

A spatial population model with random obstacles

4th Cornell Probability Summer School

János Engländer

University of California, Santa Barbara

June 26, 2008

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a)$$

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a)$$

K : Mild obstacle configuration attached to ω .

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a)$$

K : Mild obstacle configuration attached to ω .

K^c : 'Swiss cheese'

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a)$$

K : Mild obstacle configuration attached to ω .

K^c : 'Swiss cheese'

Given ω , we define P^ω as the law of the (strictly dyadic) BBM on \mathbb{R}^d , $d \geq 1$ with spatially dependent branching rate

$$\beta := \beta_1 \mathbf{1}_K + \beta_2 \mathbf{1}_{K^c}$$

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a)$$

K : Mild obstacle configuration attached to ω .

K^c : 'Swiss cheese'

Given ω , we define P^ω as the law of the (strictly dyadic) BBM on \mathbb{R}^d , $d \geq 1$ with spatially dependent branching rate

$$\beta := \beta_1 \mathbf{1}_K + \beta_2 \mathbf{1}_{K^c}$$

The process Z under P^ω is called a **BBM with mild Poissonian obstacles**.

Mild obstacles

Model.

Let ω be a Poisson point process (PPP) on \mathbb{R}^d with intensity $\nu > 0 \sim \mathbf{P}$.
 $a > 0$ and $\beta_2 > \beta_1 > 0$ fixed.

$$K = K_\omega := \bigcup_{x_i \in \text{supp}(\omega)} \bar{B}(x_i, a).$$

K : Mild obstacle configuration attached to ω .

K^c : 'Swiss cheese'

Given ω , we define P^ω as the law of the (strictly dyadic) BBM on \mathbb{R}^d , $d \geq 1$ with spatially dependent branching rate

$$\beta := \beta_1 \mathbf{1}_K + \beta_2 \mathbf{1}_{K^c}.$$

The process Z under P^ω is called a **BBM with mild Poissonian obstacles**.

Total mass process: $|Z|$.

Questions

Questions

- Growth of the **global/local** population size? How much will the absence of branching in K slow the global reproduction down? Change the exponent β_2 ?

Questions

- Growth of the **global/local** population size? How much will the absence of branching in K slow the global reproduction down? Change the exponent β_2 ?
- What are the large deviations?
 [E.g. $P(\text{atypically small population})=?$]

Questions

- Growth of the **global/local** population size? How much will the absence of branching in K slow the global reproduction down? Change the exponent β_2 ?
- What are the large deviations?
 [E.g. $P(\text{atypically small population})=?$]
- How about the **spatial spread**? How will the $\sqrt{2\beta_2}$ speed reduce?

Questions

- Growth of the **global/local** population size? How much will the absence of branching in K slow the global reproduction down? Change the exponent β_2 ?
- What are the large deviations?
 [E.g. $P(\text{atypically small population})=?$]
- How about the **spatial spread**? How will the $\sqrt{2\beta_2}$ speed reduce?

Questions can be asked in 2 diff. ways: **annealed** and **quenched** sense.

Related models in biology

Related models in biology

- (i) **Migration with unfertile areas** (Population dynamics):
Population moves in space and reproduces by binary splitting, but randomly located reproduction-suppressing areas modify the growth.

Related models in biology

- (i) **Migration with unfertile areas** (Population dynamics):
Population moves in space and reproduces by binary splitting, but randomly located reproduction-suppressing areas modify the growth.
- (ii) **Fecundity selection** (Genetics):
Reproduction and mutation. Certain randomly distributed genetic types have low fitness: even though they can be obtained by mutation, they themselves do not reproduce easily, unless mutation transforms them to different genetic types.
[‘Space’ = space of genetic types rather than physical space.]

Questions: (local and global) growth rate of the population? Once one knows the global population size, the model can be normalized by the global population size, giving a population of unit mass; then the question becomes the *shape* of the population.

Questions: (local and global) growth rate of the population? Once one knows the global population size, the model can be normalized by the global population size, giving a population of unit mass; then the question becomes the *shape* of the population.

- **Population dynamics setting:** Is there a preferred spatial location for the process to populate?

Questions: (local and global) growth rate of the population? Once one knows the global population size, the model can be normalized by the global population size, giving a population of unit mass; then the question becomes the *shape* of the population.

- **Population dynamics setting:** Is there a preferred spatial location for the process to populate?
- **Genetic setting:** Existence of a certain kind of genetic type that is preferred in the long run that lowers the risk of low fecundity caused by mutating into less fit genetics types?

Genealogical structure— exciting problem!

E.g. it seems quite possible that for large times the ‘bulk’ of the population consists of descendants of a single particle that

- decided to travel far enough (resp. to mutate many times)
- reached a less hostile environment (resp. in high fitness genetic type area), where she and her descendants can reproduce freely.

Genealogical structure— exciting problem!

E.g. it seems quite possible that for large times the ‘bulk’ of the population consists of descendants of a single particle that

- decided to travel far enough (resp. to mutate many times)

Genealogical structure— exciting problem!

E.g. it seems quite possible that for large times the ‘bulk’ of the population consists of descendants of a single particle that

- decided to travel far enough (resp. to mutate many times)
- reached a less hostile environment (resp. in high fitness genetic type area), where she and her descendants can reproduce freely.

Related phenomenon in marine systems: hypoxic patches form in estuaries because of stratification of the water.

⇒ The patches affect different organisms in different ways but are detrimental to some of them. They appear and disappear in an effectively stochastic way.

Source-sink theory: some patches of habitat are good for a species (and growth rate is positive) whereas other patches are poor (and growth rate smaller, or is zero or negative). Individuals can move between patches randomly or according to more detailed biological rules for behavior.

Source-sink theory: some patches of habitat are good for a species (and growth rate is positive) whereas other patches are poor (and growth rate smaller, or is zero or negative). Individuals can move between patches randomly or according to more detailed biological rules for behavior.

Systems with **periodic local disturbances** like e.g.

- **FORESTS** where trees sometimes fall creating gaps (which have various effects on different species but may harm some)

Source-sink theory: some patches of habitat are good for a species (and growth rate is positive) whereas other patches are poor (and growth rate smaller, or is zero or negative). Individuals can move between patches randomly or according to more detailed biological rules for behavior.

Systems with **periodic local disturbances** like e.g.

- **FORESTS** where trees sometimes fall creating gaps (which have various effects on different species but may harm some)

Source-sink theory: some patches of habitat are good for a species (and growth rate is positive) whereas other patches are poor (and growth rate smaller, or is zero or negative). Individuals can move between patches randomly or according to more detailed biological rules for behavior.

Systems with **periodic local disturbances** like e.g.

- **FORESTS** where trees sometimes fall creating gaps (which have various effects on different species but may harm some)
- **AREAS OF GRASS** or brush which are subject to occasional fires — burned areas can be expected to less suitable habitats for at least some organisms.

Result on population size

Define *average growth rate*:

$$r_t = r_t(\omega) := \frac{\log |Z_t(\omega)|}{t}.$$

Result on population size

Define *average growth rate*:

$$r_t = r_t(\omega) := \frac{\log |Z_t(\omega)|}{t}.$$

Theorem

On a set of full \mathbf{P} -measure,

$$\lim_{t \rightarrow \infty} (\log t)^{2/d} (r_t - \beta_2) = -c(d, \nu),$$

in P^ω -probability.

Result on population size

Define *average growth rate*:

$$r_t = r_t(\omega) := \frac{\log |Z_t(\omega)|}{t}.$$

Theorem

On a set of full \mathbf{P} -measure,

$$\lim_{t \rightarrow \infty} (\log t)^{2/d} (r_t - \beta_2) = -c(d, \nu),$$

in P^ω -probability.

That is, loosely speaking,

$$r_t \approx \beta_2 - c(d, \nu)(\log t)^{-2/d}.$$

Result on spatial spread

An upper estimate on the spatial spread. The order of the correction term is larger than the $\mathcal{O}(\log t)$ term in a result of Bramson, namely it is

$$\mathcal{O}\left(\frac{t}{(\log t)^{2/d}}\right).$$

Result on spatial spread

An upper estimate on the spatial spread. The order of the correction term is larger than the $\mathcal{O}(\log t)$ term in a result of Bramson, namely it is

$$\mathcal{O}\left(\frac{t}{(\log t)^{2/d}}\right).$$

Theorem

β_1 plays no role and, loosely speaking, at time t the spread of the process

$$\leq t\sqrt{2\beta_2} - c(d, \nu)\sqrt{\frac{\beta_2}{2}} \cdot \frac{t}{(\log t)^{2/d}}.$$

Problem (Shape of branching tree)

How does the discrete probability measure valued process

$$\tilde{Z}_t(\cdot) := \frac{Z_t(\cdot)}{|Z_t|}$$

look like? Is it true that \exists Unique dominant branch?

Problem (Shape of branching tree)

How does the discrete probability measure valued process

$$\tilde{Z}_t(\cdot) := \frac{Z_t(\cdot)}{|Z_t|}$$

look like? Is it true that \exists Unique dominant branch?

Problem (More general branching)

What happens when dyadic branching is replaced by a *general* one? E.g. **critical branching**. Let $\beta_1 = 0$. It is still true (nontrivial) that

$$P^\omega(\text{extinction}) = 1, \text{ a.e. } \omega \in \Omega.$$

Q: What is the order of the tail $P(\tau_{\text{ext}} > t)$?

THANK YOU FOR YOUR ATTENTION!