

Spatial statistics and Bayesian modelling Radu S. Stoica

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Spatial statistics and Bayesian modelling

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Nancy, M2 IMSD - S3

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Table of contents

Course 1. Introduction and some Mathematical background Introduction Some data sets and their related questions Mathematical background

Course 1. Introduction : course organisation, data sets and examples

About me

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Course structure

6 blocks, where 1 block = 1h30 course + 2h30 theoretical and practical exercices

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mark : exam to be defined

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and the list is far from being complete ...

What is this course about ?

Spatial data analysis : investigate and describe data sets whose elements have two components

- position : spatial coordinates of the elements in a certain space
- characteristic : value(s) of the measure(s) done at this specific location
- application domains : image analysis, environmental sciences, astronomy

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"The" questions :

- what is the data subset made of those elements having a certain "common" property ?
- what is the information contained by the data ?

"The" answer :

- generally, these "common" property and information may be described by a statistical analysis
- the spatial coordinates of the spatial data elements add a morphological component to the answer
- the data subset and the information we are looking for, they form a pattern that has relevant geometrical characteristics

Bayesian paradigm :

use prior knowledge in order to answer the question

"The" question re-formulated :

- what is the pattern hidden in the data ?
- what are the geometrical and the statistical characteristics of this pattern ?

Aim of the course : provide you with some mathematical tools to allow you formulate answers to these questions

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Examples : data sets and related questions

For the purpose of this course : software and data sets are available

- ► R library : spatstat by A. Baddeley, R. Turner and contributors → www.spatstat.org
- ► C++ library : MPPLIB by A. G. Steenbeek, M. N. M. van Lieshout, R. S. Stoica and contributors → available at simple demand

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My research partners :

- mathematics : Université de Lorraine, INRIA, Université de Lille, Universitat Jaume I Castellon, CWI Amsterdam
- astronomy and cosmology : Tartu Observatory, Observatorio Astronomico de Valencia, Observatoire de Paris IMCCE
- environmental sciences : RING and GeoRessources, INRA, CIRAD

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industry : SEDIF, St. Gobain

Forestry data (1) : the points positions exhibit attraction \rightarrow clustered distribution



Figure: Redwoodfull data from the spatstat package

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- > library(spatstat)
- > data(redwoodfull) ; plot(redwoodfull)

Forestry data (2) : the points positions exhibit neither attraction nor repulsion \rightarrow completely random distribution



Figure: Japanese data from the spatstat package

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In order to see all the available data sets

> data(package="spatstat")

Biological data (1) : the points positions exhibit repulsion \rightarrow regular distribution



Figure: Cell data from the spatstat package

```
> data(cells)
> cells
planar point pattern: 42 points
window: rectangle = [0, 1] x [0, 1] units
```

Biological data (2) : two types of cells exhibiting attraction and repulsion depending on their relative positions and types



Figure: Amacrine data from the spatstat package

```
> data(amacrine) ;
plot(amacrine,cols=c("blue","red"))
```

Geological data : two types of patterns, line segments and points \rightarrow are these patterns independent ?



Figure: Copper data from the spatstat package

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```
> attach(copper) ; L=rotate(Lines,pi/2) ;
P=rotate(Points,pi/2)
> plot(L,main="Copper",col="blue") ;
points(P$x,P$y,col="red")
```

Animal epidemiology : sub-clinical mastitis for diary herds

- ▶ points \rightarrow farms location
- to each farm \rightarrow disease score (continuous variable)
- clusters pattern detection : regions where there is a lack of hygiene or rigour in farm management



Figure: The spatial distribution of the farms outlines almost the entire French territory (INRA Avignon).

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Cluster pattern : some comments

- particularity of the disease : can spread from animal to animal but not from farm to farm
- cluster pattern : several groups (regions) of points that are close together and have the "same statistical properties"
- ► clusters regions → approximate it using interacting small regions (random disks)
- local properties of the cluster pattern : small regions where locally there are a lot of farms with a high disease score value

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problem : pre-visualisation is difficult ...

Image analysis : road and hydrographic networks



Figure: a) Rural region in Malaysia (http://southport.jpl.nasa.gov), b) Forest galleries (BRGM).

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Thin networks : some comments

- \blacktriangleright road and hydrographic networks \rightarrow approximate it by connected random segments
- topologies : roads are "straight" while rivers are "curved"
- texture : locally homogeneous, different from its right and its left with respect a local orientation
- avoid false alarms : small fields, buildings,etc.
- local properties of the network : connected segments covered by a homogeneous texture

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Cosmology (1): spatial distribution of galactic filaments



Figure: Cuboidal sample from the North Galactic Cap of the 2dF Galaxy Redshift Survey. Diameter of a galaxy \sim 30 \times 3261.6 light years.

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Cosmology (2) : study of mock catalogs



(a)

Figure: Galaxy distribution : a) Homogeneous region from the 2dfN catalog, b) A mock catalogue within the same volume

Cosmology (3) : questions and observations

Few words about the 2dF GRS and SDSS catalogues

- filaments, walls and clusters : different size and contrast
- inhomogeneity effects (only the brightest galaxies are observed)
- filamentary network the most relevant feature
- local properties of the filamentary network : connecting random cylinders containing a "lot" of galaxies "along" its main axis

Mock catalogues

- how "filamentary" they are w.r.t the real observation ?
- how the theoretical models producing the synthetic data fits the reality ?

Cosmology (4) : cluster detection



Figure: Distribution of galaxies in the 2MRS data set. Positions of galaxies are given in supergalactic coordinates, where observer is located at the origin of coordinates (marked as blue point on the figure). The thickness of the slice shown in the figure is 15 Mpc. Some galaxy clusters are marked with black ellipses to highlight the elongation of galaxy groups/clusters along the line of sight.

Cosmology (5) : questions and observations

Few words about the 2MRS data set:

- more galaxies are observed
- price to pay : lack of precision for the third coordinates
- consequence : finger-of-God effect is much more important
- galaxy groups and clusters seem elongated along the line of sight

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inhomogeneity effects

Cosmology (6) : influence of the new observations on the already detected structures



Figure: Photometric galaxies(green dots), spectroscopical galaxies (red dots) and filaments (blue) : a) photometrical galaxies projected on a sphere, b) photometrical galaxies lines of sight

Few words about the SDSS Data Release 12 data set:

- much more galaxies are observed
- price to pay : bigger lack of precision for the third coordinates
- question : how this new data set is related to the already existing detected structures
- need to compare and to relate complex patterns in 2d and 3d

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Cosmology (7) : knowing the filamentary structure what is the galaxy distribution ?



Figure: Data set : galaxy positions with spines (filaments main axes)

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Some questions related to galaxy distribution

- galaxies are spread as pearls on a necklace : (Tempel et al., 2014)
- inhomogeneity : filaments presence or interactions ?
- which of these factors controls the galaxy distribution ?
- what type of interaction : gravitational, territorial, component oriented ?

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- what is the interaction range ?
- what is the size of a cluster ?

Network science



Figure: Collaborations among researchers within the Loria laboratory : HAL data set (2018).

Description of the network :

- each node represents a researcher,
- the edges are collaborative links
- nodes' color represent the affiliation to a laboratory.
- all Loria members are coloured in yellow, while the members of the other labs are differently coloured

Related questions :

- what determines the occurrence of a collaborative link ?
- how the presence of cooperating individuals can be characterised ?
- how to quantify the cooperation behaviour of a research team ?

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Oort cloud comets (1)

The comets dynamics :

- Comets parameters (z, q, cos i, ω, Ω) → inverse of the semi-major axis, perihelion distance, inclination, perihelion argument, longitude of the ascending node
- variations of the orbital parameters
- initial state : parameters before entering the planetary region of our Solar System
- final state : current state
- question : do these perturbation exhibit any spatial pattern ?

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Oort cloud comets (2)

Study of the planetary perturbations

- $ightarrow \sim 10^7$ perturbations were simulated
- data \rightarrow set of triplets $(q, \cos i, \triangle z)$
- ▶ spatial data framework : location $(q, \cos i)$ and marks $\triangle z$
- $\triangle z = z_f z_i$:perturbations of the cometary orbital energy
- local properties of the perturbations : locations are uniformly spread in the observation domain, marks tend to be important whenever they are close to big planets orbits
- reformulated question : do these planetary observations exhibit an observable spatial pattern ?
- problem : pre-visualisation is very difficult ...

Spatio-temporal data

Time dimension available :

- the previous example may be considered snapshots
- more recent data sets have also a temporal coordinate
- question : what is the pattern hidden in the data and its spatio-temporal description ?

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Binary asteroids

Bayesian orbit determination :

- input data : *n* observations at times $t = (t_1, ..., t_n)$
- observational equation :

$$\mathsf{w} = \psi(\mathsf{s}) + \varepsilon$$

with

- $w = (x_1, y_1, \dots, x_n, y_n)$: the rectangular coordinates of the secondary asteroid with primary asteroid in the origin
- s = (a, e, i, Ω, ω, T, P) : the apparent orbit Keplerian elements and the period of revolution P
- \triangleright ε : noise and measure errors
- question : what are the best parameters s fitting the observations w
- problem : the z_1, \ldots, z_n coordinates are not observable
- our solution : elliptical regression + Bayesian modelling
Illustration example :



Figure: Orbit detection result (asteroid 283 Emma) : observed data (black points), our MCMC Bayesian detection method (blue points), N. V. Emeliyanov least square method (red points)

Roads dynamics in Central Africa region :

- in forest region with rare woods, road networks appear and disappear within the territory of an exploitation concession
- ► there is a difference between "classical" road networks and "exploitation" networks → mining galleries
- this roads dynamics may be relevant in many aspects : health of the forest, respect of rules for the enterprises, environmental behaviour and understanding

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- characterize the distribution and the dynamics of the road network
- \rightarrow video roads dynamics

Study the spatio-temporal spread of failures in a water distribution network : failures (black points), detector's activation (red points)

- information available : position, activation date, alert type, etc.
- SEDIF data and questions : do the detectors work ? do the failures form a particular pattern ?
- how to integrate the temporal dynamics ?



Paleontology : Guérande salinas - fairy rings

- growing rings : territories occupied by cyano-bacterias
- the size of a ring is proportional with its age
- what determines the spatial distribution of the rings : edge effects, water arrival, interaction ?
- how to integrate the temporal dyanmics ?



Figure: The distribution of the morphostructures and their sizes (diameter in cm)

Sismology : characterisation of earthquakes occurences

- earthquakes : space-time events
- self-excitation phenomenon
- characterize and predict the sismic activity in different regions



Figure: Representation of positions of earthquakes on the French territory

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Synthesis

Hypothesis : the pattern we are looking for can be approximated by a configuration of random objects that interact

marked points pattern : repulsive or attractive marked points

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- clusters pattern : superposing random disks
- filamentary network : connected and aligned segments

Important remark :

locally : the number of objects is finite

Marked point processes :

- probabilistic models for random points with random characteristics
- ▶ origin → stochastic geometry
- \blacktriangleright the pattern is described by means of a probability density \rightarrow stochastic modelling
- the probability density allows the computation of average quantities and descriptors (these are integrals) related to the pattern
- conversely, whenever a pattern is observed, the probabilistic framework allows the derivation of the law of parameters conditionned on the observation : Bayesian inference
- not the only integrator ...

Remarks :

- there exist also deterministic mathematical tools able to treat pattern recognition problems
- probability is cool : the phenomenon is not controlled, but understood
- probability thinking framework offers simultaneously the analysis and the synthesis abilities of the proposed method

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- probabilistic approach deeply linked with physics :
 - exploratory analysis
 - model formulation
 - simulation
 - statistical inference

- \blacktriangleright comets example : random fields \rightarrow another probabilistic mathematical tool
 - unifying random fields and marked point processes is a mathematical challenge
 - determinental point processes ... ?
- binary system example :
 - not especially a point process
 - the pattern we are looking for is a "tube" of ellipses
 - common points : Gibbsian formulation of the problem to optimize and Bayesian approach

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spatio - temporal examples :

- spatio-temporal marked point processes
- random sets theory
- ▶ general idea : new data sets require new mathematics → stochastic processes and stochastic geometry
- still, partial answers to these questions can be given using the tools presented in this course
- present challenge : big data
 - examples : particle tracking (quantum dots), cosmological data, etc.

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Important question :

- are you interested in an internship : IECL, Inria, GeoRessources, CRAN + travelling ?
- what about a phd ?

Mathematical background

Measure and integration theory \rightarrow blackboard

- $\blacktriangleright \sigma$ -algebra
- measurable space, sets, functions
- measure
- measure space, integral with respect to a measure

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probability space, probability measure

Table of contents

Course 2. Point processes, Binomial and Poisson point processes

- Definition of a point process
- Binomial point process
- Poisson point process
- Few words about self-exciting point processes
- Exercises : binomial and Poisson point processes, simulation

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Course 2. Point processes, Binomial and Poisson point processes

Construction of a point process : mathematical ingredients

• observation window : the measure space (W, B, ν), with W ⊂ ℝ^d, B the Borel σ−algebra and 0 < ν(W) < ∞ the Lebesgue measure

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▶ points configuration space : probability space $(\Omega, \mathcal{F}, \mathbb{P})$

Configuration space construction :

• state space Ω :

 $\begin{array}{ll} W_n & \text{is the set of all n-tuples} \quad \{w_1, \ldots, w_n\} \subset W \\ \Omega = \cup_{n=0}^{\infty} W_n, \quad n \in \mathbb{N} \end{array}$

• events space \mathcal{F} : the $\sigma-$ algebra given by

$$\mathcal{F} = \sigma(\{\mathsf{w} = \{\mathsf{w}_1, \ldots, \mathsf{w}_n\} \in \Omega : n(\mathsf{w}_B) = n(\mathsf{w} \cap B) = m\})$$

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for any bounded $B \in \mathcal{B}$ and $m \in \mathbb{N}$

• probability measure \mathbb{P} : the model answering our questions

Definition

A point process in W is a measurable mapping from a probability space (S, A) in (Ω, F) . Its distribution is given by

$$\mathbb{P}(X \in F) = \mathbb{P}\{\omega \in S : X(\omega) \in F\},\$$

with $F \in \mathcal{F}$. The realization of a point process is random set of points in W. We shall sometimes identify X and $\mathbb{P}(X \in F)$ and call them both a point process.

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Remarks : point process \Rightarrow random configuration of points w in a observation window W. In the following, it is considered that :

- ▶ a points configuration is w = {w₁, w₂,..., w_n}, with n the corresponding number of points
- the process is locally finite : n(w ∩ B) is finite whenever v(B) is finite

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▶ the process is simple : $w_i \neq w_j$ for $i \neq j$

Marked point processes : attach characteristics to the points \rightarrow extra-ingredient : marks probability space (M, M, ν_M)

Definition

A marked point process is a random sequence $x = \{x_n = (w_n, m_n)\}$ such that the points w_n are a point process in W and m_n are the marks corresponding for each w_n .

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Examples :

- random circles : $M = (0, \infty)$
- ▶ random segments : $M = (0, \infty) \times [0, \pi]$
- multi-type process : $M = \{1, 2, \dots, k\}$

... and all the possible combinations ... \rightarrow drawing

Stationarity and isotropy. A point process X on W is stationary if it has the same distribution as the translated proces X_w , that is

$$\{w_1,\ldots,w_n\} \stackrel{\mathcal{L}}{=} \{w_1+w,\ldots,w_n+w\}$$

for any $w \in W$.

A point process X on W is isotropic if it has the same distribution as the rotated proces rX, that is

$$\{w_1,\ldots,w_n\} \stackrel{\mathcal{L}}{=} \{\mathsf{r}w_1,\ldots,\mathsf{r}w_n\}$$

for any rotation matrix r.

- motion invariant : stationary and isotropic
- marked case : in principle easy to generalize, but take care ...
- counter example : a point process on a half plane is not stationary

Intuitive characterisation of a point process : being able to say how many points of the process X can be found in a neighbourhood in W

The mathematical tools for point processes : should be able to do the following

- count the points of a point process in a small neighbourhood of a point in W, and then extend the neighbourhood
- count the points of a point process in a small neighbourhood of a typical point of the process X, and then extend the neighbourhood
- "counting" means using a probability measure based counter

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Let X be a point process on W, and let us consider the counting variable

$$N(B) = n(X_B), \quad B \in \mathcal{B},$$

representing the number of points "falling" in B. Let us consider also the sets of the form

$$F_B = \{ \mathsf{x} \in \Omega : n(\mathsf{x}_B) = 0 \},\$$

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that are called void events.

Theorem

The distribution of a point process X on a complete, separable metric space (W, d) is determined by the finite dimensional distributions of its count function, i.e. the joint distribution of $N(B_1), \ldots, N(B_m)$ for any bounded $B_1, \ldots, B_m \in \mathcal{B}$ and $m \in \mathbb{N}$.

Theorem

The distribution of a simple point process on a complete, separable metric space (W, d) is uniquely determined by its void probabilities

$$v(B) = \mathbb{P}(N(B) = 0), \quad B \in \mathcal{B}.$$

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Choquet's theorem

Theorem

The distribution of a random closed set X is entirely determined by the functional

$$T_X(K) = \mathbb{P}(X \cap K \neq \emptyset)$$

for every compact K in W.

 random closed sets : more general object than a point process
 this object cannot be counted, but it can be observed through a compact window ...

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▶ practical applications \rightarrow Boolean model

Binomial point process

The trivial random pattern : a single random point x uniformly distributed in a compact W such that

$$\mathbb{P}(x \in B) = rac{
u(B)}{
u(W)}$$

for all $B \in \mathcal{F}$.

More interesting point pattern : n independent points distributed uniformly such that

$$\mathbb{P}(x_1 \in B_1, \dots, x_n \in B_n) = \\ = \mathbb{P}(x_1 \in B_1) \cdot \dots \cdot \mathbb{P}(x_n \in B_n) \\ = \frac{\nu(B_1) \cdot \dots \cdot \nu(B_n)}{\nu(W)^n}$$

for Borel subsets B_1, \ldots, B_n of the compact W. \rightarrow drawing Properties

- this process earns its name from a distributional probability
- ▶ the r.v. N(B) with $B \subseteq W$ follows a binomial distribution with parameters

$$n=N(W)=n(x_W)$$

and

$$p=\frac{\nu(B)}{\nu(W)}$$

the intensity of the binomial point process, or the mean number of points per unit volume

$$\rho = \frac{n}{\nu(W)}$$

the mean number of points in the set B

$$\mathbb{E}(N(B)) = np = \rho\nu(B)$$

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- the binomial point process is simple
- number of points in different subsets of W are not independent even if the subsets are disjoint

$$N(B) = m \Rightarrow N(W \setminus B) = n - m$$

the distribution of the point process is characterized by the finite dimensional distributions

$$\mathbb{P}(N(B_1) = n_1, ..., N(B_k) = n_k)$$
 for $k = 1, 2, ...$

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such that $n_1 + n_2 + \ldots + n_k \leq n$

If the B_k are disjoint Borel sets with B₁ ∪ ... B_k = W and n₁ + ... + n_k = n, the finite-dimensional distributions are given by the multinomial probabilities

$$\mathbb{P}(N(B_1) = n_1, \dots, N(B_k) = n_k) \\ = \frac{n!}{n_1! \dots n_k!} \frac{\nu(B_1)^{n_1} \dots \nu(B_k)^{n_k}}{\nu(W)^n}$$

the void probabilities for the binomial point process are given by

$$v(B) = \mathbb{P}(N(B) = 0) = \frac{(\nu(W) - \nu(B))^n}{\nu(W)^n}$$

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Stationary Poisson point process

Motivation : what happens if extend W towards \mathbb{R}^d ?

- convergence binomial towards Poisson
- \blacktriangleright \rightarrow drawing + blackboard

Definition : a stationary (homogeneous) Poisson point process X is characterized by the following fundamental properties

Poisson distribution of points counts. The random number of points of X in a bounded Borel set B has a Poisson distribution with mean ρν(B) for some constant ρ, that is

$$\mathbb{P}(N(B) = m) = \frac{(\rho\nu(B))^m}{m!} \exp(-\rho\nu(B))$$

Independent scattering. The number of points of X in k disjoint Borel sets form k independent random variables, for arbitrary k

Properties

- simplicity : no duplicate points
- the mean number of points in a Borel set B is

$$\mathbb{E}(N(B)) = \rho\nu(B)$$

- ρ : the intensity or density of the Poisson process, and it represents the mean number of points in a set of unit volume
- ▶ $0 < \rho < \infty$, since for $\rho = 0 \Rightarrow$ the process contains no points, while for $\rho = \infty$ we get a pathological case

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 if B₁,..., B_k are disjoint Borel sets, then N(B₁),..., N(B_k) are independent Poisson variable with means ρν(B₁),..., ρν(B_k). Thus

$$\mathbb{P}(N(B_1) = n_1, \dots, N(B_k) = n_k) \\ = \frac{\rho^{n_1 + \dots + n_k} \nu(B_1)^{n_1} \dots \nu(B_k)^{n_k}}{n_1! \dots n_k!} \exp\left(-\sum_{i=1}^k \rho \nu(B_i)\right),$$

- this formula can be used to compute joint probabilities for overlapping sets
- the void probabilities for the Poisson point process are given by

$$v(B) = \mathbb{P}(N(B) = 0) = exp(-\rho(\nu(B)))$$

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- the Poisson point process with ρ = ct. is stationary and isotropic
- if the intensity is a function $\rho: W \to \mathbb{R}^+$ such that

$$\int_B \rho(w) d\nu(w) < \infty$$

for bounded subsets $B \subseteq W$, then we have a inhomogeneous Poisson process with mean

$$\mathbb{E}(N(B)) = \int_{B} \rho(w) d\nu(w) = \Upsilon(B)$$

- \blacktriangleright Υ is called the intensity measure
- we have already seen that for the stationary Poisson process : $\Upsilon(B) = \rho \nu(B)$

Theorem

Conditionning a Poisson point process. Let X be a stationary Poisson point process on \mathbb{R}^d with intensity $\rho > 0$ and W a bounded Borel set with $\nu(W) > 0$. Then, conditional on the event $\{N(W) = n\}$, X restricted to W is a binomial point process of n points.

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- ▶ proof of the theorem : \rightarrow Exercice 1
- **b** bonus : \rightarrow Exercice 2, 3 and 4

Some properties of the Poisson point process

Theorem

Interval theorem. Let X be a stationary point process on $(0, \infty)$ with intensity ρ and let the points of X be written in ascending order :

$$0 < X_1 < X_2 < \ldots < X_n < \ldots$$

The the random variables :

$$Y_1 = X_1, Y_n = X_n - X_{n-1}$$

are independently, identically distributed according to $g(y) = \rho \exp(-\rho y)$ for y > 0.

- ▶ bus paradox : if to the initial process a symmetric independent copy on (-∞, 0) is added, then the interval between two consecutive points of the process containing 0 is longer than any other interval between two consecutive points
- no extension of this result for $d \ge 2$

Maybe most important marked Poisson point process : the unit intensity Poisson point process with i.i.d. marks on a compact W

- number of objects $\sim \text{Poisson}(\nu(W))$
- ► locations and marks i.i.d. : $w_i \sim \frac{1}{\nu(W)}$ and $m_i \sim \nu_M$ The corresponding probability measure : weighted 'counting' of

objects

$$\mathbb{P}(X \in F) = \sum_{n=0}^{\infty} \frac{e^{-\nu(W)}}{n!} \int_{W \times M} \cdots \int_{W \times M} \mathbb{1}_{F}\{(w_1, m_1), \dots, (w_n, m_n)\}$$
$$\times d\nu(w_1) d\nu_M(m_1) \dots d\nu(w_n) d\nu_M(m_1)$$

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for all $F \in \mathcal{F}$.

Remark : the simulation of this process is straightforward, while the knowledge of the probability distribution allows analytical computations of the interest quantities

Simulations results of some Poissonian point processes : the domain is $W = [0, 1] \times [0, 1]$ and the intensity parameter is $\rho = 100$



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Figure: Poisson based models realizations : a) unmarked, b) multi-type and c) Poisson process of segments.

Definition

A disjoint union $\bigcup_{i=1}^{\infty} X_i$ of point processes X_1, X_2, \ldots is called superposition.

Proposition

If $X_i \sim \text{Poissson}(W, \rho_i)$, i = 1, 2, ... are mutually independent and if $\rho = \sum \rho_i$ is locally integrable, then with probability one, $X = \bigcup_{i=1}^{\infty} X_i$ is a disjoint union and est $X \sim \text{Poisson}(W, \rho)$.

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 \rightarrow stable character of the Poisson process

Definition

Let be $q: W \to [0,1]$ a function and X a point process on W. The point process $X_{thin} \subset X$ obtained by including the $\xi \in X$ in X_{thin} with probability $q(\xi)$, where points are included/excluded independently of each other, is said to be an independent thinning of X with retention probabilities $q(\xi)$.

Formally, we can set

$$X_{\mathsf{thin}} = \{\xi \in X : R(\xi) \le q(\xi)\},\$$

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with the random variables $R(\xi) \sim \mathcal{U}[0,1]$, $\xi \in W$, mutually independent and independent of X.
Proposition

Suppose that $X \sim Poisson(W, \rho)$ is subject to independent thinning with retention probabilities $q(\xi)$, $\xi \in W$ and let

$$\rho_{thin} = q(\xi)\rho(\xi), \quad \xi \in W.$$

Then X_{thin} and $X \setminus X_{thin}$ are independent Poisson processes with intensity functions ρ_{thin} and $\rho - \rho_{thin}$, respectively.

Corollary

Suppose that $X \sim Poisson(W, \rho)$ with ρ bounded by a positive constant C. Then X is distributed as independent thinning of a Poisson(W, C) with retention probabilities $q(\xi) = \rho(\xi)/C$.

Some general facts concerning the Poisson point processes

- the Poisson point process is as important for spatial statistics as the Gaussian process in classical probability theory
- \blacktriangleright the law is completely known \rightarrow analytical formulas
- the Poisson process is invariant under independent thinning
- easy procedure for simulate non-stationary Poisson process
- completely random patterns : null or the default hypothesis that we want to reject
- independence \rightarrow no interaction \rightarrow no structure in the data

- two stationary Poisson point processes, they are not absolutely continuous with respect to each other, except if one process has unit intensity or if they have the same intensity
- two inhomogeneous Poisson point processes with strictly positive intensities, they are absolutely continuous with respect to each other
- ► more complicate models can be built → specifying a probability density p(x) w.r.t. the reference measure given by the unit intensity Poisson point process. This probability measure is written as

$$\mathbb{P}(X \in F) = \int_F p(\mathsf{x})\mu(d\mathsf{x})$$

with μ the reference measure.

Remark : in this case the normalizing constant is not available from an analytical point of view. To check this replace in the expression of $\mu(\cdot)$ the indicator function $1_F\{y\}$ with $p(y) \dots$ \rightarrow Exercice 5

Few words about self-exciting point processes

Hawkes processes : a point process defined by its intensity of events conditional on the past $\lambda^*(t) \rightarrow$ the intensity of the process evolves with the time depending on the points arrived in the configuration : no more indepence

$$\lambda^*(t) = \lambda + \sum_{i=1}^n \mu(t-t_i)$$

such as

- (t_i)_{1≤i≤n} the sequence of arrival times of events that have occurred up to t.
- > λ background intensity.
- $\mu : [0, +\infty[\rightarrow [0, +\infty[$ excitation function.

Remarks :

- ▶ if $\mu = 0 \Rightarrow$ classical Poisson process
- modelling : several models available for the excitation functions
- simulation : thinning method
- inference : the conditional intensity allows the consutruction of a likelihood function

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 application : sismology and epidemiology (extend the definition of the conditional intensity) Exponential model : an example of excitation function for Hawkes processes

$$\mu(t) = \alpha \exp(-\beta t)$$

with $\alpha < \beta$. The parameter α gives the instantaneous influence of events and β the rate at which it decreases.



Figure: Number of events and conditional intensity of a Hawkes process, with exponential excitation function with parameters $\alpha = 0.6, \beta = 0.8$ et $\lambda = 1.2$.

Exercises¹ : binomial and Poisson point processes, simulation

Exercise 1. Prove the Theorem related to the conditionning of a stationary Poisson point process.

Hint : Compute the void probabilities in subsets $B \subset W$.

Exercise 2. The spherical contact distribution for a point process is given by

$$F(r) = \mathbb{P}(d(w, X) \leq r)$$

where d(w, X) is the minimum distance from a given point $w \in W$ to the point process X. Compute the expression of F(r) for a stationary Poisson point process with intensity $\rho > 0$.

¹Part of the theoretical and practical exercises is due to the generous help of Zbyněk Pawlas from Charles University in Prague and Marie-Colette van Lieshout from CWI Amsterdam.

Exercise 3. Let X be a stationary Poisson point process in \mathbb{R}^2 . Denote by D_X the distance from the origin to the nearest point in X. Calculate the mean and the variance of the random variable D_X .

Exercise 4. Let $X_1, X_2, ...$ be independent and exponentially distributed with parameter ρ and define a point process on \mathbb{R}^+ by

$$X = \{X_1, X_1 + X_2, X_1 + X_2 + X_3, \ldots\}$$

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Calculate $\mathbb{P}(N((0, t]) = 0 \text{ for } t \in \mathbb{R}^+$.

Exercise 5. Install the R package **spatstat**. Download the following documents :

- Package spatstat
- Analysing spatial point patterns in 'R' by A. J. Baddeley
- a) Simulate and print 10 realizations of a Binomial point process with n = 10 points in the square $W = [0, 1]^2$.
- b) Simulate and print 10 realization of a homogeneous Poisson point processes with intensity parameter $\rho = 10$ in the square $W = [0, 1]^2$. Compare the realizations of the previous two processes. What do you observe ?

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- c) Simulate an inhomogeneous Poisson point process given by the intensity
 - $\rho(x_1, x_2) = 100(x^2 + y)$ in the domain W = [0, 2]x[0, 1].
 - $\rho(x_1, x_2) = 100 \exp[-(x^2 + y^2)]$ in the domain given by the ellipse $\frac{x^2}{4} + y^2 1 = 0$
- d) Simulate and print a realization of a multi-type marked Point process with the following parameters : the locations process is a stationary Poisson point process in [0, 1]², while the marks probability law is the uniform distribution over three point types.
- e) Simulate and print a realization of a Poisson process of random discs. The centres locations process is the previous Poisson process, while the disk radius follows an uniform distribution on the interval [0, 0.05].

- f) Simulate and print a realization of a Poisson process of random segments. The centres locations process is the previous Poisson process, while the orientation and lengths parameters are independently uniformly distributed in the intervals $[0, \pi]$ and [0, 0.2].
- g) Propose statistics that may characterize the previous processes ? Try to approximate them by simulation. How can this be used as a statistical test ?
- h) Test if the following spatstat datasets may be considered as a stationary Poisson point process : cells, swedishpines, japanesepines, redwood ?

Hint : Use the help. Some of the commands you may be interested in are ppp, psp, rpoispp, rmpoispp, owin, runifpoint.

Table of contents

Course 3. Moment and factorial moment measures

Moment and factorial moment measures, product densities Moment measures for the Poisson process Campbell measures Campbell - Mecke formula Exercises : moment and factorial moment measures

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Cours 3. Moment and factorial moment measures, product densities

Present context :

- mathematical background
- definition of a marked point process
- Binomial and Poisson point process
- important result : the point process law is determined by counts of points

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Let X be a point process on W. The counts of points in bounded Borel regions of $B \subset W$, N(B) characterize the point process and they are well defined random variables

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- it is difficult to average the pattern X
- it is possible to compute moments of the N(B)'s

The appropriate mathematical tools are :

- the moment measures
- the factorial moment measures
- the product densities

\rightarrow blackboard

Moment measures of the Poisson process

- ► stationary Poisson point processes → Exercise 6
- the n-th product density measure of an independently thinned point process is

$$\rho_{\text{thin}}^{(n)}(w_1,\ldots,w_n) = \rho^{(n)}(w_1,\ldots,w_n) \prod_{i=1}^n q(w_i)$$

this gives the invariance under independent thinning of the n-th point correlation function (van Lieshout, 2011)

$$\frac{\rho_{\text{thin}}^{(n)}(w_1,\ldots,w_n)}{\rho_{\text{thin}}(w_1)\cdot\ldots\cdot\rho_{\text{thin}}(w_n)}=\frac{\rho^{(n)}(w_1,\ldots,w_n)}{\rho(w_1)\cdot\ldots\cdot\rho(w_n)}$$

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 \blacktriangleright \rightarrow Exercise 9 : explain the envelope tests

Campbell measures

Present context :

- ► counting points *i.e.* computing moment and factorial moment measures → very interesting tool for analysing point patterns : allow the computation of average quantities
- \blacktriangleright still, compute an average pattern \rightarrow difficult and challenging problem
- \blacktriangleright idea : counting points that have some specific properties \rightarrow Campbell measures

Definition

Let X be a point process on W. The Campbell measure is

 $C(B \times F) = \mathbb{E}\left[N(B)1\{X \in F\}\right],$

for any bounded $B \in \mathcal{B}$ and $F \in \mathcal{F}$.

The first order moment measure can be expressed as a Campbell measure :

$$C(B \times \Omega) = \mathbb{E}[N(B)] = \mu^{(1)}(B).$$

► the moment and Campbell measures are not necessarily finite : their respective extension to an unique σ-finite measure can be shown (see van Lieshout, 2000) → Exercise 7 Higher order Campbell measures are constructed in a similar manner. For instance, the second ordre Campbell measure is

$$C^{(2)}(B_1 \times B_2 \times F) = \mathbb{E}\left[N(B_1)N(B_2)1\{X \in F\}\right],$$

from which we can get the second order moment measure

$$C^{(2)}(B_1 imes B_2 imes \Omega) = \mathbb{E}\left[N(B_1)N(B_2)\right] = \mu^{(2)}(B_1 imes B_2)$$

Remark :

- the moment measures allow to average functions h(x) measured in the location of a point process X : the function h does not depend on X
- the Campbell measures allow to average functions h(x,x) measured in the location of a point process X : the function h may depend on X

Campbell - Mecke formula

Theorem

Let $h: W \times \Omega \rightarrow [0, \infty)$ a measurable function that is either non-negative either integrable with respect to the Campbell measure. Then

$$\mathbb{E}\left[\sum_{w\in X}h(w,X)\right] = \int_{W}\int_{\Omega}h(w,x)dC(w,x).$$

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Proof. \rightarrow Exercise 8

A more general Campbell-Mecke formulas

Theorem

For a point process X and arbitrary nonnegative measurable function h that does not depend on X we have

$$\mathbb{E}\sum_{w_1,\ldots,w_n\in X}h(w_1,\ldots,w_n)=\int_W\cdots\int_Wh(w_1,\ldots,w_n)d\mu^{(n)}(w_1,\ldots,w_n)$$

and

$$\mathbb{E}\sum_{w_1,\ldots,w_n\in X}^{\neq}h(w_1,\ldots,w_n)=\int_{W}\cdots\int_{W}h(w_1,\ldots,w_n)d\alpha^{(n)}(w_1,\ldots,w_n)$$

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Proof.

Follow the same proof scheme as previously.

Remarks :

If the function h does not depend on the point process X, the Campbell - Mecke becomes

$$\mathbb{E}\left[\sum_{w\in X}h(w)\right]=\int_Wh(w)d\mu^{(1)}(w).$$

• point process of intensity function $\rho(w)$

$$\mathbb{E}\left[\sum_{w\in X}h(w)\right]=\int_Wh(w)\rho(w)d\nu(w).$$

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the preceding formula becomes for a stationary Poisson point process of intensity ρ > 0

$$\mathbb{E}\left[\sum_{w\in X}h(w)\right]=\rho\int_Wh(w)d\nu(w),$$

this new formula is true for any stationary point process but is difficult to relate its intensity with its distribution ...

• point process of second order intensity function $\rho^{(2)}(u, v)$

$$\mathbb{E}\left[\sum_{u,v\in X}^{\neq} h(u,v)\right] = \int_{W} \int_{W} h(u,v) \rho^{(2)}(u,v) d\nu(u) d\nu(v).$$

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Exercises : moment and factorial moment measures

Exercise 6. Let X be a stationary Poisson point process in \mathbb{R}^2 with intensity parameter ρ . Compute the first and second order moment and factorial measures. Compute the corresponding product densities and the pair correlation function.

Exercise 7. Let X be a finite point process on a compact subset $W \subset \mathbb{R}^d$ with the number of points given by $p_n = \frac{1}{n(n-1)}$ for $n \ge 2$ and zero otherwise

- a) Show that p_n is a probability function.
- b) Show that the first order moment measure on W is not finite.

Exercise 8. Prove the Campbell-Mecke Theorem. *Hint :* start by considering indicator functions $h(w,x) = 1\{(w,x) \in A \times F\}$ for some bounded Borel set $A \in B$ and some $F \in \mathcal{F}$. Exercise 9.

- a) Simulate and print a realization of a Poisson point processes with intensity parameter $\rho = 100$ on the square $W = [0, 1] \times [0, 1]$. Use the **spatstat** package to compute and print estimates of the empty space function and of the pair correlation function for a pattern of points obtained by the simulation of the previous process. Compare the obtained values with their corresponding theoretical values.
- b) Build envelope tests based on these characteristics to compare the observed pattern with the realization of a Poisson process.
- c) Analyse the data sets : redwoodfull, japanesepines and cells. How, the empty space function and the pair correlation function can be used to diagnose clustering, repulsion or completely randomness of a pattern ? Try to propose a model for one of these data sets and test it.

Hint : Use the help. Commands you may be interested in are pcf, Fest, envelope, density.

Exercise 10. χ^2 *test of homogeneity.* Let x be an observed point pattern of a finite point process in a compact $W \subset \mathbb{R}^2$. Let us consider the hypothesis test given by :

- H_O : the point process is stationary Poisson,
- H_1 : the point process is not stationary Poisson.

Divide the window W into quadrats B_1, \ldots, B_m and count the numbers of points n_1, \ldots, n_m in each quadrat. Under the null hypothesis, the n_j s are realisations of independent Poisson random variables with expected values $\mu_j = \rho a_j$ where ρ is the unknown intensity and $a_j = \nu(B_j)$.

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Given the total number of points $n = \sum_j n_j$, and the total window area $a = \sum_j a_j$, the estimated intensity is $\hat{\rho} = n/a$, and the expected count in quadrat B_j is $e_j = \hat{\rho}a_j = na_j/a$. The test statistic is

$$T = \sum_{j=1}^m \frac{(n_j - e_j)^2}{e_j}$$

and under H_0 it follows a χ^2 distribution with m-1 degrees of freedom.

- a) Simulate and print a realization of a Poisson point processes with intensity parameter $\rho = 200$ on the square $W = [0, 1] \times [0, 1]$.
- b) For a realisation of such a process visualise the decomposition in equal surface quadrats. What do you observe if you increase the number of quadrats ?
- c) For the same realisation visualise the decomposition in quadrats given by a Vornoi tesselation. What decomposition may you prefer and why ?

- d) Implement the χ^2 test of homogeneity based on equal surface quadrats.
- e) Implement the test using a Voronoi decomposition. How do you explain the obtained result ?
- f) Repeat the previous points for the realisation of an inhomogeneous Poisson point proces given by $\rho(x, y) = 100(x^2 + y)$ in the domain W = [0, 2]x[0, 1].
- g) Analyse the data sets : redwoodfull, japanesepines and cells. Based on the χ^2 test, check if these patterns tend to be rather *regular* or *clustered*.
- h) What are the advantages and the drawbacks of using this test ?

Hint : Use the help. Commands you may be interested in are quadratcount, intensity, dirichlet, quadrat.test.

Exercise 11.

- a) Simulate and print a realisation of an inhomogeneous Poisson point processes with intensity function given by $\rho(x, y) = (x^2 + y^2)$ in the square $W = [0, 5]^2$.
- b) Intensity estimator proposed by Diggle is

$$\hat{\rho}(w) = \sum_{i=1}^{n} \frac{1}{e(w)} \kappa(w - x_i)$$

with the probability density kernel $\kappa(w) \ge 0$ such that $\int_{\mathbb{R}^2} \kappa(w) d\nu(w) = 1$ and the correction for bias due to edge effects

$$e(w) = \int_W \kappa(w-u) d\nu(u)$$

Outside the window W, the estimated intensity is zero. Use the **R** function density to estimate the intensity function for the previous realisation.

- c) Plot several density estimates using different kernel's bandwidths. What do you notice? Use the R command bw.diggle in order to find an optimal estimate of the kernel's bandwidth.
- d) Estimate the intensity of some real data sets: redwoodfull, cells, japanesepines. Are these patterns stationary ? Motivate your answer.

e) The **R** spatstat data set bei contains a pattern of points and covariates. Use the help to learn about the data structure. Type the following code to visualise the data set: x11() map.data=persp(bei.extra\$elev, theta=-45, phi=18, expand=7, border=NA, apron=TRUE, shade=0.3, box=FALSE, visible=TRUE, colmap=terrain.colors(128), main="Tropical rain forest - data") perspPoints(bei, Z=bei.extra\$elev, M=map.data, pch=16, col="blue".cex=0.5)

- f) Estimate the intensity of the bei data set point pattern ? Is the pattern stationary ? Motivate your answer.
- g) Is the intensity depending of one the available covariates ? A possible manner is to use the χ^2 test based on quadrats. First of all use the function quanttess to make a tesselation of the domain depending on the quantiles of the elevation covariate. Next, use the quadrat.test to reject a Poissonian assumption. What model you test in this way ? What are the alternatives you have considered ?

- h) Another possible strategy to analyse the dependence of the intensity on a covariate is to compare the observed distribution of the values of the covariate at the data points with the values of that covariate at all spatial locations in the observation window. Do you have an explanation for it ?
- i) Use the function cdf.test.ppp to test whether the intensity of the point pattern in bei data set depends on the elevation covariate. Plot the different results.

Table of contents

Course 4. Palm theory

Interior and exterior conditionning A review of the Palm theory Slivnyak - Mecke theorem Applications : summary statistics

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Cours 4. Palm theory

Interior and exterior conditioning : present context

- counting points measures : count points in a small region in $W \rightarrow$ integrate using the Campbell Mecke formula
- idea : count points in a small region in W that is "centred" in a point of the process X → interior conditionning
- ► question : how the measures applied to a process X change, if we add or if we delete a point from the current configuration → exterior conditionning

Palm distributions

- ▶ present construction → blackboard
- the Palm distributions of X at $w \in W$ can be interpreted as

$$P_w(F) = \mathbb{P}(X \in F | N(\{w\}) > 0)$$

the Campbell - Mecke formula can be expressed as

$$\mathbb{E}\left[\sum_{w\in X}h(w,X)\right] = \int_{W}\int_{\Omega}h(w,x)dP_{w}(x)d\mu^{(1)}(w)$$

for stationary point processes

$$\mathbb{E}\left[\sum_{w \in X} h(w, X)\right]$$

= $\rho \int_{W} \int_{\Omega} h(w, x) dP_w(x) d\nu(w)$
= $\rho \int_{W} \int_{\Omega} h(w, x + w) dP_o(x) d\nu(w)$
Slivnyak - Mecke theorem

Theorem

If $X \sim \text{Poisson}(W, \rho)$, then for functions $h: W \times \Omega \rightarrow [0, \infty)$, we have

$$\mathbb{E}\sum_{w\in X}h(w,X\setminus\{w\})=\int_W\mathbb{E}h(w,X)
ho(w)d
u(w),$$

(where the left hand side is finite if and only if the right hand side is finite).

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▶ proof : \rightarrow Exercise 12

General Slivnyak - Mecke theorem

Theorem

If $X \sim Poisson(W, \rho)$, then for any $n \in \mathbb{N}$ and any functions $h: W^n \times \Omega \rightarrow [0, \infty)$, we have

$$\mathbb{E}\sum_{w_1,\ldots,w_n\in X}^{\neq} h(w_1,\ldots,w_n,X\setminus\{w_1,\ldots,w_n\})$$

= $\int_W \cdots \int_W \mathbb{E}h(w_1,\ldots,w_n,X)\prod_{i=1}^n \rho(w_i)d\nu(w_i)$

where the \neq over the summation sign means that the n points w_1, \ldots, w_n are pairwise distinct.

▶ proof : similar to the previous one + induction. For $n \ge 2$ consider

$$\tilde{h}(w, \mathsf{x}) = \sum_{w_2, \dots, w_n \in \mathsf{x}}^{\neq} h(w, w_2, \dots, w_n, \mathsf{x} \setminus \{w_2, \dots, w_n\})$$

- this theorem is a very strong result, since it allows computing averages of a Poisson point process knowing that one or several points belong to the process ...
- application in telecomunications : knowing, in this location I have a mobile phone antena, how the quality of the signal change if I add randomly more antenas ? (the group of F. Baccelli)

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 combining the Campbell-Mecke and the Slivnyak-Mecke theorem, we obtain for a Poisson proces

$$\int_{\Omega} h(\mathsf{x}) dP_w(\mathsf{x}) = \int_{\Omega} h(\mathsf{x} \cup \{w\}) dP(\mathsf{x})$$

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- in words : the Palm distribution of a Poisson process with respect to w is simply the Poisson distribution plus an added point at w
- a more mathematical formulation : the Palm distribution P[↑]_w(·) of a Poisson process of intensity measure ↑ and distribution P[↑] is the convolution P[↑] ★ δ_w of P[↑] with an additional deterministic point at w
- explanation : blackboard
- \rightarrow Exercise 13 and 14

Assumption : X is a stationary point process

the nearest neighbour distance distribution function

$$G(r) = P_w(d(w, X \setminus \{w\}) \le r) \tag{1}$$

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with P_w the Palm distribution. The translation invariance of the distribution of $X \rightarrow$ inherited by the Palm distribution \rightarrow G(r) is well-defined and does not depend on the choice of w.

▶ replacing the Palm distribution in (1) by the distribution of X → the spherical contact distribution or the empty space function

$$F(r) = \mathbb{P}(d(w, X) \leq r)$$

with \mathbb{P} the distribution of *X*.

the J function : compares nearest neighbour to empty distances

$$J(r) = \frac{1-G(r)}{1-F(r)}$$

defined for all r > 0 such that F(r) < 1

The J function describes the morphology of a point pattern with respect to a Poisson process :

$$J(r)$$
 is $\begin{cases} = 1 \quad \text{Poisson : complete random} \\ \leq 1 \quad \text{clustering : attraction} \\ \geq 1 \quad \text{regular : repulsion} \end{cases}$

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For the stationary Poisson process of intensity parameter ρ , on $W \subset \mathbb{R}^2$, these statistics have exact formulas :

$$F(r) = 1 - \exp[-\rho \pi r^2]$$

$$G(r) = F(r)$$

$$J(r) = 1$$

 \rightarrow proof the formulas at the blackboard + Exercice 17

Table of contents

Cours 5. Palm theory (continued). Papangelou conditional intensity. Reduced Palm distributions Campbell and Slivnyak theorems Summary statistics : the *K* and *L* functions Few words about summary statistics estimation Applications : summary statistics Exterior conditioning : conditional intensity Exercises : Palm theory - interior and exterior conditionning

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Cours 5. Palm theory (continued). Papangelou conditional intensity.

Reduced Palm distributions : present context

- ▶ Palm distributions : count the points in a neighbourhood centred on a point of the process → this point is counted as well
- ► idea : in some applications (telecommunications) we may wish to measure the effect of a point process in a location being a point of the process, while this particular point has no effect on the entire process → reduced Palm distributions
- the following mathematical development is rather easy to follows since it is similar to what we have already seen during until now

Reduced Campbell measure

Definition

Let X be a simple point process on the complete, separable metric space (W, d). The reduced Campbell measure is

$$C^{!}(B \times F) = \mathbb{E}\left[\sum_{w \in X \cap B} 1\{X \setminus \{w\} \in F\}\right],$$

for any bounded $B \in \mathcal{B}$ and $F \in \mathcal{F}$.

the analogue of Campbell-Mecke formula reads

$$\mathbb{E}\left[\sum_{w\in X}h(w,X\setminus\{w\})\right]=\int_W\int_\Omega h(w,\mathsf{x})dC^!(w,\mathsf{x}).$$

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assuming the first order moment measure μ⁽¹⁾ of X exists and it is σ-finite, we can apply Radon-Nikodym theory to write

$$C^!(B \times F) = \int_B P^!_w(F) d\mu^{(1)}(w),$$

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for any bounded $B \in \mathcal{B}$ and $F \in \mathcal{F}$

• the function $P^!(F)$ is defined uniquely up to an $\mu^{(1)}$ -null set

It is possible to find a version such that for fixed w ∈ W, P[!]_w(·) is a probability distribution → the reduced Palm distribution Campbell and Slivnyak theorems

the reduced Palm distribution can be interpreted as the conditional distribution

$$P_w^!(F) = \mathbb{P}(X \setminus \{w\} \in F | N(\{w\}) > 0)$$

the Campbell-Mecke formula equivalent

$$\mathbb{E}\left[\sum_{w\in X}h(w,X\setminus\{w\})\right]=\int_W\int_\Omega h(w,\mathsf{x})dP^!_w(\mathsf{x})d\mu^{(1)}(w).$$

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for stationary point processes

$$\mathbb{E}\left[\sum_{w\in X} h(w, X \setminus \{w\})\right]$$

= $\rho \int_{W} \int_{\Omega} h(w, x) dP_{w}^{!}(x) d\nu(w)$
= $\rho \int_{W} \int_{\Omega} h(w, x + w) dP_{o}^{!}(x) d\nu(w)$

► the Slivnyak-Mecke theorem : for a Poisson process on W with distribution P, we have

$$P_w^!(\cdot) = \mathbb{P}(\cdot)$$

► there is a general result linking the reduced Palm distribution and the distribution of a Gibbs process → a little bit later in this course ...

Example

The nearest neighbour distribution G(r) of stationary process can be expressed in terms of the Palm distributions

$$G(r) = 1 - P_o(X \in \Omega : N(b(o, r)) = 1),$$

and the reduced Palm distributions

$$G(r) = 1 - P_o^!(X \in \Omega : N(b(o, r)) = 0),$$

where N(b(o, r)) is the number of points inside the ball centred at the origin o of radius r.

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Summary statistics : the K and L functions

The K function : theoretical explanations \rightarrow blackboard

- maybe one of the most used summary statistic
- for a stationary process, its definition depending on the reduced Palm distribution is

$$\rho K(r) = \mathbb{E}_o^! \left[N(b(o, r)) \right]$$

the L function is

$$L(r) = \left[\frac{K(r)}{\omega_d}\right]^{1/c}$$

with $\omega_d = \nu(b(0,1))$ the volume of the *d*-dimensional unit ball

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for stationary point processes, the pair correlation function is

$$g(r) = \frac{K'(r)}{\sigma_d r^{d-1}}$$

with σ_d the surface area of the unit sphere in ℝ^d
for the stationary Poisson process we have

$$K(r) = \omega_d r^d, \quad g(r) = 1$$

and

$$L(r) = r$$

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 \blacktriangleright \rightarrow Exercise 15 and 16

Few words about summary statistics estimation

'Robbins' theorem

- spatial sampling bias
- unbiased sampling rules
- Horvitz-Thompson and spatial Horvitz-Thompson estimators

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- sampling bias for point processes
- application : Fritz Kleinschroth's work on road network dynamics and Maarja's phd results ...

 $\rightarrow \mathsf{blackboard}$

Applications : summary statistics

Road network dynamics :

► data: spatio-temporal evolution of a road network due to logging activity → you have already seen the video ...

questions:

- do the exploitation companies respect the regulations for preserving the forest ?
- does the economical activity affect the resilience capacity of the forest ?
- ► exploratory analysis: empty space function → road-less space measure
- challenge: build a stochastic model
- people: F. Kleinschroth, J. R. Healey, S. Gourlet-Fleury, F. Mortier, M. N. M. van Lieshout

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paper: (Kleinschroth et al., 2017)



Figure: Toy model for explaining the behaviour of the empty space function: the simulated roads have the length, so the same density of roads per unit of surface.



Figure: Empty space function for characterization the road network evolution in two logging companies.

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Cosmology : influence of the new observations on the already detected structure

- data: SDSS Data Release 12 photometrical galaxies
- question:
 - how this new data set is related to the already detected structure ?
- exploratory analysis: adapt the F, G and J function to establish possible dependence of different types of patterns
- challenge: the position of the photometrical galaxies is not entirely known

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- people: M. Kruuse, E. Tempel, R. Kipper
- paper: in preparation

Preliminary result :



Figure: Estimation of the bivariate J function. The considered sets were the photometric galaxies and the projection on the sphere of the filamentary spines that are rather perpendicular on the line of sight. This result indicates positive association of these two patterns.

Exterior conditioning : conditional intensity

► assume that for any fixed bounded Borel set B ∈ B, the reduced Campbell measure C[!](B × ·) is absolutely continuous with respect to the distribution P(·) of X

then

$$C^{!}(B \times F) = \int_{F} \Lambda(B; \mathsf{x}) d\mathbb{P}(\mathsf{x})$$

for some measurable function $\Lambda(B; \cdot)$ specified uniquely up to a \mathbb{P} -null set

▶ moreover, one can find a version such that for fixed x, $\Lambda(\cdot; x)$ is a locally finite Borel measure \rightarrow the first order Papangelou kernel

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If Λ(·; x) admits a density λ(·; x) with respect to the Lebesgue measure ν(·) on W, the Campbell-Mecke theorem becomes

$$\mathbb{E}\left[\sum_{w\in X} h(w, X \setminus \{w\})\right]$$

= $\int_{W} \int_{\Omega} h(w, x) dC^{!}(w, x)$
= $\mathbb{E}\left[\int_{W} h(w, X)\lambda(w; X) d\nu(w)\right]$

- the function λ(·; ·) is called the Papangelou conditional intensity
- the previous result is known as the Georgii-Nguyen-Zessin formula

the case where the distribution of X is dominated by a Poisson process is especially important

Theorem

Let X be a finite point process specified by a density p(x) with respect to a Poisson process with non-atomic finite intensity measure ν . Then X has Papangelou conditional intensity

$$\lambda(u; \mathsf{x}) = \frac{p(\mathsf{x} \cup \{u\})}{p(\mathsf{x})}$$

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for $u \notin x \in \Omega$.

Proof.

 \rightarrow Exercise 18

Importance of the conditional intensity :

intuitive interpretation :

$$\lambda(u;\mathsf{x})d\nu(u) = \mathbb{P}(\mathsf{N}(du) = 1|\mathsf{X} \cap (d\nu(u))^c = \mathsf{x} \cap (d\nu(u))^c),$$

the infinitesimal probability of finding a point in a region $d\nu(u)$ around $u \in W$ given that the point process agrees with the configuration x outside of $d\nu(u)$

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the "conditional reverse" of the Palm distributions

- \blacktriangleright describe the local interactions in a point pattern \rightarrow Markov point processes
- ► if

$$\lambda(u; \mathsf{x}) = \lambda(u; \emptyset)$$

for all patterns x satisfying $x \cap b(u, r) = \emptyset \rightarrow$ the process has 'interactions of range r at u'

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in other words, points further than r away from u do not contribute to the conditional intensity at u

- integrability of the model
- convergence of the Monte Carlo dynamics able to simulate the model
- \blacktriangleright differential characterization of Gibbs point processes \rightarrow blackboard
 - the Slivnyak-Mecke theorem links Palm distributions with the distribution of a Poisson point process
 - this characterization links the Palm distributions with the distributions of a finite point process
 - this characterization is an essential element for extending finite point process to R^d

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Exercises : Palm theory - interior and exterior conditionning

Exercise 12. Prove the Slivnyak-Mecke Theorem. *Hint :* start by considering the distribution function of Poisson point process in W and compute the desired expectation.

Exercise 13. Compute the average number of pairs of points in a stationary Poisson process of intensity ρ on the planar unit square separated by a distance that does not exceed some fixed $r < \sqrt{2}$.

- a) Do this computation conditionning on the event N(W) = n.
- b) Do this computation using the Campbell Mecke formula.

Hint : $\mathbb{E}[\mathbb{E}[X|Y]] = \mathbb{E}[X]$.

Exercise 14. Let U_1 and U_2 be two independent random variables with uniform distribution on the interval [0, r], r > 0. Define a point process X in \mathbb{R}^2 as

$$X = \bigcup_{m,n\in\mathbb{Z}} (U_1 + mr, U_2 + nr), \quad m,n\in\mathbb{Z}$$

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where $\mathbb{Z} = \{\dots, -1, 0, 1, \dots\}$. Determine the intensity measure and the Palm distributions of X.

Exercise 15. Let X be a stationary Poisson point process on \mathbb{R}^d with intensity parameter ρ . Prove that :

$$K(r) = \omega_d r^d$$
 and $L(r) = r$,

where ω_d is the volume of the unit sphere in \mathbb{R}^d .

Exercise 16. This exercice studies alternative definitions for the Palm distribution and the G-function. Let X be a stationary point process in \mathbb{R}^d with intensity ρ .

a) Show that

$$P_{\nu}(F) = \frac{1}{\rho\nu(A)} \mathbb{E} \sum_{u \in X} \mathbb{1}\{u \in A, X + \nu - u \in F\}, \quad \nu \in \mathbb{R}^{d}, F \in \mathcal{F}$$

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for an arbitrary set $A \subset \mathbb{R}^d$ with $0 < \nu(A) < \infty$.

b) Show that

$$G(r) = rac{1}{
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u(A)} \mathbb{E} \sum_{u \in X} \mathbb{1} \{ u \in A, (X \setminus \{u\}) \cap b(u, r) \neq \emptyset \}, \quad r > 0,$$

for an arbitrary set $A \subset \mathbb{R}^d$ with $0 < \nu(A) < \infty$.

Hint : use the Campbell-Mecke theorem

Exercise 17.

- a) Simulate and print a realization of a Poisson point processes with intensity parameter $\rho = 100$ on the square $W = [0, 1] \times [0, 1]$. Use the **spatstat** package to compute and print estimates of the *G*, *K* and of the *J* function for a pattern of points obtained by the simulation of the previous process. Compare the obtained values with their corresponding theoretical values.
- b) Build envelope tests based on these characteristics to compare the observed pattern with the realization of a Poisson process.

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- c) Analyse the data sets : redwoodfull, japanesepines and swedishpines.
- d) Analyse the behaviour of the different tree species in the data set : lansing.

Hint : help(lansing). For the last two points use all the sumary statistics you know.

Exercise 18. Prove the Papangelou conditional intensity Theorem. *Hint :* use the Campbell-Mecke and the Georgii-Nguyen-Zessin formulas.

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Table of contents

Cours 6. Poisson based point processes : Cox and Boolean processes

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Cox processes Cluster processes Boolean model Capacity functional.Choquet theorem Exercises : Poisson based point processes Cours 6. Poisson based point processes.

Direct application of counting measures and Palm distributions for point process analysis

- \blacktriangleright counting measures \rightarrow summary statistics for point pattern characterization
- two categories : interpoint distances (F, G and J) and second order characteristics (ρ, K and L)
- possible extension of the summary statistics : marks, non-stationary processes, different observation spaces W case and spatio-temporal

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- non-parametrical estimation of the summary statistics : kernel estimation and management of the border effects + numerical sensitivity and computational cost
- central limit available : statistical tests
- envelope simulation based tests : reject a particular model
- summary statistics for parameter estimation of a given model :
 - these statistics are an "equivalent" of the moments in probability theory, hence they do not entirely determine the model to be estimated

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- (Baddeley and Silverman, 1984)
- (Bedford and Berg, 1997)

- good exploring tool : spatstat
- outline important characteristics of a point pattern : clustering, repulsion, completely randomness
- it is difficult to differentiate between interaction and inhomogeneity if only one realisation is available
- need for models able to reproduce these characteristics
- counting or choosing a typical point is not always obvious : consider random measure theory

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Cox processes

Definition

Let Υ be a random locally finite diffuse measure on (W, \mathcal{B}) . If the conditional distribution of X given Υ is a Poisson process on W with intensity measure Υ , X is said to be a Cox point process with driving measure Υ . Sometimes X is also called doubly stochastic Poisson process.

Remarks :

▶ if there exists a random field $Z = \{Z(w), w \in W\}$ such that

$$\Upsilon(B) = \int_B Z(w) d\nu(w)$$

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then X is a Cox process with driving function Z

► the conditional distribution of X given Z = z is a distribution of the Poisson process with intensity function z ⇒

$$\mathbb{E}[N(B)|Z=z] = \int_{B} z(w) d\nu(w)$$

the first order factorial moment measure is obtained using the law of the total expectation

$$\mu^{(1)}(B) = \alpha^{(1)}(B) = \mathbb{E}[N(B)]$$

= $\mathbb{E}[\mathbb{E}[N(B)|Z = z]] = \mathbb{E}\left[\int_{B} Z(w)d\nu(w)\right]$
= $\mathbb{E}[\Upsilon(B)] = \int_{B} \mathbb{E}Z(w)d\nu(w)$

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• if $\rho(w) = \mathbb{E}Z(w)$ exists then it is the intensity function

smilarly, it can be shown that the second order factorial moment measure is

$$\begin{aligned} \alpha^{(2)}(B_1 \times B_2) &= \mathbb{E}\left[\Upsilon(B_1)\Upsilon(B_2)\right] \\ &= \mathbb{E}\left[\int_{B_1} Z(u)d\nu(u)\int_{B_2} Z(v)d\nu(v)\right] \\ &= \mathbb{E}\left[\int_{B_1}\int_{B_2} Z(u)Z(v)d\nu(u)d\nu(v)\right] \\ &= \int_{B_1}\int_{B_2} \mathbb{E}\left[Z(u)Z(v)\right]d\nu(u)d\nu(v)\end{aligned}$$

• if $\rho^{(2)}(u,v) = \mathbb{E}Z(u)Z(v)$ exists, then it is the product density

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the pair correlation function is

$$g(u,v) = \frac{\rho^{(2)}(u,v)}{\rho(u)\rho(v)} = \frac{\mathbb{E}\left[Z(u)Z(v)\right]}{\mathbb{E}\left[Z(u)\right]\mathbb{E}\left[Z(v)\right]}$$

(日)

depending on Z it is possible to obtain analytic formulas for the second order characteristics (g, K and L) and the interpoint distance characteristic (F, G and J) the variance VarN(B) is obtained using the total variance law, and it is

$$VarN(B) = \mathbb{E}N(B) + Var\left[\int_{B} Z(w)d
u(w)
ight] \geq \mathbb{E}N(B)$$

⇒ over - dispersion of the Cox process counting variables
 the void probabilities of Cox processes are

$$\mathbb{P}(N(B) = 0)) = \mathbb{E}\{N(B) = 0\}$$

= $\mathbb{E}[\mathbb{E}\{N(B) = 0\}|Z = z]] = \mathbb{E}[\mathbb{P}(N(B) = 0|Z = z)]$
= $\mathbb{E}\left[\exp\left(-\int_{B} Z(w)d\nu(w)\right)\right]$
= $\mathbb{E}[\exp(-\Upsilon(B))]$

Trivial Cox process : mixed Poisson processes

- Z(w) = Z₀ a common positive random variable for all locations w ∈ W
- \triangleright X |Z₀ follows a homogeneous Poisson process with intensity Z₀
- the driving measure is $\Upsilon(B) = Z_0 \nu(B)$

Thinning of Cox processes :

- X is a Cox process driven by Z
- ▶ $\Pi = {\Pi(w) : w \in W} \subseteq [0,1]$ is a random field which is independent of (X, Z)
- X_{thin} | Π → the point process obtained by independent thinning of the points in X with retention probabilities Π

► \Rightarrow X_{thin} is a Cox process driven by $Z_{thin}(w) = \Pi(w)Z(w)$

Log Gaussian Cox processes

- introduced independently by astronomers (Coles and Jones, 1991) and statisticians (Møller et. al., 1998)
- consider $Y = \log Z$ is a Gaussian field
 - ▶ for any integer n > 0, locations $\xi_1, \ldots, x_n \in \mathbb{R}^d$ and numbers $a_1, \ldots, a_n \in \mathbb{R}$, $\sum_{i=1}^n a_i Y(\xi_i)$ follows a normal distribution

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the Cox process X driven by Z = exp(Y) is a log Gaussian Cox Process (LGCP) the distribution of (X, Y) is entirely determined by the mean and the covariance function

$$m(\xi) = \mathbb{E}Y(\xi)$$
 and $c(\xi, \eta) = Cov(Y(\xi)Y(\eta))$

covariance function :

for simplicity it may be considered translation invariant

$$c(\xi,\eta)=c(\xi-\eta)$$

of the form

$$c(\xi) = \sigma^2 r(\xi/\alpha)$$

► the function r : ℝ^d → [-1, 1] is a correlation function for a Gaussian field iif r is positive definite

$$\sum_{i=1}^n a_i a_j r(\xi_i, \xi_j) > 0 \quad \text{for all} \quad \xi_1, \dots, x_n \in \mathbb{R}^d, a_1, \dots a_n \in \mathbb{R}$$

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weak conditions are required on m and r in order to get

$$\Upsilon(B) = \int_B Z(\xi) d\nu(\xi)$$

for bounded $B \subset \mathbb{R}^d$. For instance, we may require $\xi \to Y(\xi)$ continuous almost surely

as example, this is satisfied by continuous m and r such that

$$r(\xi) = \exp(-\parallel \xi \parallel^{\delta}), \quad 0 \le \delta \le 2$$

with δ controlling the smoothness of the realizations of the Gaussian field

- $\delta = 1$: exponential correlation function
- $\delta = 1/2$: stable correlation function
- $\delta = 2$: Gaussian correlation function

there is a one-to-one correspondence between (m, c) and (g, ρ) Rightarrow the distribution of (X, Y) is uniquely determined by (ρ, g)

 \rightarrow Exercise 19

Cluster processes

Definition

Let C be a point process (parent process), and for each $c \in C$ let X_c be a finite point process (daughter process). Then

 $X = \bigcup_{c \in C} X_c$

is called a cluster point process.

Definition

Let X be a cluster point process such that C is a Poisson point process and conditional on C, the processes $X_c, c \in C$ are independent. Then X is called a Poisson cluster point process.

Neyman-Scott processes

Definition

Let X be a Poisson cluster point process such that centred daughter processes $X_c - c$ are independent of C. Given C, let the points of $X_c - c$ be i.i.d. with probability density function k on \mathbb{R}^d and $N(X_c)$ be i.i.d. random variables. Then X is called a Neyman-Scott process. If moreover $N(X_c)$ given C has a Poisson distribution with intensity α , then X is a Neyman-Scott Poisson process.

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 \rightarrow drawing + Exercice 20

Theorem

Let X be a Neyman-Scott Poisson process such that C is a stationary Poisson process with intensity κ . Then X is stationary process with intensity $\rho = \alpha \kappa$ and pair correlation function

$$g(u) = 1 + \frac{h(u)}{\kappa},$$

where

$$h(u) = \int k(v)k(u+v)d\nu(v)$$

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is the density for the difference between two independent points distributed according to k.

Proof. \rightarrow Exercise 21

Other very known cluster point processes

Matérn cluster process (Matérn 1960,1986)

$$k(u) = \frac{1\{\parallel u \parallel \le r\}}{\omega_d r^d}$$

is the uniform density on the ball b(o, r)

Thomas process (Thomas 1949)

$$k(u) = \frac{\exp\left(-\frac{\|u\|^2}{2\omega^2}\right)}{(2\pi\omega^2)^{d/2}}$$

is the density for $\mathcal{N}_d(0, \omega^2 I_d)$, i.e. for *d* independent normally distributed variables with mean 0 and variance $\omega^2 > 0$

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both kernels are isotropic

the Thomas process pair correlation function is

$$g(u) = 1 + rac{1}{\kappa (4\pi\omega^2)^{d/2}} \exp\left[-rac{\parallel u \parallel^2}{4\omega^2}
ight]$$

and its K-function for d = 2 is

$$\mathcal{K}(r) = \pi r^2 + \frac{1 - \exp[-r^2/(4\omega^2)]}{\kappa}$$

- other summary statistics can be also computed
- the expressions of the summary statistics are more complicated for the Matérn process
- \rightarrow drawing the processes \ldots

Remarks :

- usually in applications Z is unobserved
- one cannot distinguish a Cox process X from its corresponding Poisson process X|Z whenever a single realisation of X is available
- open question : which of the two models might be most appropriate, i.e. whether Z should be random or "systematic"/deterministic

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- prior knowledge of the observed phenomenon
- ► Bayesian setting of the intensity function ⇒ Cox processes
- if we want to investigate the dependence of certain covariates associated to Z, these may be treated as systematic terms, while unobserved effects may be treated as random terms
- Cox process: more flexible models for clustered patterns than inhomogeneous Poisson point processes

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- parameter estimation methods: minimum contrast, Palm distributions, composite likelihood
 - based on the K function and the Palm distributions
 - spatstat: the kppm function

 \rightarrow Exercise 22

Boolean model

Random objects "centred" around Poissonian points \rightarrow germs and grains

- germs : a stationary Poisson point process X of intensity ρ on \mathbb{R}^d
- grains : a sequence of i.i.d. random compact sets Γ₁, Γ₂,... and independent of X

The Boolean model is the random set obtained by the replacement of the germs by the appropriately shifted corresponding set, and taking the set union as it follows

$$\Gamma = \bigcup_{n=1}^{\infty} (\Gamma_n + w_n) = (\Gamma_1 + w_1) \cup (\Gamma_2 + w_2) \cup \dots$$

The random set Γ_0 is said to be the typical grain. The set Γ is also called the Poisson germ-grain model.

The Boolean model observation is an incomplete observation of a marked point process, since the locations points is not available



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Figure: Boolean model of random discs : marked point process (complete) and random sets perspectives.

Remarks :

- classical references : Matheron (1975), Molchanov (1997), Lantuéjoul (2002), Chiu et al. (2013)
- \blacktriangleright important practical applications \rightarrow one of the first models of complex pattern
- Neyman-Scott processes are Boolean models as well
- ► very convenient models since they allow the analytical computation of quantities describing them → this is due mainly to the independence properties allowed in the construction of the process
- independence \leftrightarrow no objects interactions \leftrightarrow no structure

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Capacity functional.Choquet theorem

- ▶ in general, for random sets it is rather difficult to use moment and factorial measures ↔ it is not possible to "count" points
- un-marked and marked point processes are particular random sets

Definition

The capacity functional the random closed set Γ is

$$T_{\Gamma}(K) = \mathbb{P}(\Gamma \cap K \neq \emptyset)$$

for K an element of the family \mathcal{K} of compact sets in \mathbb{R}^d .

Theorem

(Choquet theorem). The distribution of a random closed set Γ is completely determined by the capacity functionals $T_{\Gamma}(K)$ for all $K \in \mathcal{K}$.

Capacity functional of the Boolean model

Proposition

The capacity functional of the Boolean model Γ is

$$T_{\Gamma}(K) = 1 - \exp\left[-\rho \mathbb{E}(\nu(\check{\Gamma_0} \oplus K))\right].$$

the reflection of the typical grain :

$$\check{\Gamma_0} = -\Gamma_0 = \{-w : w \in A\}, \text{ for } A \subset \mathbb{R}^d$$

the Minkowski addition :

$$A \oplus B = \{u + v : u \in A, y \in B\}, \text{ for } A, B \subset \mathbb{R}^d$$

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Proof. \rightarrow blackboard: Exercise 23

Basic characteristics of the Boolean model

 the volume fraction : the mean fraction of volume occupied by Γ in a region of unit volume

$$p = \mathbb{P}(o \in \Gamma) = 1 - \exp[-\rho \mathbb{E}(\nu(\Gamma_0))]$$

the non-centred covariance

 $C(r) = \mathbb{P}(o \in \Gamma \text{ and } r \in \Gamma) = 2p - 1 + (1 - p)^2 \exp[
ho \eta_{\Gamma_o}]$

with $\eta_{\Gamma_o} = \mathbb{E}[\nu(\Gamma_o \cap (\Gamma_o - r))]$

contact distribution

$$H_B(r) = 1 - \frac{1 - T_{\Gamma}(rB)}{1 - p}$$

 \rightarrow Exercise 24 + some more explanations for the contact distribution

Stability of the Boolean model

Proposition

The following properties are satisfied :

i) the union of two independent Boolean models is a Boolean model

ii) a Boolean model dilated by a non-empty compact subset of \mathbb{R}^d is a Boolean model

iii) the intersection between a Boolean model and a compact subset of \mathbb{R}^d is a Boolean model

iv) the cross-section of a Boolean model by an *i*-flat is a Boolean model

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Proof.

 \rightarrow blackboard: Exercise 25

Exercises : Poisson based point processes

Exercise 19. The intensity and pair correlation function of an LGCP are given by

$$\rho(\xi) = \exp(m(\xi) + c(\xi,\xi)/2), \quad g(\xi,\eta) = \exp(c(\xi,\eta)).$$

Hint : the moment generating function of the normal distribution with mean *m* and variance σ^2 is $\exp(mt + \sigma^2 t^2/2)$.

Exercise 20. In the case of a Neyman-Scott Poisson process as defined during the course, show that X_c given C are independent Poisson processes with intensity function $\rho(w) = \alpha k(w - c)$. *Hint* : compute the void probabilities.

Exercise 21. Prove the stationarity of a Neyman-Scott Poisson process and the computation of its pair-correlation function. *Hint :* compute the void probabilities

Exercise 22. Use the **spatstat** package to compute and print estimates of the known summary statistics (second order and interpoint distances) for

- a) Thomas process with parameters $\alpha = 10$, $\kappa = 10$ and $\omega^2 = 0.01$ in a window $W = [0, 1] \times [0, 1]$.
- b) Matérn cluster process with parameters $\alpha = 10, \kappa = 10$ and r = 0.1 in a window $W = [0, 1] \times [0, 1]$.
- c) LGCP with exponential correlation function and parameters $m=4,\sigma^2=0.2,\alpha=0.1$

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- d) What is the theoretical intensity of these processes ? Do you see any differences between two realizations of these two processes, respectively ? How, can you use these observations in order to chose an appropriate model for a given data set ?
- e) Estimate the model parameters of the three precedent models on the data set redwood.
- f) Compare the three results using an envelope test based on the L function.

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Hint : Spatstat commands : rThomas, rMatClust, rLGCP,kppm, envelope. Install the package RandomFields. Play with model parameters in order to obtain different configuration topologies. *Exercise 23.* Prove the capacity functional formula of the Boolean model.

Exercise 24. Let Γ be a Boolean process on \mathbb{R}^d with intensity parameter ρ . Prove the formulas presented in the course, for the volume fraction, the covariance and the contact distribution function.

Exercise 25. Prove the stability properties of the Boolean model.

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Table of contents

Cours 7. Gibbs models.

Gibbs models Construction of the probability density Markov point processes Exercises : Gibbs models

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Gibbs models

Construction of the probability density of a point process :

- the independence property of the Poisson based processes does not allow to introduce point interactions
- Gibbs models are flexible point processes that allow the specification of point interactions via a a probability density
- In the following, let us consider (W, d) a complete, separable metric space such that W ⊂ ℝ^d and 0 < ν(W) < ∞</p>
- let μ be the unit intensity Poisson point process on W
- the condition that W has to be finite is required in order to obtain a well defined probability density for the considered Gibbs model

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- the probability density of a Gibbs model is a Radon-Nykodim derivative w.r.t the Poissonian reference measure μ
- within a statistical physics context, the probability density of such a point process has the form

$$p(\mathbf{x}) = rac{\exp[-U(\mathbf{x}|\theta)]}{\alpha(\theta)}$$

with $U(x|\theta)$ the energy function, θ the model parameters and

$$\alpha(\theta) = \int_{\Omega} \exp[-U(\mathbf{x}|\theta)] d\mu(\mathbf{x})$$

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the normalizing constant or the partition function

 under these circumstances, the probability distribution of a Gibbs model writes as

$$\mathbb{P}(X \in F) = \int_F p(\mathsf{x}) d\mu(\mathsf{x})$$

and by introducing the expression of $\boldsymbol{\mu},$ it is further expressed as

$$\mathbb{P}(X \in F) =$$

$$= \sum_{n=0}^{\infty} \frac{\exp[-\nu(W)]}{n!} \int_{W} \cdots \int_{W} \mathbb{1}(\{x_1, \dots, x_n\} \in F) \times p(\{x_1, \dots, x_n\}) d\nu(x_1) \dots d\nu(x_n),$$

whenever n > 0. If n = 0, we take $\exp[-\nu(W)]1(\emptyset \in F)p(\emptyset)$. If $\nu(W) = 0$, then $P(X = \emptyset) = 1$. For applications, we always assume that $\nu(W) > 0$.

the marked case writes in a similar way by introducing also the marks distribution v_M ► usually the probability density is known only up to a constant : p ∝ h = exp(-U)

the normalizing constant or the partition function is given by

$$\alpha = \int_{\Omega} h(\mathsf{x}) d\mu(\mathsf{x})$$

that becomes

$$\alpha = \sum_{n=0}^{\infty} \frac{\exp[-\nu(W)]}{n!} \int_{W} \cdots \int_{W} h(\{x_1, \dots, x_n\}) d\nu(x_1) \dots d\nu(x_n)$$
(2)

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- the previous quantity is not always available under analytical closed form
- this is the main difficulty to be solved while ausing this approach ...

Normalizing constant for the Poisson process : Let ρ be the intensity function of a Poisson point process on W. Its probability density up to a normalizing constant is

$$p(\mathsf{x}) \propto \prod_{x_i \in \mathsf{x}} \rho(x_i).$$

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Let $\Upsilon(B) = \int_B \rho(w) d\nu(w)$ be the associated intensity measure. By using (2), we get

$$\alpha = \exp[-\nu(W)] \sum_{n=0}^{\infty} \frac{\Upsilon(W)^n}{n!} = \exp[\Upsilon(W) - \nu(W)],$$

that gives for the complete probability density

$$p(\mathsf{x}) = \exp[\nu(W) - \Upsilon(W)] \prod_{\mathsf{x}_i \in \mathsf{w}} \rho(\mathsf{x}_i)$$

If the process is stationary $\rho(x) = \rho = \text{ct.}$, then the probability density is

$$p(\mathbf{x}) = \exp[(1-\rho)\nu(W)]\rho^n$$

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Remarks :

- the probability density is specified only for finite point processes
- ► the extension to ℝ^d of a finite point process specified by a probability density is possible under some conditions (see (Møller and Waagpetersen, 2004, section 6.4))
- two such conditions that are equivalent :
 - Iocal specification for Markov point processes
 - differential characterisation of Gibbs point process : link Palm distribution and conditional intensity

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phase transition - if such an extension is possible, does it surely leads to an unique probability measure ?

Construction of the probability density

• specify the interaction functions $\phi^{(k)}: \Omega \to \mathbb{R}^+$

$$\phi(x_{i_1},\ldots,x_{i_k})^{(k)}$$

for any k-tuplet of objects

the un-normalized probability density is the product of all these functions

$$h(\mathbf{x}) = \prod_{x_i \in \mathbf{x}} \phi(x_i)^{(1)} \cdots \prod_{\{x_{i_1}, \dots, x_{i_k}\} \in \mathbf{x}} \phi(x_{i_1}, \dots, x_{i_k})^{(k)}$$
(3)

- clearly, the energy function is obtained by taking
 U(x) = -log h(x)
- $\blacktriangleright \alpha$ the normalizing constant is difficult to be determined : untractable mathematical formula

 the un-normalized probability densities (3) are suitable for modelling provided they are integrable on Ω; that is

$$lpha = \int_{\Omega} h(\mathsf{x}) d\mu(\mathsf{x}) < \infty.$$

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► the following results ensure the integrability of the probability density of a marked point process → the Ruelle stability conditions

Definition

Let X be a marked point process given by the un-normalized probability density h w.r.t the reference measure μ . The process X is stable in the sense of Ruelle, if it exists $\Lambda > 0$ such that

$$h(\mathbf{x}) \leq \Lambda^{n(\mathbf{x})}, \quad \forall \mathbf{x} \in \Omega.$$
 (4)

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Proposition

The un-normalized probability density of a stable point process is integrable.

Proof.

The integrability of h(x) follows directly from the preceding condition :

$$\begin{split} \int_{\Omega} h(\mathbf{x})\mu(d\mathbf{x}) &\leq \int_{\Omega} \Lambda^{n(\mathbf{x})}\mu(d\mathbf{x}) \\ &= \sum_{n=0}^{\infty} \frac{\exp[-\nu(W)][\Lambda\nu(W)])^n}{n!} = \exp[\nu(W)(\Lambda-1)]. \end{split}$$

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Definition

Under the same hypotheses as in Prop. 5, a marked point process is said to be locally stable if it exists $\Lambda > 0$ such that

$$h(\mathsf{x} \cup \{\eta\}) \leq \Lambda h(\mathsf{x}), \quad \forall \mathsf{x} \in \Omega, \eta \in W \times M \setminus \mathsf{x}$$
(5)

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Proposition

A locally stable point process is stable in the sense of Ruelle.

Proof.

It is easy to show by induction that

$$h(\mathbf{x}) \leq h(\emptyset) \Lambda^{n(\mathbf{x})}, \quad \forall \mathbf{x} \in \Omega.$$

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The local stability of a point process (5) implies its integrability (4).

the conditional intensity for a point process X with probability density p is

$$\lambda(\eta; \mathsf{x}) = \frac{p(\mathsf{x} \cup \{\eta\})}{p(\mathsf{x})} = \frac{h(\mathsf{x} \cup \{\eta\})}{h(\mathsf{x})}, \quad \mathsf{x} \in \Omega, \eta \in W \times M \setminus \mathsf{x},$$

taking a/0 = 0 for $a \ge 0$

- the conditional intensity is also known in the literature as the Papangelou intensity condition (we have already meet it)
- we shall often consider functions h : Ω → [0,∞[which are hereditary

$$h(x) > 0 \Rightarrow h(y) > 0$$
, for $y \subset x$.

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 if p is hereditary, then there is a one-to-one correspondence between p and λ

Importance of the conditional intensity : key element in modelling

- plays a similar role as the conditional probabilities for Markov random fields
- integrability
- convergence properties of the MCMC algorithms used to sample from p
- the process X is attractive if $x \subseteq y$ implies

$$\lambda(\eta; \mathsf{x}) \leq \lambda(\eta; \mathsf{y}),$$

and repulsive otherwise

$$\lambda(\eta; \mathsf{x}) \geq \lambda(\eta; \mathsf{y}),$$

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- attractive processes tend to cluster the points, while the repulsive ones tend to distance the points
- these conditions are important also for exact MCMC algorithms
- there exist processes that are neither attractive nor repulsive
- there are processes that are integrable but not locally stable : Lennard - Jones (statistical physics)

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Markov point processes

The conditional intensity of an interacting point process is given by

$$\lambda(\eta;\mathsf{x}) = \phi(\eta)^{(1)} \prod_{x_i \in \mathsf{x}} \phi(x_i, \eta)^{(2)} \cdots \prod_{\{x_{i_1}, \dots, x_{i_k}\} \in \mathsf{x}} \phi(x_{i_1}, \dots, x_{i_k}, \eta)^{(k+1)}$$

- difficult to manipulate
- ▶ possible simplifications : limit the order of interactions → only pairs of points for instance
- limit the range of the interaction : a point interact only with its closest neighbours

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Let \sim be a symmetrical and reflexive relation between points belonging to $W \times M$. This may be a typical neighbourhood relation based on a metric (Euclidean, Hausdorff) or on sets intersection.

Definition

A clique is a configuration $x \in \Omega$ such that $\eta \sim \zeta$ for all $\eta, \zeta \in x$. The empty set is a clique.

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Definition

Let X be a marked point process on $W \times M$ with probability density p w.r.t the reference measure μ . The process X is Markov if for all $x \in \Omega$ such that p(x) > 0, the following conditions are simultaneously fulfilled :

(i)
$$p(y) > 0$$
 for all $y \subseteq x$ (hereditary)
(ii) $\frac{p(x \cup \{\zeta\})}{p(x)}$ depends only on ζ and $\partial(\zeta) \cap x = \{\eta \in x : \eta \sim \zeta\}.$

This process is known in the literature as the Ripley-Kelly Markov process.

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Example : The probability density w.r.t to μ of a marked Poisson process on $W \times M$ with constant intensity function $(\rho(\eta) = \beta > 0)$ is

$$p(\mathbf{x}) = \beta^{n(\mathbf{x})} \exp[(1-\beta)\nu(W)].$$

Clearly p(x) > 0 for all configurations x. Its Papangelou conditional intensity is

$$\lambda(\eta; \mathsf{x}) = \beta \mathbb{1}\{\eta \notin \mathsf{x}\}.$$

Hence, the Poisson process is Markov, independently of the interaction functions $\phi^{(k)}$. This agrees with the choice of the Poisson process for modelling a completely random structure.

The following result is known as the spatial Markov property. \rightarrow drawing

Theorem

Let X be a Markov point process with density $p(\cdot)$ on W and consider a Borel set $A \subseteq W$. Then the conditional distribution of $X \cap A$ given $X \cap A^c$ depends only on X restricted to the neighbourhood

$$\partial(A) \cap A^c = \{ u \in W \setminus A : u \sim a \text{ for some } a \in A \}.$$

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Proof. \rightarrow Exercise 26

The following result is known as the Hammersley-Clifford theorem.

Theorem

A marked point process density $p: \Omega \to \mathbb{R}^+$ is Markov with respect to the interaction relation \sim if and only if there is a measurable function $\phi_c: \Omega \to \mathbb{R}^+$ such that

$$p(\mathsf{x}) = \prod_{\text{cliques}} \phi_{\mathsf{c}}(\mathsf{y}), \quad \alpha = \phi(\emptyset) \tag{6}$$

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for all $x \in \Omega$.

Proof.

 \rightarrow Exercise 27

Remarks :

- the previous result simplifies the writing of the probability density of an interacting point process
- ▶ taking φ_c(y) = 1 whenever y is not a clique leads us to the equivalence of (3) and (6)
- Markov point processes are known in physics community as Gibbs point processes

$$p(\mathbf{x}) = \frac{1}{Z} \exp\left[-U(\mathbf{x})\right] = \frac{1}{Z} \exp\left[-\sum_{\text{cliques}} U_c(\mathbf{z})\right],$$

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with Z the partition function, U the system energy and $U_c = \log \phi_c$ the clique potential

- all the Markov processes are Gibbs
- the reciprocal is not true

Poisson process as a Markov process : the probability density of a Poisson point process is

$$p(\mathbf{x}) = e^{(1-\beta)\nu(W)} \prod_{\mathbf{x} \in \mathbf{x}} \beta.$$

Hence, the interactions functions applied to cliques are

$$\phi_c(\emptyset) = e^{(1-\beta)\nu(W)}$$

$$\phi_c(\{u\}) = \beta$$

with $\phi_c \equiv 1$ for the cliques made of more than one object. The potential of the cliques made of a single object is

$$U_c(u) = -\log\beta,$$

while $U_c = 0$ otherwise. This confirms the lack of interaction in the Poisson process. It validates also, the choice of this process to model patterns exhibiting no particular morphological structure.

Distance interaction model - Strauss model : (Strauss, 1975), (Kelly and Ripley, 1976)

$$p(\mathsf{x}) = \alpha \beta^{n(\mathsf{x})} \gamma^{\mathsf{s}_r(\mathsf{x})}, \quad \alpha, \beta > 0, \gamma \in [0, 1]$$



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Figure: Strauss model realisations for different parameter values : a) $\gamma = 1.0$, b) $\gamma = 0.5$ and c) $\gamma = 0.0$.

The interaction function $\gamma: W \times W \rightarrow [0,1]$ is

$$\gamma(u, v) = \begin{cases} \gamma & \text{if } d(u, v) \leq r \\ 1 & \text{otherwise} \end{cases}$$

The conditional intensity of adding a point η to $x \setminus \{\eta\}$ is

$$\lambda(u; \mathsf{x}) = \beta \gamma^{\operatorname{card}\partial(u)}$$

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where $\partial(u) = \{v \in x : d(u, v) \le r\}$

The Strauss model is a locally stable model with $\Lambda = \beta$ and Markov with interaction range *r*. The interaction functions applied to cliques are

$$\phi_{c}(\emptyset) = \alpha$$

$$\phi_{c}(\{u\}) = \beta$$

$$\phi_{c}(\{u,v\}) = \gamma(u,v)$$

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and $\phi_c \equiv 1$ if the cliques have three or more objects. The interaction potentials are obtained taking $U_c = -\log \phi_c$.

Multi-type pairwise interaction processes



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Figure: Bivariate pairwise interaction processes with r = 0.05 and : a) $\gamma_{1,2} = \gamma_{2,1} = 1.0$, b) $\gamma_{1,2} = \gamma_{2,1} = 0.75$ and c) $\gamma_{1,2} = \gamma_{2,1} = 0$. Circles around the points have a radius of 0.025.

Widom-Rowlinson or penetrable spheres model : this model is described by the mark space $M = \{1, 2\}$ and the density

$$p(x) = \alpha \prod_{(w,m)\in x} \beta_m \prod_{(u,1), (v,2)\in x} 1\{ \| u - v \| > r \}$$
(7)

w.r.t the standard Poisson point process on $W \times M$ with $\nu_M(1) = \nu_M(2)$. The parameters $\beta_1 > 0$ and $\beta_2 > 0$ control the number of particles of type 1 and 2, respectively. The conditional intensity for adding $(w, 1) \notin x$ to the configuration x is

$$\lambda((w,1);\mathsf{x}) = \beta_1 \mathbb{1}\{d(u,w) > r \text{ for all the } (u,2) \in \mathsf{x}\}.$$

A similar expression is available for adding an object of type 2.

The Widom-Rowlinson is hereditary and locally stable with

$$\Lambda = \max\{\beta_1, \beta_2\}.$$

Furthermore, $\lambda((w, m); x') \ge \lambda((w, m); x)$ for all $x' \subseteq x$ and $(w, m) \in W \times M$. The interaction functions are

$$\begin{aligned} \phi_c(\emptyset) &= \alpha \\ \phi_c(\{(w,m)\}) &= \beta_m \\ \phi_c(\{(u,1),(v,2)\}) &= 1\{d(u,v) > r\} \end{aligned}$$

and $\phi_{\rm c} \equiv 1$ if the cliques have two or more objects of the same type.

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Multi-type pairwise interaction process : consider $M = \{1, ..., I\}$ with $I \in \mathbb{N}$ and ν_M the uniform distribution on M. The probability density w.r.t the standard multi-type process is

$$p(\mathbf{x}) = \alpha \prod_{(w,m)\in\mathbf{x}} \beta_m \prod_{(u,i)\neq(v,j)\in\mathbf{x}} \gamma_{ij}(d(u,v)).$$
(8)

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- ► the parameters β_m > 0, m ∈ M control the intensity of the points of type m.
- ► the measurable functions γ_{ij} : [0,∞) → [0,1] describe the interaction between each type pair of objects i, j ∈ M

Symmetric functions :
$$\gamma_{ij} \equiv \gamma_{ji}$$
 for all $i, j \in M$

For $(w, m) \notin x$, the conditional intensity is

$$\lambda((w,m); \mathsf{x}) = \beta_m \prod_{(u,i) \in \mathsf{x}} \gamma_{im}(d(u,w)).$$

This process is locally stable with $\Lambda = \max_{m \in M} \beta_m$, anti-monotonic and Markov under smooth assumptions on the functions γ_{ij} . The interaction functions are

$$\begin{aligned} \phi_c(\emptyset) &= \alpha \\ \phi_c(\{(w,m)\}) &= \beta_m \\ \phi_c(\{(u,i),(v,j)\}) &= \gamma_{ij}(d(u,v)) \end{aligned}$$

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with $\phi_c \equiv 1$ for cliques of three objects and more.

Area interaction model : (Baddeley and van Lieshout, 1995)

$$p(\mathbf{x}) \propto \beta^{n(\mathbf{x})} \gamma^{-\nu[\Gamma(\mathbf{x})]}, \quad \beta, \gamma > 0$$
(9)

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Figure: Area interaction model realisations for different parameter values : a) $\gamma = 1.0$, b) $\gamma > 1.0$ and c) $\gamma < 1.0$.

Remarks :

- ► the first probability density based point process producing clusters → alternative to the Strauss process ...
- the model should be re-parametrized in order to be identifiable

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Proposition

The area interaction process given by (9) is a Markov point process.

Proof. \rightarrow Exercice 28 + Exercice 29

Candy model :

(van Lieshout and Stoica, 2003), (Stoica, Descombes and Zerubia, 2004)

$$p(\mathbf{x}) \propto \gamma_f^{n_f(\mathbf{x})} \gamma_s^{n_s(\mathbf{x})} \gamma_d^{n_d(\mathbf{x})} \gamma_o^{n_o(\mathbf{x})} \gamma_r^{n_r(\mathbf{x})},$$

with $\gamma_f, \gamma_s, \gamma_d > 0$ and $\gamma_o, \gamma_r \in [0, 1]$



Figure: Candy model realisations.

Bisous model : (Stoica, Gregori and Mateu, 2005)

 $p(\mathbf{x}) \propto \left[\prod_{s=0}^{q} \gamma_{s}^{n_{s}(\mathbf{x})} \right| \prod_{\kappa \in \Gamma \subset \mathcal{R}} \gamma_{\kappa}^{n_{\kappa}(\mathbf{x})} \quad \gamma_{s} > 0, \gamma_{\kappa} \in [0, 1]$



Figure: Random shapes generated with Bisous model.

Remarks :

- ► Candy and Bisous are based on compound interactions → drawing + explanations
- connections are produced by giving different weights for the repulsive interactions
- the conditional intensity is bounded

$$\lambda(\zeta; \mathsf{x}) \leq \prod_{s=0}^{q} \max\{\gamma_{s}, \gamma_{s}^{-1}\}^{12} = \mathsf{\Lambda}.$$

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this gives the name of the model \rightarrow kissing number

- \blacktriangleright \rightarrow blackboard Candy
- Markov range : $4r_h + 2r_a$
- the models are locally stable but the exact simulation is sometimes difficult ...

Compare two random sets : idea inspired by work with M. N. M. van Lieshout and classical literature in mathematical morphology





Figure: Realizations of the Candy model obtained with different samplers.

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Empty space function : these probability distributions should be similar \Rightarrow Kolmogorov-Smirnov p- value is higher than 0.8



Empty space functions for Candy patterns

Figure: Estimation of the empty space function for the previous Candy realizations

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Exercises : Gibbs models

Exercise 26. Prove the Spatial Markov property Theorem.

Exercise 27. Prove the Hammersley - Clifford Theorem.

Exercise 28. Let us consider the area-interaction point process as defined in the course.

- a) Write its conditional intensity.
- b) Prove that the model is locally stable and for $\beta, \gamma > 0$.
- c) What kind of pattern is generated if $\gamma < 1$, $\gamma = 1$ and $\gamma > 1$?

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d) Prove the model is Markov and exhibit its interaction functions. What order are these interactions ?

Exercise 29.

- a) Simulate attractive and repulsive area-interaction processes in $W = [0, 1]^2$. Use the R spatat function : rmh.
- b) To an attractive and repulsive pattern obtained previously fit the area-interaction process. What do you notice ? Are the differences between the theoretical and estimated parameters important ? Why ? Use the R spatat function : ppm.
- c) Consider now the data set redwood. After en exploratory analysis of the data set, fit an area-interaction process to it. Test the model using envelope tests. How do you interpret the results ? How do you compare these results with the ones obtained at the Exercice 22 ?

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Table of contents

Cours 8. Markov chains. Definitions, properties, algorithms.

Monte Carlo simulation Markov chains : a little bit of theory Metropolis-Hastings algorithm MH algorithm for sampling marked point processes Perfect or exact simulation Spatial birth-and-death processes Exercises : Markov chains. Definitions, properties, algorithms.

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Markov chain Monte Carlo algorithms

Problem : sampling or simulation probability distributions

$$\pi(A) = \int_A p(\mathsf{x}) d\mu(\mathsf{x})$$

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that are not available in closed form \leftrightarrow normalising constant analytically intractable

Basic MCMC algorithm

Algorithm

$$\begin{array}{l} x = My \ \textit{first} \ \textit{MCMC} \ \textit{sampler} \ (T) \\ 1. \ \textit{choose an initial condition} \ x_0 \\ 2. \ \textit{for} \ i = 1 \ \textit{to} \ T, \ \textit{do} \\ & \begin{cases} \\ & x_i = \textit{Update}(x_{i-1}) \\ & \end{cases} \end{array} \end{array}$$

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3. return x_T .

Principles of the MCMC algorithm :

- simulates a Markov chain
- the Update function reproduces the transition kernel of the Markov chain
- ▶ the output x_T is asymptotically distributed according to π whenever $T \to \infty$
- \blacktriangleright if the simulated Markov chain has good properties \rightarrow statistical inference is possible
- several solutions : Gibbs sampler, Metropolis-Hastings, birth and death processes, stochastic adsorption, RJMCMC, exact simulation (CFTP, clan of ancestors, etc.)

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Markov chains : a little bit of theory

Let $(\Omega, \mathcal{F}, \mu)$ a probability space.

Markov chain : a sequence of random variables $\{X_n\}$ such that :

$$\mathbb{P}(X_{n+1}|X_0,\ldots,X_n)=\mathbb{P}(X_{n+1}|X_n)$$

The chain is homogeneous if the probabilities from going from one state to another do not change in time.

Transition kernel : a mapping $P : \Omega \times \mathcal{F} \rightarrow [0, 1]$ such that

- ► $P(\cdot, A)$ is measurable for any $A \in \mathcal{F}$
- ► $P(x, \cdot)$ is a probability measure on (Ω, \mathcal{F}) for any fixed $x \in \Omega$

As on discrete spaces the *n*-step transition probability kernel is defined iteratively. Let $P^0(x, A) = \delta_x(A)$ the Dirac measure defined by

$$\delta_{\mathsf{x}}(A) = \begin{cases} 1 & \mathsf{x} \in A \\ 0 & \mathsf{x} \notin A, \end{cases}$$

and, for $n \ge 1$ the n-step transition probability kernel is defined inductively

$$P^n(\mathsf{x}, A) = \int_{\Omega} P(\mathsf{x}, d\mathsf{y}) P^{n-1}(\mathsf{y}, A) \quad \mathsf{x} \in \Omega, A \in \mathcal{F}.$$

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Theorem: Chapman-Kolmogorov equations. For any m with $0 \le m \le n$,

$$P^n(\mathsf{x},A) = \int_\Omega P^m(\mathsf{x},d\mathsf{y})P^{n-m}(\mathsf{y},A), \quad \mathsf{x}\in\Omega, A\in\mathcal{F}.$$

Proof: See (Meyn and Tweedie 2009, Thm. 3.4.2 pp.61). It is easy to see that

$$P^{n}(\mathbf{x}, A) = \int_{\Omega} P(\mathbf{x}, d\mathbf{y}) P^{n-1}(\mathbf{y}, A)$$
$$\int_{\Omega} P(\mathbf{x}, d\mathbf{y}) \int_{\Omega} P(\mathbf{y}, d\mathbf{z}) P^{n-2}(\mathbf{z}, A)$$
$$\int_{\Omega} P^{2}(\mathbf{x}, d\mathbf{z}) P^{n-2}(\mathbf{z}, A)$$
$$\vdots$$
$$\int_{\Omega} P^{m}(\mathbf{x}, d\mathbf{y}) P^{n-m}(\mathbf{y}, A).$$

- in a very intuitive way, this result states that to get from x to A in n + m steps, the chain must pass through some y on the nth step
- in the discrete case the Chapman-Kolmogorov equations may be interpreted as a matrix product
- in the general case the kernel Pⁿ is an operator acting on both bounded measurable functions and σ-finite measures μ on F via

$$P^n f(\mathsf{x}) = \int_{\Omega} P^n(\mathsf{x}, d\mathsf{y}) f(\mathsf{y}), \quad \mu P^n(A) = \int_{\Omega} \mu(d\mathsf{x}) P^n(\mathsf{x}, A).$$

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Definition: invariant measure. A σ -finite measure on (Ω, \mathcal{F}) is invariant, if it verifies

$$\pi(A) = \int_\Omega \pi(d\mathsf{x}) \mathsf{P}(\mathsf{x},A), \quad A \in \mathcal{F}.$$

- if an initial condition is sampled according to the invariant distribution π, then the action of the transition kernel will produce new states always distributed according to π
- simple verification:

$$\pi(A) = \int_{\Omega} \pi(d\mathsf{x}) P(\mathsf{x}, A) = \int_{\Omega} \left[\int_{\Omega} \pi(d\mathsf{y}) P(\mathsf{y}, d\mathsf{x}) \right] P(\mathsf{x}, A)$$
$$= \int_{\Omega} \pi(d\mathsf{y}) \left[\int_{\Omega} P(\mathsf{y}, d\mathsf{x}) P(\mathsf{x}, A) \right] = \int_{\Omega} \pi(d\mathsf{y}) P^{2}(\mathsf{y}, A)$$
$$\vdots$$
$$= \int_{\Omega} \pi(d\mathsf{y}) P^{n}(\mathsf{y}, A),$$

for any *n* and all $A \in \mathcal{F}$.

- \blacktriangleright now, consider any starting distribution μ
- if a limiting measure γ_μ exists in a suitable topology on the space of probability measures, such as

$$\gamma_{\mu}(A) = \lim_{n \to \infty} \int_{\Omega} \mu(d\mathsf{x}) P^{(n)}(\mathsf{x}, A)$$

for sets $A \in \mathcal{F}$

then

$$egin{array}{rcl} \gamma_{\mu}(\mathcal{A}) &=& \lim_{n o \infty} \int_{\Omega} \mu(d\mathsf{x}) \int_{\Omega} \mathcal{P}^{n-1}(\mathsf{x},d\mathsf{y}) \mathcal{P}(\mathsf{y},\mathcal{A}) \ &=& \int_{\Omega} \gamma_{\mu}(d\mathsf{y}) \mathcal{P}(\mathsf{y},\mathcal{A}), \end{array}$$

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hence γ_{μ} is an invariant probability measure.

- \blacktriangleright if the chain has an unique invariant measure π
- then
 - the limit γ_{μ} equals π
 - $\blacktriangleright\,$ it is independent of any initial condition $\mu\,$
- in this case, the invariant measure is the equilibrium distribution of the chain

Definition: reversibility. A transition kernel P(x, A) is said to be reversible with respect to a measure π , if the integral

$$\int_{A} \int_{B} \mathbf{1}_{A}(\mathsf{x}) \mathbf{1}_{B}(\mathsf{y}) \pi(d\mathsf{x}) P(\mathsf{x}, d\mathsf{y})$$
(10)

is symmetric under the interchange of A and B.

 applying the previous definition to the formula (10) leads directly to

$$\int_{A} \pi(d\mathsf{x}) P(\mathsf{x}, B) = \int_{B} \pi(d\mathsf{x}) P(\mathsf{x}, A)$$

hence, the probability of going from A to B equals the probability of going backwards from B to A. • if $P(\mathbf{x}, \Omega) = 1$ and $B = \Omega$, it follows immediately

$$\pi(A) = \int_{\Omega} \pi(d\mathsf{x}) P(\mathsf{x}, A),$$

hence, the reversibility of the transition kernel guarantees the invariance of the considered measure

- practical point of view: the Update() mechanism should reproduce a reversible transition kernel
- ► the reversibility condition alone → only the construction of a sampling algorithm that needs the initial conditions to be chosen according to the distribution of interest
- problem solved: if the Markov chain is irreducible, aperiodic and recurrent

Irreducibility

- crucial property in the setup of MCMC algorithms
- Intuitive description: guarantees that from any initial point x ∈ Ω, the chain may reach any region A ∈ F of the configuration space
- first measure of the sensitivity of the Markov chain to the initial conditions
- it leads to convergence conditions of the simulated chain towards the desired equilibrium distribution.
- for presenting it, we need first the definition of the stopping time of the chain in a set A.

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Definition: stopping time at A. For any set $A \in \mathcal{F}$, the quantity

$$\tau_A = \min\{n \ge 1 : X_n \in A\}$$

is called the stopping time at A. If $X_n \notin A$ for every *n*, then by convention $\tau_A = +\infty$.

 discrete case: irreducibility means, that all the chain states communicate, that is

$$\mathbb{P}_{\mathsf{x}}(\tau_{\mathsf{y}} < \infty) > 0 \quad \forall \mathsf{x}, \mathsf{y} \in \Omega.$$

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- continuous state space: $\mathbb{P}_{x}(\tau_{y} < \infty) = 0$
- ▶ in order to correctly define irreducibility for general state spaces, an auxiliary measure φ is needed

Definition: ϕ -irreducibility. The Markov chain (X_n) is ϕ -irreducible if there exists a measure ϕ on \mathcal{F} such that, whenever $\phi(A) > 0$, we have $\mathbb{P}_x(\tau_A < \infty) > 0$ for all $x \in \Omega$.

An equivalent formulation of the definition of the ϕ -irreducibility is that for all $x \in \Omega$, whenever $\phi(A) > 0$, there exists n > 0 such that $P^n(x, A) > 0$.

- a φ-irreducible Markov chain is able to reach any set A which is "big enough", independently of the initial condition
- in the following we will give the tools needed for establishing the irreducibility of a transition kernel

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Definition: small sets. A set *C* is a small set if there exists $n \in \mathbb{N}^*$ and a nonzero measure ν_n such that

$$P^{n}(\mathbf{x}, A) \ge \nu_{n}(A) \tag{11}$$

for all $x \in C$ and $A \in \mathcal{F}$. Whenever the equation (11) is verified, the set C is called ν_n -small.

Theorem. Let (X_n) be a ϕ -irreducible chain. For every set $A \in \mathcal{F}$ such that $\phi(A) > 0$, there exists $n \in \mathbb{N}^*$ and a small set $C \subset A$ such that $\nu_n(C) > 0$. Moreover, Ω can be decomposed in a countable partition of small sets.

- ▶ proof: (Meyn and Tweedie, 2009) Prop. 5.2.4 pp.105-106
- the small sets are tool for "discretizing" a continuous space state

- communicating states (*i.e.* irreducibility): desirable property whenever building Markov chain for sampling probability distributions
- cyclic behaviour: naturally considered as undesirable

Definition: aperiodic chain. Let $P(\cdot, \cdot)$ be a ϕ -irreducible transition kernel. Suppose that there is a set $A \in \mathcal{F}$, a probability measure ν with $\nu(A) = 1$, a constant $\epsilon > 0$ and an integer $n_0 \ge 1$ such that

$$P^{n_0}(\mathsf{x},\cdot) \geq \epsilon \nu(\cdot) \quad \forall \mathsf{x} \in A.$$

The induced Markov chain is aperiodic if

 $gcd\{m: \exists \quad \epsilon_m > 0 \quad such that \quad P^m(\mathbf{x}, \cdot) \geq \epsilon_m \nu(\cdot) \quad \forall \mathbf{x} \in A\} = 1.$

 practical consequence: aperiodic Markov chains may be constructed by simply allowing the Update procedure to remain in the initial state with a positive probability Theorem. Suppose the chain (X_n) has invariant probability measure π . Assume that the chain is π -irreducible and aperiodic. Then there is a set $\Omega' \subseteq \Omega$ such that $\pi(\Omega') = 1$ and

$$\sup_{A\in\mathcal{F}} |P^n(\mathbf{x},A) - \pi(A)| \to 0$$
 (12)

for each $x \in \Omega'$.

- proof: (Athreya et al. 1996), Thm. 1 pp. 72
- the results allows to build Update() procedures to sample from π
- the result does not prevent the Markov chain of a set of configurations Ω" with π(Ω") = 0 for which the limit in (12) differs from zero

Recurrence

- the recurrence is the Markov chain property that guarantees the non-existence of null-sets, such as Ω".
- it guarantees the independence of a MCMC sampling algorithm with respect to the initial conditions

Definition: number of passages. For any set $A \in \mathcal{F}$, the quantity

$$\eta_A = \sum_{n=1}^{\infty} \mathbb{1}_A(X_n)$$

is the number of passages of (X_n) in A.

Definition: Harris recurrence. A set A is Harris recurrent if $\mathbb{P}_{x}(\eta_{A} = \infty) = 1$ for all $x \in A$. The chain (X_{n}) is Harris recurrent if it is ϕ -irreducible and if any set $A \in \mathcal{F}$ such that $\phi(A) > 0$ is Harris recurrent.

- ▶ Harris recurrence is a "better" property than ϕ -irreducibility
- \blacktriangleright in practice, the Update mechanisms that are $\phi-{\rm irreducible},$ they are also Harris recurrent
- the following result presents: mathematical tool for proving Harris recurrence, based on the so-called drift condition

Theorem. Let (X_n) be a ϕ -irreducible Markov chain and suppose there exist a small set $C \in \mathcal{F}$ and function $V : \Omega \to (0, \infty)$ such that the level sets

$$C_V(\alpha) = \{ \mathsf{x} \in \Omega : V(\mathsf{x}) \le \alpha \}$$

are small. The chain is Harris recurrent if the drift riangle V(x) = PV(x) - V(x) is negative for any $x \notin C$, that is :

$$PV(\mathsf{x}) = \int_{\Omega} P(\mathsf{x}, d\mathsf{y})V(\mathsf{y}) \leq V(\mathsf{x}), \quad \mathsf{x} \notin C.$$

▶ proof: (Meyn and Tweedie, 2009), Thm. 9.1.8 pp. 206.

Ergodicity

- whenever a Harris recurrent Markov chain is simulated through an Update procedure, LLN and CLT can be used with the obtained samples, only after the chain reaches the equilibrium
- ▶ for the non-perfect MCMC algorithms, *i.e.* this is impossible
- the only thing we know: the chain should reach the equilibrium, but we do not know exactly when this will happen
- \blacktriangleright possible solution to this problem \rightarrow build an ergodic chain
- no need to wait till equilibrium
- the only thing needed: enough samples in order to be able to apply the LLN or the CLT apply the LLN or the CLT

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Definition: ergodicity. The Markov chain (X_n) is ergodic if it is both Harris recurrent and aperiodic.

- for ergodic chains, (12) holds independently of the initial conditions
- ► the speed of convergence of the chain may be the same for all the initial conditions → the chain is uniformly ergodic
- if the speed of convergence depends on the starting a point, we may have a geometrically ergodic chain

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Definition: total variation norm. The total variation norm of a bounded signed measure ν on (Ω, \mathcal{F}) is defined as

$$\parallel \nu \parallel = \sup_{A \in \mathcal{F}} \nu(A) - \inf_{A \in \mathcal{F}} \nu(A).$$

The total variation distance between two such measures ν_1 and ν_2 is $\parallel \nu_1 - \nu_2 \parallel$.

Theorem. Let (X_n) be a Markov chain ϕ -irreducible and aperiodic. The chain is geometrically ergodic if there exists a function $V : \Omega \to [1, \infty)$, constants $b < \infty$ and a < 1, and a small set $C \in \mathcal{F}$ such that

$$PV(\mathbf{x}) \le aV(\mathbf{x}) + b\mathbf{1}_{C}(\mathbf{x}), \quad \forall \mathbf{x} \in \Omega.$$
 (13)

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- proof: (Meyn and Tweedie, 2009), Thm. 15.0.1 pp. 363
- geometric ergodicity means that the iterations of the transition kernel approach the equilibrium distribution at a geometric speed
- it can be shown, that the sets C_V(α) are small for any α > 0
 → the geometric ergodicity drift condition implies the drift condition for Harris recurrence.

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Metropolis-Hastings algorithm

Principle :

- consider the chain in the state $x_i = x$
- ▶ propose a new state $x_f = y$ using the proposal density $q(x_i \rightarrow x_f)$
- accept this new state with probability

$$\alpha(x,y) = \min\left\{1, \frac{p(y)q(y \to x)}{p(x)q(x \to y)}\right\}$$

if not remain in the previous state

iterate as many times as we need (... in theory till infinity ...)

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Properties

- ▶ $\alpha(\cdot, \cdot)$ is a solution of the detailed balance equation \rightarrow reversibility is preserved
- ▶ very few conditions are required for q(· → ·) so that the chain has all the convergence properties

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- ▶ $q(\cdot \rightarrow \cdot)$ should be simple to calculate and to simulate
- the knowledge of the normalizing constant of $p(\cdot)$ is not needed

\rightarrow Exercise 30

MH algorithm for sampling marked point processes

Idea : the transition kernel propose to add an object to the configuration with probability p_b or propose to delete an object from the configuration with the probability p_d

Birth : add an object

• initial state : $x_i = x$ an object configuration

• final state :
$$x_f = \mathbf{x} \cup \{\zeta\}$$

proposal density to add an object : choose uniformly its location in W and its mark independently according to v_M

$$q(x_i \to x_f) = q(\mathsf{x} \to \mathsf{x} \cup \{\zeta\}) = p_b \frac{1\{\zeta_w \in W\}}{\nu(W)}$$

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Proposal density to remove an object : choose uniformly an object from x ∪ {ζ}

$$q(x_f \rightarrow x_i) = q(\mathsf{x} \cup \{\zeta\} \rightarrow \mathsf{x}) = p_d \frac{1\{\zeta \in \mathsf{x} \cup \{\zeta\}\}}{n(\mathsf{x}) + 1}$$

acceptance probability

$$\alpha(\mathsf{x} \to \mathsf{x} \cup \{\zeta\}) = \min\left\{1, \frac{p_d p(\mathsf{x} \cup \{\zeta\})}{p_b p(\mathsf{x})} \times \frac{\nu(W)}{n(\mathsf{x}) + 1}\right\} \quad (14)$$

Death : remove an object

- the inverse movement of birth
- acceptance probability

$$\alpha(\mathsf{x} \to \mathsf{x} \setminus \{\zeta\}) = \min\left\{1, \frac{p_b p(\mathsf{x} \setminus \{\zeta\})}{p_d p(\mathsf{x})} \times \frac{n(\mathsf{x})}{\nu(K)}\right\}$$
(15)

A transition kernel doing these transformations is

$$P(\mathbf{x}, A) = p_b \int_{\mathcal{K}} b(\mathbf{x}, \eta) \alpha(\mathbf{x}, \mathbf{y} := \mathbf{x} \cup \{\eta\}) \mathbf{1}\{\mathbf{y} \in A\} d\sigma(\eta)$$

+ $p_d \sum_{\eta \in \mathbf{x}} d(\mathbf{x}, \eta) \alpha(\mathbf{x}, \mathbf{y} := \mathbf{x} \setminus \{\eta\}) \mathbf{1}\{\mathbf{y} \in A\}$
+ $\mathbf{1}\{\mathbf{x} \in A\} \left[1 - p_b \int_{\mathcal{K}} b(\mathbf{x}, \eta) \alpha(\mathbf{x}, \mathbf{x} \cup \{\eta\}) d\sigma(\eta) - p_d \sum_{\eta \in \mathbf{x}} d(\mathbf{x}, \eta) \alpha(\mathbf{x}, \mathbf{x} \setminus \{\eta\}) \right],$

where $K = W \times M$, $d\sigma(\eta) = d\sigma((w, m)) = d\nu(w) \times d\nu_M(m)$ et $0 < p_b + p_d \le 1$. The birth rate is $b(x, \eta) = \frac{1}{\nu(W)}$ and the death rate is $d(x, \eta) = \frac{1}{n(x)}$

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Remarks :

▶ the Papangelou intensity appears in the acceptance probability

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- local stability property guarantees good convergence properties of the Markov chain
- \blacktriangleright \rightarrow blackboard : discuss reversibility

Algorithm

- y = Update(x)
 - Choose "birth" or "death" with probabilities p_b and p_d, respectively.
 - If "birth" was chosen, then generate a new object following b(x, η). Accept the new configuration, y = x ∪ {η} with the probability α(x, y) given by (14).
 - If "death" was chosen, then select the object to be removed using d(x, η). Accept the new configuration, y = x \ {η} with the probability α(x, y) given by (15).

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4. Return the present configuration.

Theorem. Let be b, d and q as described previously. Assume that $b(x, \eta)$ and $d(x, \eta)$ are strictly positive on their corresponding definition domain, respectively, and

$$\lim_{n\to\infty} u_n = \lim_{n\to\infty} \left[\sup_{\eta\in W\times M, x\in\Xi_n} \frac{d(x\cup\{\eta\},\eta)}{b(x,\eta)} \right] \to 0.$$

Fix $p_b, p_d \in (0, 1)$ with $p_b + p_d \leq 1$ and let p(x) be the probability density of a marked point process on $W \times M$. The point process is locally stable and p(x) is built w.r.t the standard Poisson process μ . The MH sampler defined previously simulates a Markov chain with invariant measure $\pi = \int p d\mu$ who is ϕ -irreducible, Harris recurrent and geometric ergodic.

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 \rightarrow proof: Exercise 31

Remark :

► the same result holds if change moves are introduced with care ... → explain ...

Optimality of the MH dynamics

- theoretical convergence properties
- local computation
- no need of the normalising constant
- highly correlated samples : only one element changed per accepted transition
- allows improvements : transition kernels that "help" the model

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Tailored to the model proposal distribution

$$b(\mathbf{x},\eta) = \frac{p_1}{\nu(K)} + p_2 b_a(\mathbf{x},\eta),$$

with $p_1 + p_2 = 1$ and $b_a(x, \eta)$ a probability density given by

$$b_a(\mathbf{x},\eta) = rac{1}{n(A(\mathbf{x}))} \sum_{\mathbf{x} \in A(\mathbf{x})} \tilde{b}(\mathbf{x},\eta).$$

- the role of b_a(x, η) : propose the birth of a new pointin those regions where the interactions between the new born and the other configuration members is favoured or not penalised by the model
- A(x) : the set of marked points in a configuration that are not exhibiting yet "good" interactions



Figure: Extremities marked by triangles are connected and further than $\frac{1}{2}I_{\text{max}} + r_c$ to the boundary, those labeled by a black disk are closer than $\frac{1}{2}I_{\text{max}} + r_c$ to the boundary of K.
MH algorithm for sampling the Candy model : dynamics behaviour through the sufficient statistics analysis



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- great adaptability and theoretical convergence
- easy to use
- appropriate solutions need to be found for each new model
- the general framework, even if it has good theoretical properties, it is not always the most efficient from a numerical point of view
- still, if no good theoretical properties are available, the results will be always bad

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 \rightarrow Exercise 32

Perfect or exact simulation

+ la schioapa sunt \ldots

- \blacktriangleright classical MCMC methods \rightarrow need a theoretical infinite time till convergence
- dependence on the initial conditions
- perfect or exact MCMC methods indicate by themselves whenever the convergence is attained
- these methods are perfect within the limits of the random number generators of the computers

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historical paper : (Propp and Willson, 1996)

Let us build a MCMC sampler for π defined on the discrete state space $\Omega = \{\omega_1, \omega_2, \dots, \omega_m\}$. The induced Markov chain (X_n) is represented by its transition functiones $\phi(\cdot, \cdot)$ such that

$$X_{n+1} = \phi(X_n, V_n), \tag{16}$$

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where V_n are i.i.d random variables.

Key idea :

- consider *m*, $(X_n(\omega_i))$ all initialised with a different state, that evolve from -T < 0 till 0
- the chains are coupled : they use the same V_n s
- If at a certain moment n ∈ −T,...,0 all the chains are in the same state or they coalesced, that is

$$X_n \equiv x$$

then they will all remain in the same state, till the time 0

- the influence of the initial conditions just ... vanished
- ▶ if the chains are started before the time T, at infinite, the chains will be all in the same state, at the same moment
- it comes out that X_0 is a perfect sampler from the equilibrium distribution π
- \rightarrow blackboard : drawing

Extraordinary smart idea :

- launching m parallel chains is not always feasible
- if Ω can be ordered

$$\omega_{\min} = \omega_{(1)} < \omega_{(2)} < \ldots < \omega_{(m)} = \omega_{\max}$$

and if the transition kernel respect this order relation

$$\omega \le \omega' \quad \Rightarrow \quad \phi(\omega) \le \phi(\omega')$$

then only the states ω_{\min} and ω_{\max} are needed

the behaviour of the other chains is bounded by the extremal chains X_n(ω_{min}) and X_n(ω_{max})

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This idea is known under the name Coupling From The Past (CFTP).

Theoretical background : continuous time Markov chains \rightarrow the very nice book of S. Resnick (2005)

- history : the first MCMC sampler for marked point processes
- the simulation of locally stable marked point process is rather simple : thinning procedure

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the simulated pattern is "hidden" in a dominating Poisson process

Algorithm

Let p be the probability density w.r.t. the standard Poisson process, of a locally stable marked point process on $W \times M$. Its corresponding Papangelou conditional intensity bound is Λ . Set $X^{(0)} = x_0$ for some configuration $x_0 \in \Omega$ with $p(x_0) > 0$. For $n = 0, 1, ..., \text{ if } X^{(n)} = x, \text{ do } :$

- ► the sojourn time T⁽ⁿ⁾ in the state X⁽ⁿ⁾ is exponentially distributed with mean 1/(n(x)+Aν(W))
- the next transition is a death with probability n(x)/n(x)+Λν(W), obtained by selecting one point η from the current configuration with probability 1/n(x), and then deleting it, that is x = x \ {η}
- with probability $\frac{\Lambda\nu(W)}{n(x)+\Lambda\nu(W)}$ the next transition is a birth, obtained by generating a new point η according to the probability density $\frac{1}{\nu(W)}$ and accepting it with probability $\frac{\lambda(\eta;x)}{\Lambda}$, hence $x = x \cup \{\eta\}$,

Remarks :

- this algorithm has equivalent convergence properties compared to MH algoritm (van Lieshout 2000, Thm. 3.3 pp. 86; Møller and Waagepetersen 2004, Prop. G7 pp. 276)
- nevertheless, for strong interactions Λ may be very high
- Candy and Bisous models with strong interactions cannot be simulated with this algorithm
- but ... extremely important tool in building perfect simulation algorithms

Perfect algorithms for sampling marked point processes

Coupling From The Past algorithms :

- dominating process :
 - ▶ stationary Poisson \rightarrow (CFTP and clan of ancestors), Λ parameter is important
 - it runs into the past, and then into the future till time 0 : doubling time scheme till coalescence
- the coupling and transition function ingredients : the V_ns, conditional intensity and Λ
- spatial birth-and-death based algorithms : CFTP and clan of ancestors
- Metropolis-Hastings based algorithms : different dominant process + control of the birth and death proposal probabilities
- apply to locally stable point processes
- monotonic and anti-monotonic point processes

Gibbs sampler algorithm :

- ► discrete probability densities → approximation of the continuous marked point processes probability density
- does not require : order relation, dominating process, monotonic or anti-monotonic relation
- still the algorithm is more efficient if these properties are exhibited by the considered model
- Potts like models

Remarks :

CFTP algorithm is implemented within the spatstat package

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all these algorithms are implemented within the MPPLIB
 C++ package

Strauss model : convergence speed for exact sampling methods (van Lieshout and Stoica, 2006)



Figure: Exact simulation algorithms applied to Strauss model : a) CFTP, b) clan of ancestors, c) Metropolis-Hastings and d) Gibbs.

Comparison exact simulation and MH algorithm for the Strauss model (1)



Comparison Strauss statistics distributions

Figure: Boxplots comparison for the n and s_r statistics distributions : white - the distributions obtained using the exact algorithm, pink (dark couloured) - the distributions obtained using the Metropolis - Hastings algorithm.

Comparison exact simulation and MH algorithm for the Strauss model (2)



Figure: Qqplot comparison for the n and s_r statistics distributions.

 \rightarrow Exercise 33

Open questions MCMC methods :

Classical algorithms - MH based dynamics

- good convergence properties but convergence at infinity
- burning-in methods + de-correlation techniques
- great adaptability : tailored to the model moves
- manipulate several objects during one move : work of X. Descombes
- Ink with RJMCMC : great adaptability, but difficult to state convergence proofs, hence difficult to use ...

Perfect simulation algorithms

- the simulated chain indicates by itself whenever convergence is reached
- parameter dependence : can be applied in practice only to a restricted range of parameters
- neither change moves, nor tailored moves
- study existing algorithm : Fill algorithm, forward simulation and simulated tempering
- ► challenging perspective : synthesis of both families of algorithms → exact algorithms able to be tailored to the model

Exercises : Markov chain Monte Carlo algorithms

Exercise 30. Look at the following code lines and explain the role of each function :

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```
q.prop = function (x)
ł
delta=0.1;
lim=0.5*delta;
res=runif(1,x-lim,x+lim);
}
q.density = function (x,y)
Ł
delta=0.1;
lim=0.5*delta;
```

```
res=dunif(y,x-lim,x+lim);
}
```

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```
p.density = function (y)
ſ
 d1=100;
 d2=100;
 if(y>=0)
   { res=df(y,d1,d2); }
 else
   { res=0; }
}
x0=0.5;
m=10;
n=1000;
x=1:n;
for (i in 1:n)
ł
  x[i]=algo.mh(x0,m);
  x0=x[i];
}
```

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Using the previous code, answer the following questions :

- a) Simulate n = 1000 random variables distributed according to a Fisher distribution $F(\nu_1, \nu_2)$ of parameters $\nu_1 = \nu_2 = 100$.
- b) Plot the histogram of the obtained values. On the same plot, add the theoretical density. Plot the empirical cross-correlation function.
- c) For reducing the correlation of the samples obtained using the Metropolis-Hasting algorithm, one common techniques is to separate the samples; Re-do this exercice, by taking the samples every $m = \{1, 5, 10, 50, 100\}$ iterations. Interpretation of the obtained results.

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- d) Repeat the exercise for $\delta = 0.001, 1.0, 100$. If needed the very first samples given by the algorithm can be droped of. This is called *burn-in* time. Try different values for it.
- e) Simulate n = 1000 random variables following a Fisher distribution with parameters $\nu_1 = \nu_2 = 100$. In this case, we should obtain E[X] = 1.02. What is the value of the variance we should obtain ? Build confidence intervals for the samples mean. Make a statistical test to verify that the simulated variables have the desired theoretical mean.
- f) Plot the evolution of the mean, the confidence intervalls, the p-values and the empirical levels for the confidence intervals and the tests, while the number of samples increases.
- g) Suppose that we do not know what is the sampled distribution, build a test to verify the values obtained from the empirical mean. As before, build and plot confidence intervals, statistical tests and empirical levels for the samples mean.

Exercise 31. Prove the convergence of the MH sampler for marked point processes.

Exercice 32. Look at the following code lines and explain the role of each function :

a) The previous setting of the MH dynamics saves all the intermediate patterns. To have access to these patterns use : patternX=attr(X,"saved")

Make a plot of the evolution of the sufficient statistics of the model.

- b) If $\gamma = 1$ in a Strauss model a Poisson point process is simulated. Use the sufficient statistics plots in order to find a parameter setting of the MH dynamics that may indicate convergence of the algorithm.
- c) Simulate different realisations of the Strauss model for different model parameters. item For $\beta = 100$ and $\gamma = 1$ build confidence intervals and statistical tests in order to verify the obtained values for the mean of the sufficient statistics.

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d) Repeat the previous question but for $\beta=$ 50, $\gamma=$ 0.5 and r= 0.1.

Exercice 33. Consider a Strauss process in $W = [0, 1]^2$ with the following parameters $\beta = 100, \gamma = 0.25$ and r = 0.005

- a) Simulate 1000 samples using a Metropolis-Hastings dynamics and plot the time series evolution and diagnostics corresponding to the sufficient statistics.
- b) Simulate 1000 samples using the CFTP dynamics and plot the time series evolution and diagnostics corresponding to the sufficient statistics.
- c) Compare the sufficient statistics samples using exploratory analysis tools (boxplots, qqplots, histograms) and also statistical tests (Kolmogorov-Smirnoff).
- d) How many samples do we need in order to obtain Metropolis-Hastings based patterns that may be considered "good enough" ?

Table of contents

Cours 9. Parameter estimation. Model validation.

Statistical inference problems Parameter estimation based on pseudo-likelihood Monte Carlo Maximum likelihood estimation Parameter estimation using the ABC Shadow algorithm Model validation : residual analysis for point processes Exercises : Parameter estimation. Model validation.

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Parameter estimation and model validation : the statistical inference problems

Problem I : parameter estimation

- observe the pattern x and find the model parameters θ able to statistically reproduce it
- complete and incomplete data : pseudo-likelihood, Monte Carlo maximum likelihood, EM, ABC Shadow
- open problem : sampling $p(\theta|\mathbf{x})$

Problem II : pattern detection

- observe the data d and find x "hidden"
- the model parameters are : hidden, modelled, unknown
- open problem : the does the detected pattern really exist ?

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Problem III : sample the joint law of the pattern and the parameters $p(x, \theta)$

- ▶ shape modelling \rightarrow "crystal ball" ?
- look at a phenomenon and propose a model providing statistical prediction and validation of the observed pattern
- needs also the time dimension
- open problem : how to introduce time
- ► what it is the time "quanta" → generating element and interactions as for marked point processes → a twisted spaghetti - see pasta geometry of Max Tegmark ?

Parameter estimation based on pseudo-likelihood

The pseudo-likelihood of a marked point process X with conditional intensity $\lambda_{\theta}(\zeta; x)$ observed on the bounded set W is expressed as

$$PL_{W}(\theta; \mathbf{x}) = \left[\prod_{x_{i} \in \mathbf{x}} \lambda_{\theta}(x_{i}; \mathbf{x})\right] \exp\left[-\int_{W \times M} \lambda_{\theta}((w, m); \mathbf{x})\nu(dw)\nu_{M}(dm)\right].$$

 \rightarrow blackboard : construction of the pseudo-likelihood

The pseudo-likelihood estimator is given by the solution of the equation :

$$\frac{\partial PL_W(\theta; \mathsf{x})}{\partial \theta} = \mathsf{0}$$

Remarks :

- the PL is concave for exponential models
- no normalising constant needed ...
- it "amplifies" the interaction weights : check the formula for a Strauss process the interactions are counted twice ...
- consistency and asymptotic normality of the estimator : if we observe the model in a finite window, then it converges towards the parameters estimated using the pseudo-vraisemblance based on the observation of the "whole" window (Jensen and Møller, 1991)

- it can be used to have a "good" initial condition for other more elaborate methods
 - "good" results for mild interactions : (Mateu and Montes, 2001)
- lacks of statistical significance : there is no real link with the true model behind the pattern
 - except for the Poisson process : in this case the pseudo-likelihood is the true likelihood
- easy to be implemented :
 - this was the motivation to introduce it in the middle of 70s (Besag, 1975)
 - see (Baddeley, Rubak and Turner, 2016) for implementation details within the spatstat package

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 \rightarrow Exercise 34 + Exercice 35

Implementation within R spatstat package

stationary Strauss process :

$$\log \lambda_{\theta}(\zeta; \mathsf{x}) = \log \beta + (\log \gamma) t(\zeta, \mathsf{x})$$

with $t(\zeta, \mathbf{x})$ the number of pairs of objects closer than the distance r in the configuration \mathbf{y}

general structure of the conditional intensity

$$\log \lambda_{\theta}(\zeta; \mathsf{x}) = \eta S(\zeta) + \phi V(\zeta, \mathsf{x}),$$

with the 'first order term' S(u) that describes spatial inhomogeneity and/or covariates effects and the 'higher order' term that describes interobject interaction

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refer to the **spatstat** documentation

 \rightarrow caution : definition of the model \ldots

Applications

Pseudo-likelihood profile analysis : the range parameters
>radius = data.frame(r=seq(0.05,0.11, by=0.01))
>pradius = profilepl(radius, Strauss, japanesepines)
>plot(pradius,main="Strauss : PL analysis")



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Fitting the model to the pattern :

> ppm(japanesepines, 1,Strauss(r=0.08),rbord=0.08)
Stationary Strauss process
First order term: beta 77.93567
Interaction: Strauss process interaction distance:
0.08
Fitted interaction parameter gamma: 0.7953

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Monte Carlo Maximum likelihood estimation

Exponential family models :

- very general framework
- the point processes models that were presented are given by

$$p(\mathbf{x}|\theta) = \frac{h(\mathbf{x}|\theta)}{Z(\theta)} = \frac{\exp\langle t(\mathbf{x}), \theta\rangle}{Z(\theta)}$$

where $h(x|\theta)$, t(x) and θ represent the un-normalized probability density w.r.t. the standard Poisson process, the sufficient statistics vector and the model parameters vector, respectively. The normalising constant $Z(\theta)$ is unknown.

 \rightarrow blackboard : general notions about the exponential family models

The configuration x is entirely observed, hence the log-likelihood with respect a known parameter ψ can be written as follows :

$$I(heta) = \langle t(\mathsf{x}), heta - \psi
angle - \log rac{Z(heta)}{Z(\psi)}$$

It is easy to check, that the normalizing constants ratio is

$$\frac{Z(\theta)}{Z(\psi)} = \mathbb{E}\left[\exp\langle t(\mathbb{X}), \theta - \psi\rangle\right],$$

since we have

$$\begin{aligned} \frac{Z(\theta)}{Z(\psi)} &= \frac{1}{Z(\psi)} \int_{\Omega} h(\mathbf{x}|\theta) d\mu(\mathbf{x}) \\ &= \frac{1}{Z(\psi)} \int_{\Omega} h(\mathbf{x}|\theta) \frac{h(\mathbf{x}|\psi)}{h(\mathbf{x}|\psi)} d\mu(\mathbf{x}) \\ &= \int_{\Omega} \frac{h(\mathbf{x}|\theta)}{h(\mathbf{x}|\psi)} \frac{h(\mathbf{x}|\psi)}{Z(\psi)} d\mu(\mathbf{x}) \\ &= \mathbb{E}\left[\frac{h(\mathbb{X}|\theta)}{h(\mathbb{X}|\psi)}\right] \end{aligned}$$

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The Monte Carlo approximation of the normalizing constants ratio gives :

$$\frac{Z(\theta)}{Z(\psi)} \approx \frac{1}{n} \sum_{i=1}^{n} \exp\langle t(\mathbb{X}_i), \theta - \psi \rangle,$$

where X_1, X_2, \ldots, X_n are samples obtained from $p(y|\psi)$. Hence, the Monte-Carlo counterpart of the log-likelihood is :

$$I_n(\theta) = \langle t(\mathbf{x}), \theta - \psi \rangle - \log \left(\frac{1}{n} \sum_{i=1}^n \exp \langle t(\mathbb{X}_i), \theta - \psi \rangle \right).$$

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Theorem

The log-likelihood of an exponential family model is a convex function.

- proof : see (Monfort 1997, Thm.3, pp. 61)
- $I_n(\theta) \rightarrow I(\theta)$ almost sureley
- ▶ all these suggest that local optimisation procedures applied to $I_n(\theta)$ may give interesting results

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\rightarrow Exercise 36

MCMC local optimisation procedures

The gradient of the MCMC log-likelihood is

$$\nabla I_n(\theta) = t(\mathsf{x}) - \mathbb{E}_{n,\theta,\psi}[t(\mathbb{X})]$$

where

$$\mathbb{E}_{n,\theta,\psi}[t(\mathbb{X})] = \frac{\sum_{i=1}^{n} t(\mathbb{X}_i) \exp\langle t(\mathbb{X}_i), \theta - \psi \rangle}{\sum_{i=1}^{n} \exp\langle t(\mathbb{X}_i), \theta - \psi \rangle}$$

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that is the Monte Carlo importance sampling approximation of $\mathbb{E}_{\theta} t(\mathbb{X})$.

Similarly, the Hessian can be computed too :

$$-\nabla^2 I_n(\theta) = \mathbb{V}ar_{n,\theta,\psi}[t(\mathbb{X})]$$

where

$$\mathbb{V}ar_{n,\theta,\psi}[t(\mathbb{X})] = \mathbb{E}_{n,\theta,\psi}[t(\mathbb{X})t(\mathbb{X})^t] - \mathbb{E}_{n,\theta,\psi}[t(\mathbb{X})]\mathbb{E}_{n,\theta,\psi}[t(\mathbb{X})^t].$$

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Newton-Raphson method :

$$\theta_{k+1} = \theta_k - [\nabla^2 I_n(\theta_k)]^{-1} \nabla I_n(\theta_k)$$
(17)

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for k = 1, 2, ...,

- ▶ $I_n(\cdot)$ is computed using *n* samples from $p(x|\psi)$
- the computation of the gradient and Hessian inverse is numerically unstable
- useful only if the initial value is close enough from the solution

Iterative gradient method :

$$\begin{cases} I_n(\theta_k + \rho(\theta_k) \nabla I_n(\theta_k)) = \max_{\rho \in \mathbb{R}} I_n(\theta_k + \rho \nabla I_n(\theta_k)) \\ \theta_{k+1} = \theta_k + \rho(\theta_k) \nabla I_n(\theta_k) \end{cases}$$

where $\rho(\theta_k)$ is the optimal step (Descombes et al. '99, Stoica '01).

- ▶ re-sampling if $\parallel \theta_k \psi \parallel >$ threshold
- \blacktriangleright obtain a reference value θ_0 close enough to the maximum likelihood estimator

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Stochastic gradient :

$$\theta_{k+1} = \theta_k + \epsilon_k [t(\mathbf{x}) - t(\mathbf{X}_k)]$$

where $\epsilon_k > 0$ is a decreasing sequence while \mathbb{X}_k is a sample of $p(\mathbf{x}|\theta_k)$

very simple, but finding an optimal sequence {\epsilon_k} is an open problem

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- L. Younes, G. Winkler : Markov random fields
- R. Moyeed and A. Baddeley : point processes

Asymptotic results MCMCML estimation

The random variable $\sqrt{n}(\hat{\theta}_n - \hat{\theta})$ whenever $n \to \infty$, it converges in distribution towards a normal random variable of zero mean and variance $I(\hat{\theta})^{-1} \Gamma I(\hat{\theta})^{-1}$:

$$\sqrt{n}(\widehat{\theta}_n - \widehat{\theta}) \to \mathcal{N}(0, I(\widehat{\theta})^{-1} \Gamma I(\widehat{\theta})^{-1}).$$

the matrix

$$I(\widehat{\theta}) = \mathbb{V}ar_{\widehat{\theta}}[t(\mathbb{X})] = -\nabla^2 I(\widehat{\theta})$$

is the Fisher information of $\widehat{\theta}$

It he matrix Γ is the matrix of the asymptotic covariance of the normalised Monte Carlo gradient √n∇l_n(θ̂)

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- ▶ the variance of the components of $\hat{\theta} \theta_0$ can be estimated by taking the diagonal elements of the inverse of $-\nabla^2 I_n(\hat{\theta}_n)$
 - it represents the error between the maximum likelihood estimate and the true model parameters

► the variance of the components of $\sqrt{n}(\hat{\theta}_n - \hat{\theta})$ can be estimated by taking the diagonal elements of $I(\hat{\theta})^{-1}\Gamma I(\hat{\theta})^{-1}$

 it represents the error between the maximum likelihood estimate and its Monte Carlo counterpart

 refer to (Monfort, 1997), (Geyer, 1999) and (van Lieshout and Stoica, 2003) for the computation of these matrices

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 \rightarrow show the next example + blackboard : incomplete data observation - EM algorithm

MCML example

Candy model : (van Lieshout and Stoica, 2003)



Figure: Realization (left) of the reference model given by the parameters in the middle table. The observed values of the sufficient statistics are listed at right.

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Results : estimation of the parameters from the reference configuration given by the Candy model

Initial param-	Iterative	Monte Carlo
eters	method	MLE
$\theta_f^i = -9.5$	$\hat{\theta}_{f}^{0} = -8.37$	$\hat{\theta}_f^n = -8.32$
$\theta_s^i = -4.0$	$\hat{ heta}_{s}^{0} = -2.74$	$\hat{\theta}_s^n = -2.73$
$\theta^i_d = 1.5$	$\hat{\theta}_{d}^{0} = 2.46$	$\hat{\theta}_d^n = 2.47$
$\theta_o^i = -3.5$	$\hat{ heta}_o^0 = -2.13$	$\hat{\theta}_o^n = -2.17$
$\theta_r^i = -3.5$	$\hat{\theta}_{r}^{0} = -2.42$	$\hat{\theta}_r^n = -2.42$

Asymptotics : estimation errors (central limit theorems available)

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Asymptotic standard	MCSE
deviation of MLE	
0.51	0.002
0.23	0.003
0.17	0.001
0.30	0.002
0.31	0.005

Log-likelihood ratio approximation :



Figure: Monte Carlo approximation of the log likelihood function. The X axis represents the variation of a single component. The Y axis represents the values of the Monte Carlo log likelihood with all other components of $\hat{\theta}^0$ fixed : a - $\theta_f \in [-11, -7]$, b - $\theta_s \in [-5, -1]$, c - $\theta_d \in [1, 5]$, d - $\theta_o \in [-4.5, -0.5]$, e - $\theta_r \in [-4.5, -0.5]$.

Parameter estimation using the ABC Shadow algorithm

Parameter estimation : the pattern detection dual problem

let p(θ|y) be the conditional distribution of the model parameters given the patten

$$p(\theta|\mathbf{y}) = \frac{\exp[-U(\mathbf{y}|\theta)]p(\theta)}{Z(\mathbf{y})c(\theta)}$$

with $p(\theta)$ the prior density for the model parameters and Z(y) the normalization constant

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► the posterior law is defined on Θ a compact region in ℝ^m; the parameter space Θ is endowed with the Borel algebra T_Θ the parameter estimator is

$$\widehat{ heta} = rg\max_{ heta \in \Theta} \{ p(heta | \mathsf{y}) \}$$

- optimisation procedure : requires sampling $p(\theta|y)$
- Sampling the posterior law is not straightforward → requires the evaluation of the ratio c(θ)/c(ψ)
- if p(θ) is the uniform distribution over Θ then θ is the maximum likelihood estimator

Remark : the model parameters taken into account by the posterior distribution are the "interaction" parameters. And not the "range" parameters ...

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Sampling posterior laws : theoretical solution

Auxiliary variable method : given by (Møller, Pettitt, Reeves and Berthelsen, 2006)

▶ principle : use an auxiliary variable X defined by the probability density a(x|θ, y) → sample the joint distribution

$$\pi(\theta, \mathsf{x}|\mathsf{y}) = \mathsf{a}(\mathsf{x}|\theta, \mathsf{y})\mathsf{p}(\theta|\mathsf{y}) = \mathsf{a}(\mathsf{x}|\theta, \mathsf{y})\frac{\exp[-U(\mathsf{y}|\theta)\mathsf{p}(\theta)}{Z(\mathsf{y})\mathsf{c}(\theta)}$$

present context : the auxiliary variable → auxiliary pattern ...
 key idea : the final mathematical construction leads to the simplification of the normalizing constants ratio in the MH algorithm sampling π(θ, x|y)

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\rightarrow blackboard: explain the algorithm

Remarks :

- very elegant solution
- exact sampling of the auxiliary variable
- choice of the auxiliary variable density : behaviour of the simulated chain ...

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Sampling posterior laws : approximate solution

Approximate Bayesian Computation

- generic name for numerical simulation methods allowing model selection based on an approximate sampling from the posterior distribution p(θ|y)
- \blacktriangleright idea originated in Montpellier \rightarrow statistics and environmental sciences communities

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- (Grelaud, Robert, Marin, Rodolphe and Taly, 2009)
- (Marin, Pudlo, Robert and Ryder, 2012)

Algorithm ABC : assume the observed pattern is y, fix a tolerance threshold ϵ and an integer value *n*.

- 1. For i = 1 to n do
 - Generate θ_i according to $p(\theta)$.
 - Generate x_i according to the probability density $p(x|\theta_i) = \frac{\exp[-U(x|\theta_i)]}{c(\theta_i)}$
- 2. Return all the θ_i 's such that the distance between the statistics of the observation and those of the simulated pattern is small, that is

$$d(t(y), t(x_i)) \leq \epsilon$$

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Figure: Graphical representation of the outputs of an ABC algorithm.

Theoretical result

- (Blum, 2009) : gives the bias and the variance of the posterior distribution estimate
- (G. Biau, F. Cérous and A. Guyader , 2015) : give asymptotic features of the outputs of a slightly different algorithm
- ideas : kernel and k-nearest neighbour estimation

Remarks :

- exact sampling from $p(x|\theta)$ is needed
- choice of the statistics vector
 - exponential family models \rightarrow the sufficient statistics
- appropriate setting :
 - distance d
 - precision parameter ϵ
 - number of neighbours k_n and bandwidth parameter h_n

Synthesis :

 ABC algorithms are useful if enough samples x_i are "close" to the observed pattern y

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ABC Shadow algorithm

Key points

- need : an algorithm with outputs "close" enough to the posterior distribution
- tool : build a Markov chain evolving "close" to an equilibrium regime given by p(θ|y)
- plan : use the auxiliary variable method ideas

Ideal MCMC sampling of the posterior : general MH algorithm

- \blacktriangleright assume the system is in the state θ
- choose a new value ψ according to a proposal density $q(\theta \rightarrow \psi)$
- \blacktriangleright the value ψ is accepted with probability

$$\alpha_i(\theta \to \psi) = \min\left\{1, \frac{p(\psi|\mathbf{y})p(\psi)}{p(\theta|\mathbf{y})p(\theta)} \frac{q(\psi \to \theta)}{q(\theta \to \psi)}\right\}$$

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consider the proposal density

$$q(\theta \to \psi) = q_{\Delta}(\theta \to \psi | \mathsf{x}) = \frac{f(\mathsf{x} | \psi) / c(\psi)}{I(\theta, \Delta, \mathsf{x})} \mathbb{1}_{b(\theta, \Delta/2)} \{\psi\}$$

with

- x : outcome of a marked point process driven by the probability density p(x|v) where v is any value in Θ.
- ▶ pattern detection context : $f(x|\psi) = exp[-U(x|\psi)]$
- $\Delta > 0$: control parameter
- 1_{b(θ,Δ/2)}{·} is the indicator function over b(θ, Δ/2), which is the ball of centre θ and radius Δ/2

$$I(\theta, \Delta, \mathbf{x}) = \int_{b(\theta, \Delta/2)} f(\mathbf{x}|\phi) / c(\phi) \, d\phi.$$

- this choice guarantees the ideal chain to be uniformly ergodic and avoids the evaluation of the ratios c(θ)/c(ψ)
- ▶ but, it requires the computation of integrals $I(\theta, \Delta, x)$...
- \rightarrow blackboard : drawing

Shadow chain : approximation of the ideal chain

Theorem : if $p(\mathbf{x}|\theta)$ is a continuously differentiable function in θ For any fixed $\theta \in \Theta$ and $A \in \mathcal{T}_{\Theta}$, we have

$$\lim_{\Delta\to 0_+}\int_{\mathcal{A}}|q_{\Delta}(\theta\to\psi)-U_{\Delta}(\theta\to\psi)|d\psi=0$$

For any fixed $\theta \in \Theta$, we have

$$\lim_{\Delta \to 0_+} \sup_{\psi \in \Theta} \left| \frac{q_{\Delta}(\theta \to \psi | \mathsf{x})}{q_{\Delta}(\psi \to \theta | \mathsf{x})} - \frac{\frac{f(\mathsf{x}|\psi)}{c(\psi)} \mathbf{1}_{b(\theta, \Delta/2)}(\psi)}{\frac{f(\mathsf{x}|\theta)}{c(\theta)} \mathbf{1}_{b(\psi, \Delta/2)}(\theta)} \right| = 0$$

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uniformly in $\theta \in \Theta$, with

Application : simulate the shadow chain that approximate the ideal chain

- First part : use U_Δ(θ → ψ) instead of q_Δ(θ → ψ) for proposing new values
- second part : approximates the computation of the proposal density ratio while simplifying the normalizing constant ratio
- the shadow Markov chain accepts new states with the probability :

$$\alpha_{s}(\theta \to \psi) = \min\left\{1, \frac{p(\psi|\mathsf{y})p(\psi)}{p(\theta|\mathsf{y})p(\theta)} \times \frac{f(\mathsf{x};\theta)c(\psi)\mathbf{1}_{b(\psi,\Delta/2)}\{\theta\}}{f(\mathsf{x};\psi)c(\theta)\mathbf{1}_{b(\theta,\Delta/2)}\{\psi\}}\right\}$$

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Corollary : the acceptance probabilities of the ideal and shadow chains are uniformly as closed as desired whenever $\triangle \rightarrow 0_+$

Proposition : Let P_i and P_s be the transition kernels for the ideal and the shadow Markov chains using a general $\Delta > 0$ and a configuration $x \in \Omega$. Then, for every $\epsilon > 0$ and every $n \in \mathbb{N}$, there exists $\Delta_0 = \Delta_0(\epsilon, n) > 0$ such that for every $\Delta \leq \Delta_0$

$$|P_i^{(n)}(\theta,A) - P_s^{(n)}(\theta,A)| < \epsilon$$

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uniformly in $\theta \in \Theta$ and $A \in \mathcal{T}_{\Theta}$.

Algorithm ABC Shadow : assume the observed pattern is y and fix values for Δ and *n* and the current state θ_0

- 1. Generate x according to $p(x|\theta_0)$
- 2. For k = 1 to n do
 - Generate a new candidate ψ following $U_{\Delta}(\theta_{k-1} \rightarrow \psi)$.

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► The new state $\theta_k = \psi$ is accepted with probability $\alpha_s(\theta_{k-1} \to \psi)$, otherwise $\theta_k = \theta_{k-1}$

3. Return θ_n

Remarks :

- if several samples are needed, re-start the procedure for the same Δ and n, with $\theta_0 = \theta_n$.
- depending on Δ, the algorithm approaches the equilibrium regime of the ideal chain :

$$\|\mathcal{P}_{s}^{(n)}(\theta, A) - \pi(A)\| \leq M(\mathsf{x}, \Delta)\rho^{n} + \epsilon.$$

with $\pi(A) = \int_A p(\theta|\mathbf{y}) d\theta$; *M* and ρ : ergodicity parameters of the ideal chain

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caution : this is not convergence

Application : sampling the posterior of a Gaussian model

The posterior of a Normal model with mean θ_1 and variance θ_2 is

$$p(heta_1, heta_2|\mathsf{y}=\mathbb{Y}(\omega))\propto rac{\exp\left(rac{ heta_1}{ heta_2}\mathbb{Y}(\omega)-rac{\mathbb{Y}^2(\omega)}{2 heta_2}
ight)}{c(heta_1, heta_2)}p(heta_1, heta_2)$$

with

 y = 𝔅(ω) : observation issued from the supposed model
 t(y) = (𝔅(ω), 𝔅²(ω)) : the sufficient statistics vector
 if the sample size is *m* then : t(y) = (∑_{i=1}^m 𝔅_i(ω), ∑_{i=1}^m 𝔅²_i(ω))

Experiment :

- Simulate 1000 i.i.d. Normal r. v.'s with parameters $\theta = (\mu, \sigma^2) = (2, 9)$
- t(y) = (1765.45, 12145.83)
- ▶ $p(\theta_1, \theta_2)$ the uniform distribution over $[-100, 100] \times [0, 200]$
- compare 1000 samples of the MH and ABC Shadow algorithms
- $\Delta = (0.005, 0.025)$ and n = 500

Summary statistics for Normal posterior sampling						
Algorithm	Q_5	Q ₂₅	Q_{50}	$\bar{ heta}$	Q ₇₅	Q_{95}
$MH \ \theta_1$	1.60	1.69	1.75	1.76	1.82	1.92
ABC θ_1	1.60	1.70	1.76	1.76	1.82	1.91
MH θ_2	8.45	8.80	9.07	9.08	9.33	9.76
ABC θ_2	8.35	8.78	9.03	9.06	9.33	9.83

Table: Empirical quantiles and mean for the posterior of the Normal model.



Figure: Boxplots and qqplots of the MH and ABC Shadow outpts.



Figure: Sample path for the Normal posterior. Left colum: the MH algorithm results - from top to bottom the joint parameter path and the θ_1 time series. Right column: the ABC Shadow procedure - from top to bottom the joint parameter path and the θ_1 time series.

Parameter estimation for marked point processes

Strauss model :

$$p(\mathbf{y}|\theta) \propto \beta^{n(\mathbf{y})} \gamma^{s_r(\mathbf{y})},$$
 (18)

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with

t(y) = (n(y), s_r(y)) : sufficient statistics
 θ = (log β, log γ) : model parameters

Experiment :

- domain $W = [0, 1]^2$ and range parameter r = 0.1
- ▶ simulate 1000 realisations of the model $\theta = (4.60, -1.60)$ using the CFTP algorithm
- ► the empirical means of the sufficient statistics $\overline{t}(y) = (\overline{n}(y), \overline{s_r}(y)) = (34.33, 5.31)$
- ▶ ABC Shadow "sampling" of $p(\theta|\bar{t}(y)) \rightarrow 1000$ samples

•
$$\Delta = (0.01, 0.01)$$
 and $n = 200$

▶ $p(\theta)$ the uniform distribution over $[3.5, 5.5] \times [-5, 0]$

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Figure: Strauss posterior sampling using the ABC procedure : histograms and kernel density estimates of the marginals. Maximum of the kernel estimated density is $\hat{\theta} = (4.63, -1.53)$.

Summary statistics for Strauss posterior sampling				
Algorithm	Q_{50}	$ar{ heta}$		
ABC $\log \beta$	4.606	4.603		
ABC $\log \gamma$	-1.669	-1.700		

Table: Empirical median and mean for the posterior of the Strauss model.

Candy model :

$$p(\mathbf{y}|\theta) \propto \exp\langle\theta_d n_d(\mathbf{y}) + \theta_s n_s(\mathbf{y}) + \theta_f n_f(\mathbf{y}) + \theta_r n_r(\mathbf{y})\rangle$$
(19) with

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Experiment :

- domain W = [0,3] × [0,1], segment length I = 0.12, connection range r_c = 0.01, curvature parameters τ_c = τ_r = 0.5 radians
- simulate 1000 realisations of the model θ = (10,7,3,-1) using an Adapted MH algorithm
- the empirical means of the sufficient statistics $\overline{t}(y) = (51.10, 101.06, 19.97, 72.89)$
- ▶ ABC Shadow "sampling" of $p(\theta|\bar{t}(y)) \rightarrow 1000$ samples

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- $\Delta = (0.01, 0.01, 0.01, 0.01)$ and n = 500
- $p(\theta)$ the uniform distribution over $[2, 12]^3 \times [-7, 0]$
| Summary statistics for Candy posterior sampling | | | | | |
|-------------------------------------------------|----------|-------------|--|--|--|
| Algorithm | Q_{50} | $ar{	heta}$ | | | |
| ABC $\log \theta_d$ | 9.995 | 9.998 | | | |
| ABC $\log \theta_f$ | 2.977 | 2.975 | | | |
| ABC $\log \theta_s$ | 7.005 | 7.008 | | | |
| ABC $\log \theta_r$ | -1.014 | - 1.016 | | | |

Table: Empirical median and mean for the posterior of the Candy model. Maximum of the kernel estimated density is $\hat{\theta} = (9.96, 3.02, 6.99, -1.00)$.

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Real data application : cosmology

short range correlations for the galaxy distribution



Figure: Sample of a cosmological data set. The points represent galaxy positions in a region of our Universe.

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► exploratory analysis : summary statistics → cluster behaviour



Figure: Summary statistics envelope test (100 simulations) for the galaxy pattern: shaded (gray) region - the Monte Carlo envelopes, dotted line (red) - the theoretical statistics, continuous (black) line - the observed statistics.

- sampling the posterior of an area-interaction model conditionally on different range parameters
- ► the posterior sampling → the probability that the model parameters indicate clustering

Data for the Galaxy pattern								
r	0.01	0.02	0.03	0.04	0.05	0.06	0.07	
n(y) = 163								
$-a_r(y)$	135.91	114.05	96.44	82.23	69.85	59.05	49.73	

Table: The observed sufficient statistics computed for the galaxy pattern, depending on the range parameter r. For all these parameters n(y) remains constant, while $a_r(y)$ depends on r.

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▶ the area interaction model : $p(y|\theta) \propto \beta^{n(y)} \gamma^{a_r(y)}$ with

$$a_{r}(y) = -\frac{\nu \left[A_{r}(y)\right]}{\pi r^{2}} = -\frac{\nu \left[\bigcup_{i=1}^{n} b(w_{i}, r)\right]}{\pi r^{2}}$$



Figure: Box-plots of the posterior distributions for the parameters of the area-interaction process estimated from the galaxy pattern, given different values for the interaction radius.

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Estimation errors								
r	0.01	0.02	0.03	0.04	0.05	0.06	0.07	
Asymptotic standard deviation								
$\widehat{\sigma}_{\log\beta}$	0.20	0.17	0.13	0.11	0.10	0.10	0.08	
$\widehat{\sigma}_{\log \gamma}$	0.26	0.28	0.27	0.30	0.34	0.40	0.52	
Monte Carlo standard deviation								
$\widehat{\sigma}_{\log \beta}^{MC}$	0.001	0.002	0.001	0.002	0.002	0.002	0.003	
$\widehat{\sigma}_{\log \gamma}^{MC}$	0.002	0.003	0.004	0.006	0.008	0.012	0.024	

Table: Estimation errors computed for the MAP estimates obtained from the galaxy pattern.

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Synthesis parameter estimation

Pseudo-likelihood estimation :

- easy to compute
- good alternative whenever nothing else can be done
- consistency and central limit theorems : difficult to interpret
- no properties concerning the sufficient statistics of the model using the PL estimates of the parameters
- work of J. Mateu and P. Montes : comparison with maximum likelihood

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Monte Carlo maximum likelihood :

- general statistical framework
- \blacktriangleright numerically unstable \rightarrow but re-sampling is guaranteed to convergence, since the log-likelihood is convex
- the asymptotics are related to the true model
- property : the expectation of the sufficient statistics computed by the model with the ML parameters equals the observed sufficient statistics

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ABC Shadow parameter estimation :

approximate algorithm that samples "close" to the posterior

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- interesting numerical results : comparable with classical MCMCML inference
- relatively low computational cost : MCMCML needs re-sampling
- ► complementary tool → applications :
 - initialisation point for more rigorous methods
 - statistical testing
 - model validation
- open problem : range parameters

Open questions :

- range parameters
- parameters of the mark distribution
- posterior sampling
- ► incomplete data : EM algorithms converges towards the first local maximum → a lot of references available ...
- ► ABC methods : empirical methods for parameter estimation → control the sufficient statistics

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Model validation : residual analysis for point processes Let X be a locally stable marked point process on $W \times M$.

h-Innovations : for nonnegative functions h and $A \subseteq W \times M$

$$I(A,h,\lambda) = \sum_{x_i \in \mathbb{Y}_A} h(x_i,\mathbb{X}\setminus x_i) - \int_A \lambda(\eta;\mathbb{X})h(\eta,\mathbb{X})(
u imes
u_M)(d\eta)$$

assuming the sum and the integral in the definition have finite expectations, the Georgii-Nguyen-Zessin formula gives

$$\mathbb{E}I(A,h,\lambda)=0$$

- I is a signed measure
- ► $\triangle I(x_i) = h(x_i, \mathbb{X} \setminus \eta)$: the innovation increment ('error') attached to a point $\eta \in \mathbb{X}$
- ► $dI(\eta) = -\lambda(\eta; \mathbb{X})h(\eta, \mathbb{X})$: the innovation increment attached to a background location $\eta \in W \times M$

h-Residuals : for $h \ge 0$ functions and $A \subseteq W \times M$

$$\begin{split} R(A,\widehat{h},\widehat{\theta}) &= I(A,\widehat{h},\widehat{\lambda}) \\ &= \sum_{x_i \in \mathsf{x}_A} \widehat{h}(x_i,\mathsf{x} \setminus x_i) - \int_A \widehat{\lambda}(\eta;\mathsf{x}) \widehat{h}(\eta,\mathsf{x})(\nu \times \nu_M)(d\eta) \end{split}$$

since the function h may depend on the model, \hat{h} denotes an estimate.

Application idea :

- consider a parametric model for a marked point process X observed within A
- estimate the model parameters (maximum likelihood, pseudo-likelihood)
- expect the residuals R(A) to be close to 0 if the model is appropriate

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Building residuals : several possible choices for h

▶ raw residuals
$$h(\eta, x) = 1$$

$$R(A,1,\widehat{\theta}) = n(\mathsf{x} \cap A) - \int_{A} \widehat{\lambda}(\eta;\mathsf{x})(\nu \times \nu_{M})(d\eta)$$

 inverse residuals h(η, x) = 1/λ(η; x) (equivalent with the Stoyan-Grabarnik diagnostic)

$$R(A,\frac{1}{\widehat{\lambda}},\widehat{\theta}) = \sum_{x_i \in \mathsf{x}_A} \frac{1}{\widehat{\lambda}(x_i;\mathsf{x}_A)} - \int_A \mathbb{1}\{\widehat{\lambda}(\eta;\mathsf{x}) > 0\}(\nu \times \nu_M)(d\eta)$$

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▶ Pearson residuals $h(\eta, x) = 1/\sqrt{\lambda(\eta; x)}$ (analogy with Poisson log-linear regression)

$$R(A, \frac{1}{\sqrt{\widehat{\lambda}}}, \widehat{\theta}) = \sum_{x_i \in x_A} \frac{1}{\sqrt{\widehat{\lambda}(x_i; x_A)}} - \int_A \sqrt{\widehat{\lambda}(\eta; x)} (\nu \times \nu_M) (d\eta)$$

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Remark : the inverse and Pearson residuals we need $\lambda_{\theta(x)}(x_i; x) > 0$ for all $x_i \in x$ for any pattern x, while $\lambda_{\theta(x)}(\eta; x) = 0$ is allowed for $\eta \notin x$

Properties :

expectation

$$\begin{split} &\mathbb{E}\left[R(A,\widehat{h},\widehat{\theta})\right] \\ &= \int_{\mathcal{A}}\mathbb{E}\left[h_{\widehat{\theta}(\mathbb{X}\cup\{\eta\})}(\eta,\mathbb{X})\lambda(\eta,\mathbb{X}) - h_{\widehat{\theta}(\mathbb{X})}(\eta,\mathbb{X})\lambda_{\widehat{\theta}(\mathbb{X})}(\eta,\mathbb{X})\right] \end{split}$$

- variance : more complicate structures but very nice formulas for Poisson processes (Baddeley, Moller and Pakes 2008)
- ► these residuals do not have independent increments → the raw innovations for Markov point processes are conditionnaly independent and uncorrelated (Baddeley, 2005)
- consistency and asymptotic normality for the residuals of stationary Gibbs point processes (Coeurjolly and Lavancier, 2013)

Application : smoothed residuals to test several models for japanesepines datasets

- Strauss process : only repulsion
- area-interaction process : repulsion or attraction (competition for ressources)



Figure: Raw residual analysis, from left to right : $Strauss(r{=}0.08)$ and $AreaInt(r{=}0.09)$

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R code:

>mjp=ppm(japanesepines, 1,Strauss(r=0.08),rbord=0.08)
>rjp=residuals(mjp,type="raw") >plot(rjp)

QQ plots: comparison of empirical quantiles of the smoothed residuals with the expected quantiles under the estimated model

- interpretation in the spirit of K and F functions
- if the data pattern is more clustered than the model: heavier tails especially in the left-hand tail
- if the data pattern is more inhibited than the model: lighter tails especially in the right-hand tail

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R code: qqplot.ppm(rjp, type="raw")



Figure: Q-Q plot analysis, from left to right : Strauss(r=0.08) and AreaInt(r=0.09)

- Strauss(r=0.08): over-estimates repulsion, but under-estimates close attraction
- AreaInter(r=0.09): very well for the close attraction, underestimate the repulsion
- the best model for the entire data set : polynomial inhomogeneity and soft-core interaction

Remarks :

- the theory is wonderful
- but the numerical results are obtained using the PL estimators
- see the remark of J. Besag
- visualisation of residuals difficult for higher dimensions
- ► the qq plots very informative → link with the central limit theorems for computing confidence intervals

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open question : validating pattern detection result ... ?

\rightarrow Exercise 37

Exercise 34. Let X be a homogeneous Poisson point process on a compact $W \subset \mathbb{R}^d$ with unknown intensity parameter θ .

a) Write its probability density w.r.t. the reference given by the unit intensity Poisson process. Specifiy its normalising constant.

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b) Write the log-likelihood function and give the maximum likelihood estimate of θ .

Exercice 35. Let X be an inhomogeneous Poisson point process on a compact $W \subset \mathbb{R}^d$ with the intensity function λ depending on the parameter θ .

- a) Write its probability density w.r.t. the reference given by the unit intensity Poisson process. Specifiy its normalising constant.
- b) Write the log-likelihood function and compare it with the log pseudo-likelihood of a point process.
- c) In this case, what is the difference between the intensity of the process and its conditional intensity ?

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Exercice 36. Let X be a Strauss point process on a compact $W \subset \mathbb{R}^d$ given by the probability density w.r.t. the unit rate Poisson point procees :

$$p(\mathsf{x};\theta) = \alpha(\theta)\beta^{n(\mathsf{x})}\gamma^{s(\mathsf{x})}$$

where n(x) denotes the cardinality of x, s(x) the number of r-close pairs in x and $\theta = (\beta, \gamma)$ with $\beta > 0$ and $\gamma \in (0, 1)$ is the parameter vector of interest

- a) Specify the normalising constant and write the log-likelihood function.
- b) Write the system of equations for obtaining θ .
- c) Show that the maximum likelihood estimator solves :

$$(n(x), s(x)) = (\mathbb{E}_{\theta} n(X), \mathbb{E}_{\theta} s(X))$$

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Exercice 37. The data set waterstriders within the package **spatstat** contains three point patterns representing the positions of water striders in a pound. Try to answer the following question : what is the more important factor in the relative position of these insects - the distance w.r.t to its neighbour or the occupied territory ? In order to answer the question, do the following steps :

- a) Plot the data. Do a summary statistics analysis in order to propose at least two candidates for modelling ? What type of interaction is exhibited by all these patterns ?
- b) Use the profile of the pseudo-likelihood and the previous analysis in order to propose a good candidate for the interaction radius.
- c) Fit the models using ppm.
- d) Verify the models using envelope tests.
- e) Choose a model using the residual analysis : residuals, qqplot

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Table of contents

Cours 10. Pattern detection and characterisation.

Statistical pattern detection Does the detected pattern really exist ? Exercises : Pattern detection and characterisation.

Pattern detection and characterisation

The pattern detection problem :

- \blacktriangleright a spatial data set is observed \rightarrow pattern hidden in the data ?
- hypothesis : the pattern is the realization of a random process (MRFs, marked point processes, etc.)
- \blacktriangleright the Gibbsian modelling framework \rightarrow write a probability density model

$$p(\mathbf{y}| heta) = rac{\exp[-U(\mathbf{x}| heta)]}{c(heta)}$$

with $U: \Omega \to \mathbb{R}^+$ the energy function, θ the model parameters and $c(\theta)$ the normalizing constant

- ▶ parameters knowledge \rightarrow prior density $p(\theta)$
- the pattern estimator is

$$(\widehat{\mathbf{x}},\widehat{\mathbf{\theta}}) = \arg \max_{(\mathbf{x},\mathbf{\theta})\in\Omega imes\Theta} \{p(\mathbf{x}|\mathbf{\theta})p(\mathbf{\theta})\} = \arg \max_{(\mathbf{x},\mathbf{\theta})\in\Omega imes\Theta} \{p(\mathbf{x},\mathbf{\theta})\}.$$

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Statistical pattern detection

Build the pattern model : probability density construction conditionally on the data observation

$$p(\mathbf{x}, \theta | \mathbf{d}) \propto \exp \left[-\frac{U_{\mathbf{d}}(\mathbf{x}|\theta) + U_{i}(\mathbf{x}|\theta)}{Z(\theta)} + \log p(\theta) \right]$$

- interaction energy U_i(x|θ) → objects interactions (geometrical shape of the structure)
- ► data energy $U_d(\mathbf{x}|\theta)$ induced by the data field d \rightarrow object locations
- if the interaction parameters are unknown \rightarrow prior model $p(\theta)$

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role of the interaction and data energies :



Figure: Influence of the energy components of the model : a) original image SPOT ; and results obtained using only : the data term (b), the interaction term (c), the complete model (d)

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setting the model parameters :



Figure: Two segments configurations : a) the connectivity is favored over alignment, b) connectivity and alignment have equivalent potentials.

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Pattern estimator : the object configuration that maximises the probability density

$$(\widehat{\mathbf{x}},\widehat{\theta}) = \arg\min_{\Omega imes \Psi} \left\{ \frac{U_{\mathsf{d}}(\mathbf{x}|\theta) + U_i(\mathbf{x}|\theta)}{Z(\theta)} - \log p(\theta) \right\}$$

with Ψ the model parameters space.

Simulated annealing : global optimisation technique

- ▶ sampling from $p(x, \theta)^{1/T}$ while slowly $T \rightarrow 0$
- convergence towards the uniform distribution on the configuration subspace minimizing U(x, θ) (Stoica, Gregori and Mateu, 2005)

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inhomogeneous Markov chain

Algorithm SA : x = Simulated Annealing (T_0, δ, T)

1. choose an initial condition x_0

2. for
$$i = 1$$
 to T do

$$\begin{cases} \\ \mathsf{x}_i = \mathsf{Update} (\mathsf{x}_{i-1}, \mathcal{T}_{i-1}, \delta) \\ \mathcal{T}_i = \mathcal{T}_0 / [\mathsf{log}(i) + 1] \\ \end{cases}$$

3. return x_T .

 \blacktriangleright slow algorithm \rightarrow an alternative cooling schedule :

$$T_{n+1} = cT_n$$
 with $c \in [0.95, 1[$

simulated tempering : improving mixing properties

Level sets estimators :

visit maps for compact regions in W :

$$\{T(w) > \alpha\} \Rightarrow \{T_n(w) > \alpha\}$$

with $T(w) = \mathbb{P}(w \in X)$ the probability that the structure hits a point in W

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link with the capacity functional and volumic fraction

- two challenges : discretisation and Monte Carlo approximations
- Vorob'ev expectation : the level set with volume equal to the mean volume of the random set
 - demands the knowledge of the behaviour of an unknown random set, but still manageable in practice ...
- average behaviour of the pattern (fixed temperature)
- ▶ (Heinrich, Stoica and Tran, 2012) prove the convergence L¹ of these estimators

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\rightarrow Exercise 38

Build the machine ...

Filaments detection in galaxies catalogues :

- interaction energy : Bisous model (random cylinders)
- data energy : local tests (density and spread of galaxies inside a cylinder)



Figure: Locating interacting cylinders in a field of points.

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Cluster detection in galaxies catalogues :

- interaction energy : Strauss and Area-interaction models (random object : trunck of a cone + two hal-spheres)
- data energy : local minimum number of galaxies insid the considered random object tests



Figure: Cross-section of the considered random object : two half-spheres connected with a truncated cone. The object is fully determined by its centre position, radius r and shape parameter $t \ge 1$. Shape parameter t gives the aspect ratio of the object along and perpendicular to the line of sight; for t = 1 the object is a ball. For a given r and t the height of the truncated cone is defined as h = 2r(t - 1). The shape of the truncated cone is defined by the lines of sights, which are indicated by dashed lines on the figure. The observer is located at far left from the object.

Cluster detection in epidemiological data :

- interaction energy : Strauss and Area-interaction models (random disks)
- data energy : local statistical test (the average score of the farms covered by a disk)



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Figure: Data \rightarrow field of marked points : a) observed clusters, b) clusters approximated by random disks.

Orbit determination for binary systems (1)

interaction energy :

$$U_l(\theta) = \log p(\theta) = \sum_{i=1}^7 \log p(\theta_i),$$

where $\theta = (a, e, i, \Omega, \omega, \tau, P)$ is the vector of orbital parameters

Jeffreys' principle - non-informative independent priors

- our choice : uniform distributions over bounded intervals
- the intervals were chosen taking into account the a priori knowledge of the objects to be detected

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perspective : introduce dependence of the parameters
Orbit determination for binary systems (2)

data energy : sum of the distances between the observed positions and the computed positions ; these last ones are computed using the given model parameters

$$U_{\mathsf{d}}(\varphi|\theta) = \sum_{i=1}^{n} \left[-\left(|x_i^{\mathsf{o}} - x_i^{\mathsf{c}}|^{l} + |y_i^{\mathsf{o}} - y_i^{\mathsf{c}}|^{l} \right)^{k/l} \right] =$$

where

- d = {(x_i^o, y_i^o)}, (i = 1, 2, ..., n) : the n observed positions of the secondary asteroid with respect to the primary
- {(x_i^c, y_i^c)}: the computed positions at the same time *i* as the corresponding observations, given the current θ
- ▶ *k*, *l* : pre-fixed model parameters
 - k = l = 2: Gaussian character of the data model
 - k = l = 1: Laplacian character of the data model

perspective : model choice

Road network extraction in satellite and aerial images

(Stoica, Descombes, van Lieshout and Zerubia, 2002)



Figure: Rural region in Malaysia : a) original image; b) obtained results.

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Forest galleries : verifying the results (Stoica, Descombes and Zerubia, 2004)



Figure: Forest galleries extraction : a) original image ; b) ground truth ; c)-d) obtained results. Data provided by BRGM.

Filaments detection (1) : (Tempel, Stoica et. al., 2014)



Figure: Detected filamentary pattern (cylinder axes) in a small sample volume within a pattern of galaxies (points).

Filaments detection (2)

(Tempel, Stoica et. al., 2014) The movie, showing the MCMC in action is available at :http://www.aai.ee/ elmo/sdss-filaments/

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Cluster detection in cosmology : (Tempel, Stoica et. al., 2018)



Figure: The distribution of galaxies in supergalactic coordinates (points) and the visit map obtained using the cluster detection process. The thickness of the slice is 4 Mpc around SGX = 0. Red points show galaxies in the Coma cluster while the other coloured points show galaxies in some clusters with at least five members. The grey points show all remaining galaxies.

Cluster detection in epidemiology : sub-clinical mastitis data

(Stoica, Gay and Kretzschmar, 2007)



Figure: Disease data scores and coordinates for the year 1996 : a) disk configuration obtained using the simulated annealing algorithm ; b) cover probabilities.

Orbit determination (1) : (Kovalenko, Stoica and Emelyanov, 2017)



Figure: Simulated observations (black points): Δx and Δy correspond to relative positions of the secondary with respect to the primary. Lines show a search for the optimal solution during SA algorithm.

Orbit determination (2) :



Figure: Resulting distributions of semi-major axis *a*, eccentricity *e*, inclination *i* and longitude of the ascending node Ω (referenced to J2000 equatorial frame) obtained for simulated observations with the likelihood model. Solid line represents the true parameter value. The doted lines represent the 2.5% and 97.5% quantiles of the resulting sample.

Does the detected pattern really exist ?

Idea : the sufficient statistics of the model \rightarrow morphological descriptors of the shape hidden by the data

- turn the machine at constant temperature T = 1
- compute the average of the sufficient statistics
- compare with the maximum value obtained for the permuted data

Sufficient statistics :

Bisous model (pattern of connected cylinders) : free cylinders, cylinders with one extremity connected, cylinders with both extremities connected

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Test for the filaments existence in galaxy catalogs

Permuted data : keeping the same number of galaxies while spreading them uniformly (binomial point process)

	Data			
Sufficient statistics	NGP150	NGP200	NGP250	
n ₂	4.13	5.83	9.88	
$\bar{n_0}$	15.88	21.19	35.82	
$\bar{n_1}$	21.35	35.58	46.49	

Sufficient statistics	Simulated data (100 binomial catalogs)			
	NGP150	NGP200	NGP250	
max $ar{n_2}$	0.015	0.05	0.015	
max $ar{n_0}$	0.54	0.27	0.45	
$\max \bar{n_1}$	0.39	0.24	0.33	

Test for the cluster existence epidemiological data

Permuted data : keeping the same farm locations while exchanging the score disease

Results :

sufficient statistics for the data from the year 1996 :

$$\bar{n}(y) = 74.10, \quad \bar{\nu}[Z(y)] = 312.46, \quad \bar{n}_o = 555.08$$

 maximum values of the sufficient statistics for 100 simulated data fields

$$\bar{n}(y) = 2.36$$
, $\bar{\nu}[Z(y)] = 13.83$, $\bar{n}_o = 2.62$

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Interpretation : this test does not say if the pattern is well detected, but it says that there is something to be detected ...

Orbit determination validation : position prediction for the obtained parameter values



Figure: The calculated positions (black circles) are compared with given observed positions (crosses) by the x and y coordinates on sky-plane. Black bars denote the 2.5%-97.5% quantiles interval. Dotted line corresponds to the calculated positions for the orbit, obtained with the entire set of observations.

How similar are two data sets ?

Cosmology : compare the sufficient statistics for 22 mock catalogues with the ones for the observation (Stoica, Martinez and Saar, 2010)

Discussion

- mock catalogues exhibit filaments
- mock filaments are generally shorter, more fragmented and more dense
- Bisous model : good for testing the filamentary structure

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Figure: Comparison of the sufficient statistics distributions for the real data (dark box plot) and the mock catalogues.

Some conclusion and perspectives

Spatial models and random geometry :

- Markov marked point processes allow statistical and morphological description of the pattern
- good synthesis properties
- limitations : models remain just models ...

Perspectives :

- ► stochastic processes and random geometry (marked point processes, random fields) → modelling, simulation, statistical inference
- temporal dimension ...
- applications : astronomy and environmental sciences

Acknowledgements : this work was done together with wonderful co-authors and also with the precious help of some very generous persons ...

Exercises : Pattern detection and characterisation.

Exercise 38.

a) Build an Metropolis-Hastings algorithm for sampling the probability distribution :

 $f(x; k, \theta) \propto \Gamma(k, \theta) + 0.75\Gamma(3k, 2\theta)$

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with k = 6 and $\theta = 0.5$. Please explain the choice of the algorithm parameters.

b) Find the maximum of this distribution using the *Simulated Annealing* algorithm.

c) Compare the algorithm performances depending of the following cooling schedules

$$T_{k+1} = cT_k, \quad c \in \{0.99, 0.999, 0.9999, 0.99999\}$$

and

$$T_{k+1} = \frac{T_0}{\log(k+1)+1}, \quad k \in \mathbb{N}.$$

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d) Test the algorithm dependence on the set of initial conditions given by $x_0 = \{0.5, 8, 25\}$.