Random Point Processes

Models, Characteristics and Structural Properties

Volker Schmidt

Department of Stochastics University of Ulm

Introduction Aims

- 1. Explain some basic ideas for stochastic modelling of spatial point patterns
 - stationarity (homogeneity) vs. spatial trends
 - isotropy (rotational invariance)
 - complete spatial randomness
 - interaction between points (clustering, repulsion)

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 - Intensity measure, conditional intensities
 - Pair correlation function, Ripley's K-function, etc.

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- 2. Describe some basic characteristics of point-process models
 - Intensity measure, conditional intensities
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- 3. Present techniques for the statistical analysis of spatial point patterns



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 - Nonparametric estimation of model characteristics
 - Maximum pseudolikelihood estimation

Examples *Point patterns in networks*



Street system of Paris



Cytoskeleton of a leukemia cell

Examples Random tessellations



Poisson line (PLT)



Poisson-Voronoi (PVT)



Poisson-Delaunay (PDT)

Simple and iterated tessellation models



Examples *Model fitting for network data*



Model fitting for telecommunication networks

Examples Analysis of biological networks



a) Actin network at the cell periphery



b) Lamellipodium



c) Region behind lamellipodia

Actin filament networks

Examples Analysis of biological networks



SEM image (keratin)

Graph structure

(PVT/PLT)

Model fitting for keratin and actin networks



SEM image (actin)



Graph structure



Tessellation model

(PLT/PDT)

Other examples Modelling of tropical storm tracks



Storm tracks of cyclons over Japan 1945-2004



Estimated intensity field for initial points of storm tracks over Japan

Other examples Point patterns in biological cell nuclei

Heterochromatin structures in interphase nuclei



3D-reconstruction of NB4 cell nuclei (DNA shown in gray levels) and centromere distributions (shown in red)

Other examples Point patterns in biological cell nuclei





Projections of the 3D chromocenter distributions of an undifferentiated (left) NB4 cell and a differentiated (right) NB4 cell onto the xy-plane

Other examples Point patterns in biological cell nuclei







Capsides of cytomegalovirus

Extracted point pattern

Model for a point field (cluster)

- 1. Mathematical definition of spatial point processes
 - Let $\{X_1, X_2, ...\}$ be a sequence of random vectors with values in \mathbb{R}^2 and
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- 2. A point process $\{X_1, X_2, ...\}$ is called
 - stationary (homogeneous) if the distribution of {X₁, X₂, ...} is invariant w.r.t translations of the origin,
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 - isotropic if the distribution of $\{X_n\}$ is invariant w.r.t rotations around the origin



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 - Equivalent notions: point field instead of spatial point process
 - Point processes are not necessarily dynamic
 - Dynamics (w.r.t. time/space) can be added => spatial birth-and-death processes

Models Stationary Poisson process

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 - Conditional uniformity: Given X(B) = n, the locations of the n points in B are independent and uniformly distributed random variables



Simulated realization of a stationary Poisson process



Simulated realization of a stationary Poisson process

convincingly shows

Complete spatial randomness



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- Complete spatial randomness
- No spatial trend



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No interaction between the points

- 1. Definition of general (non-homogeneous) Poisson processes
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 - Intensity function $\lambda(x) \ge 0$ if $\Lambda(B) = \int_B \lambda(x) \, dx$



Realisation of a Poisson hardcore process



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Constructed from stationary Poisson processes (by random deletion of points)



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- Constructed from stationary Poisson processes (by random deletion of points)
- Realizations are relatively regular point patterns (with smaller spatial variability than in the Poisson case)

Description of the model

Start from a stationary Poisson process (with some intensity $\lambda > 0$)

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- Two-parametric model (with parameters λ and R)



Realization of a Matern-cluster process



Realization of a Matern-cluster process

Constructed from stationary Poisson processes (of so-called cluster centers)



Realization of a Matern-cluster process

- Constructed from stationary Poisson processes (of so-called cluster centers)
- Realizations are clustered point patterns (with higher spatial variability than in the Poisson case)

Description of the model

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 - general Poisson process: $\lambda(u, \mathbf{x}) = \lambda(u)$
- Stationary Strauss process $\lambda(u, \mathbf{x}) = \lambda \gamma^{t(u, \mathbf{x})}$, where $t(u, \mathbf{x}) = \#\{n : |u x_n| < r\}$ is the number of points in $\mathbf{x} = \{x_n\}$ that have a distance from u less than r

More advanced models Strauss process

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• If $0 < \gamma < 1$, then softcore process (repulsion)

Probability density w.r.t. stationary Poisson processes
=> simulation algorithm

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• $s(\mathbf{x})$ the number of pairs $x, x' \in \mathbf{x}$ with |x - x'| < r





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 - Four-parametric model (with parameters λ, γ, r, R)

Probability density $f(\mathbf{x})$ w.r.t. stationary Poisson process in an open bounded set $W \subset \mathbb{R}^2$:

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where $d(X_n)$ is the distance from X_n to its nearest neighbor

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The maximum pseudolikelihood estimate $\hat{\theta}$ of θ is the value which maximizes $PL(\theta; \mathbf{x})$

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 - which includes all the data points x_j as well as some "dummy" points
 - Let $z_j = 1$ if u_j is a data point, and $z_j = 0$ if u_j is a dummy point

Then

$$\log PL(\theta; \mathbf{x}) = \sum_{j=1}^{m} \left(z_j \log \lambda_{\theta}(u_j, \mathbf{x}) - w_j \lambda_{\theta}(u_j, \mathbf{x}) \right)$$
$$= \sum_{j=1}^{m} w_j (y_j \log \lambda_j - \lambda_j)$$

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- Thus, standard statistical software for fitting generalized linear models can be used to compute $\hat{\theta}$

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