-\log \alpha}^{\delta_0})\right]$ .

Theorem 9. (Greven, Klimovsky and W.) If the initial states converges in  $\mathbb{U}$  and  $\sum_{x \in \mathbb{Z}^d} \hat{a}(0,x)e^{\lambda x} < \infty$  for some  $\lambda > 0$ , then

$$(\widehat{\mathcal{U}}_{\alpha}^{\Lambda,a_N})_{\alpha\in(0,1]} \overset{\text{f.d.d.}}{\Longrightarrow} (\mathcal{U}_{-\log\alpha}^{\delta_0})_{\alpha\in(0,1]}.$$

# Local point of view

#### Infinite geographic space

Countable infinite geographic space requires  $\sigma$ -finite sample measures.

**Localization.** Fix a sequence  $G_n \uparrow G$  with  $\#G_n < \infty$ . We refer to  $(U, r, \mu, \kappa)$  as marked mm-space iff for every  $n \in \mathbb{N}$ , the restriction  $(U_n, r_n, \kappa_n)$  to all individuals with a spatial mark in  $G_n$  together with  $\mu_n := \frac{1}{\#G_n} \mu\big|_{U_n}$  is a marked metric probability space.

Define the spatial tree-valued  $\Lambda$ -Fleming-Viot

$$(\mathcal{U}_t^{\Lambda,M,a})_{t\geq 0} := ((U_t, r_t, \{\mu_t^x; x \in G\}, \kappa_t))_{t\geq 0}$$

via the look-down process and local approximation.

#### The associated measure-valued $\Lambda$ -Fleming-Viot

 $\overline{\mathcal{U}^{\Lambda,M}}$ , a; tree-valued  $\Lambda$ -Fleming-Viot.

Put for each  $x \in G$ ,

$$X_t^{\Lambda,M,a,x} := \mu_t \{ u \in U_t : \kappa_t(u) = \cdot \times \{x\} \} \in \mathcal{M}_1(K).$$

Theorem 10. (Greven, Klimovsky & W.) Assume that the underlying symmetrized random walk is irreducible, and that  $\{X_0^{\Lambda,M,a,x}; x \in G\}$  is translation invariant and ergodic with intensity  $\theta \in \mathcal{M}_1(K)$ . Then there is a translation invariant measure  $\nu_{\theta}$  with intensity  $\theta$  such that

$$X_t^{\Lambda,M,a} \Longrightarrow_{t\to\infty} \nu_{\theta}.$$

#### **Dichotomy**

Transient symmetrized walks imply  $\nu_{\theta}$  is spatially ergodic.

Recurrent symmetrized walk.  $\nu_{\theta} = \int_{K} \theta(\mathrm{d}k) (\delta_k)^{\otimes G}$ .

#### **Tree-valued equilibrium (including mutation)**

since distances can tend to  $\infty$ , put  $\tilde{r}_t(x,y) := 1 - e^{-r(x,y)}$ 

write  $\widetilde{U}_t^{\Lambda,M,a}$  for the tree-valued dynamics with shrinked distances

Theorem 10. (Greven, Klimovsky & W.) For every intensity  $\theta \in \mathcal{M}_1(K)$  there is a invariant measure

$$\widetilde{\mathcal{U}}_{\infty}^{\Lambda,M,a,\downarrow,\theta}$$
.

If the associate measure-valued process of the initial state is **translation invariant and ergodic** with intensity measure  $\theta$ , then

$$\widetilde{U}_t^{\Lambda,M,a} \Longrightarrow_{t \to \infty} \widetilde{\mathcal{U}}_{\infty}^{\Lambda,M,a,\downarrow,\theta}.$$

**Representation.** Here again we can read of the genealogiy of an infinitely old population from the look-down representation of the spatial  $\Lambda$ -coalescent measure tree, and then assign every "tree" a type in K by sampling i.i.d. from  $\theta$ .

#### The local finite system scheme; $d \ge 3$

Theorem 11. (Greven, Klimovsky & W.) If the associated measure-valued process is initially translation invariant and ergodic with intensity  $\theta$ , then for all t > 0,

$$\left(\widetilde{\mathcal{U}}_{t\cdot(2N+1)^d+s}^{\Lambda,M,a_N}\right)_{s\geq 0}\underset{N\to\infty}{\Longrightarrow} \mathcal{L}^{\widetilde{\mathcal{U}}_{\infty}^{\Lambda,M,a,\downarrow,\theta_t}}\left[\left(\widetilde{\mathcal{U}}_s^{\Lambda,M,a}\right)_{s\geq 0}\right],$$

where the intensity  $\theta_t$  of the equilibrium equals in law a (non-spatial) rate  $\kappa$ s Fleming-Viot started in  $\theta$ .

# Many thanks