# GIST 4302/5302: Spatial Analysis and Modeling Point Pattern Analysis

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Spring 2017

# Spatial Point Patterns



#### Characteristics:

- set of n point locations with recorded "events", e.g., locations of trees, disease or crime incidents  $S = \{s_1, \dots, s_i, \dots, s_n\}$
- point locations correspond to all possible events or to subsets of them
- attribute values also possible at same locations, e.g., tree diameter, magnitude of earthquakes (marked point pattern)
   W = {w<sub>1</sub>,..., w<sub>i</sub>,..., w<sub>n</sub>}

# Analysis objectives:

- detect spatial clustering or repulsion, as opposed to complete randomness, of event locations (in space and time)
- if clustering detected, investigate possible relations with nearby "sources"



#### Further issues:

- analysis of point patterns over large areas should take into account distance distortions due to map projections
- boundaries of study area should not be arbitrary
- analysis of sampled point patterns can be misleading
- one-to-one correspondence between objects in study area and events in pattern

# Simple Descriptive Statistics

# Mean center of a point pattern:

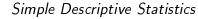
• point with coordinates  $\bar{s} = (\bar{x}, \bar{y})$ :

$$\bar{x} = \frac{\sum_{i=1}^{n} w_i x_i}{\sum_{i=1}^{n} w_i}$$
 and  $\bar{y} = \frac{\sum_{i=1}^{n} w_i y_i}{\sum_{i=1}^{n} w_i}$ 

• center of point pattern, or point with average x and y-coordinates

## Median center of a point pattern:

- <u>both</u> of the following two centers are called <u>median centers</u>, although they are essentially different (confusing!)
  - $\blacktriangleright$  the intersection between the median of the x and the y coordinates
  - ightharpoonup center for minimum distance:  $s_c \in \{s_1, \ldots, s_n\}$ s.t.min  $\sum\limits_{i=1}^n |s_i s_c|$
- the first type of *median center* is not unique, and there is <u>no</u> closed form for the second type
- p-median problem (a typical problem in spatial optimization): the problem of locating p "facilities" relative to a set of "customers" such that the sum of





# Changes of population center (year 1790-2000):





# Descriptive Statistics



#### Standard distance of a point pattern:

 average squared deviations of x and y coordinates from their respective mean:

$$d_{std} = \sqrt{\frac{\sum_{i=1}^{n} (x_i - \bar{x})^2 + \sum_{i=1}^{n} (y_i - \bar{y})^2}{n-2}}$$

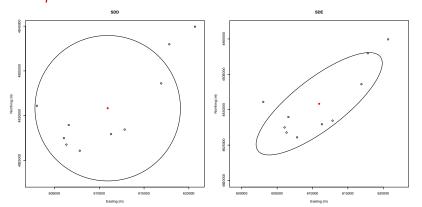
• related to standard deviation of coordinates, a summary circle (centered at  $\overline{s}$  with radius  $d_{std}$ ) of a point pattern

# Standard deviational ellipse:

- Taking directional effects into account for anisotropy cases
- Please refer to Levine and Associates, 2004 for calculations

# Descriptive Statistics

# Examples:



#### Remarks:

- indicates overall shape and center of point pattern
- do not suffice to fully specify a spatial point pattern



# 1st order (i.e., intensity): absolute location of events on map:

- Quadrat methods
- Density Estimation (KDE)
- Moran's I and Geary's C

# 2nd order (i.e., interactions): interaction of events:

- Nearest neighbor distance
- Distance functions G, K, F, L
- Getis-Ord Gi\* and Anselin local Moran's I

### Quadrat methods

Consider a point pattern with n events within a study region A of area |A| Global intensity:

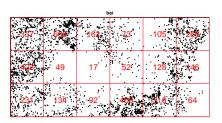
$$\hat{\lambda} = \frac{n}{|A|} = \frac{\text{\#of events within}A}{|A|}$$

## Local intensity via quadrats

- 1. partition A into L sub-regions  $A_l$ ,  $l=1,\ldots,L$  of equal area  $|A_l|$  (also called quadrats)
- 2. count number of events  $n(A_I)$  in each sub-region  $A_I$
- 3. convert sample counts into estimated intensity rates as:

$$\hat{\lambda}(A_I) = \frac{n(A_I)}{|A_I|}$$

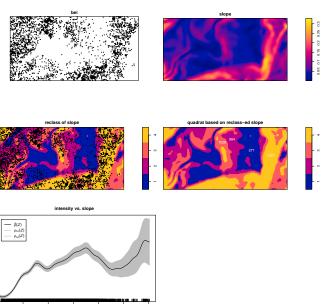




- estimated rates  $\hat{\lambda}(A_I)$  over set of quadrats
- reveal large-scale patterns in intensity variation over A
- larger quadrats yield smoother intensity maps; smaller quadrats yield 'spiky' intensity maps
- size, origin, and shape of quadrats is critical (recall: MAUP)
- only first-order effects are captured



# Dependence of intensity on a covariate (Inhomogeneous cases)





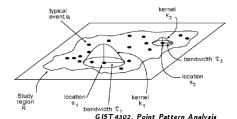
#### Kernel Density Estimation

# Procedure of Kernel Density Estimation (KDE)

- 1. define a kernel K(s; r) of radius (or bandwidth) r centered at any arbitrary location s
- 2. estimate local intensity at s as:

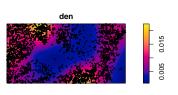
$$\hat{\lambda}(\mathbf{s}) = \frac{1}{n} \sum_{i=1}^{n} K(\mathbf{s}_{i} - \mathbf{s}; r)$$

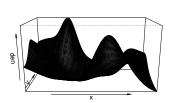
repeat estimation for all points s in the study region to create a density map





# Example for the previous dataset:

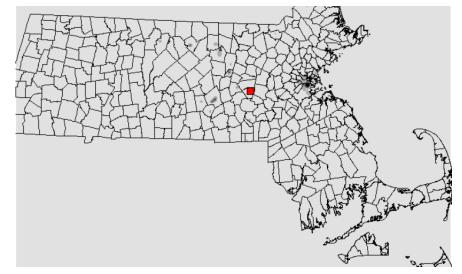




den

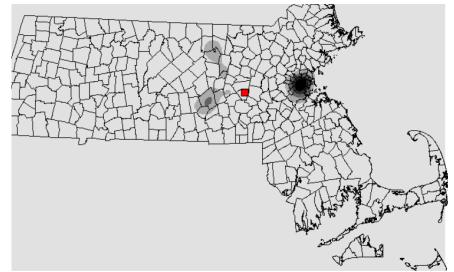


# Example with 2km bandwidth



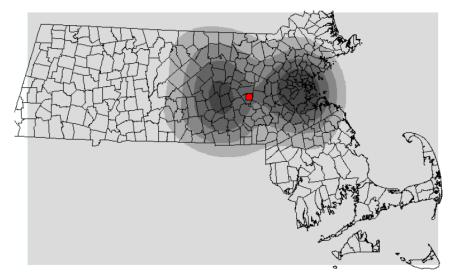


# Example with 10km bandwidth





# Example with 40km bandwidth



### Kernel Density Estimation



#### Comments

- Choice of kernel function is not critical (Diggle, 1985)
- Choice of bandwidth, or degree of smoothing critical:
  - ► Small bandwidth → spiky results
  - ▶ Large bandwidth  $\rightarrow$  loss of detail
- Multi-scale analyses can use these bandwidth characteristics to investigate both broad trends and localized variation
- How to choose bandwidth: choose the degree of smoothing subjectively, by eye, or by formula (Diggle)
- could define local bandwidth based on function of presence of events in neighborhood of s (i.e., adaptive kernel estimation)

What does the output of KDE means?



# Distance-based Descriptors of Point Patterns

- Distances: accessing second order effects
  - Event-to-event distance: distance d<sub>ij</sub> between event at arbitrary location s<sub>i</sub> and another event at another arbitrary location s<sub>j</sub>:

$$d_{ij} = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2}$$

Point-to-event distance: distance  $\tilde{d}_{pj}$  between a randomly chosen point at location  $\tilde{s}_p$  and an event at location  $s_i$ :

$$\tilde{d}_{pj} = \sqrt{(\tilde{x}_p - x_j)^2 + (\tilde{y}_i - y_j)^2}$$

► Event-to-nearest-neighbour distance: distance  $d_{min}(s_i)$  between an event at location  $s_i$  and its nearest neighbor event:

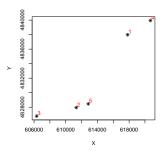
$$d_{min}(s_i) = min\{d_{ii}, j \neq i, j = 1, \dots, n\}$$

Point-to-nearest-neighbour distance (i.e., empty space distance): distance d<sub>min</sub>(s̄<sub>p</sub>) between a randomly chosen point at location s̄<sub>p</sub> and its nearest neighbor event:

$$\tilde{d}_{min}(\tilde{s}_{p}) = min\{\tilde{d}_{pi}, j = 1, \dots, n\}$$



# Event-to-Nearest-Neighbor Distances



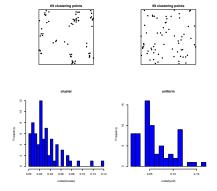
	1	2	3	4	5
1	0.00	11947.70	16042.65	3481.22	10742.98
2	11947.70	0.00	5126.79	15219.58	1599.07
3	16042.65	5126.79	0.00	19481.59	6720.59
4	3481.22	15219.58	19481.59	0.00	13913.70
5	10742.98	1599.07	6720.59	13913.70	0.00

Table: Euclidean distance matrix



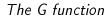
# Event-to-Nearest-Neighbor Distances

# Nearest neighour distances



• Mean nearest neighbour distance: Average of all  $d_{min}(s_i)$  values

$$\bar{d}_{min} = \frac{1}{n} \sum_{i=1}^{n} d_{min}(s_i)$$

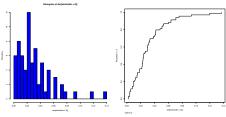


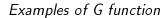


• Definition: nearest neighbour distance function, i.e., proportion of event-to-nearest-neighbor distances  $d_{min}(s_i)$  no greater than given distance cutoff d, estimated as:

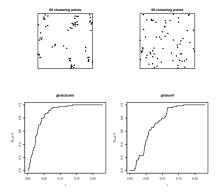
$$\hat{G}(d) = \frac{\#\{d_{min}(s_i) < d, i = 1, ..., n\}}{n}$$

- alternative definition: cumulative distribution function (CDF) of all n event-to-nearest-neighbor distances; instead of computing average d<sub>min</sub> of d<sub>min</sub> values, compute their CDF
- the G function provides information on event proximity
- example for previous clustering point pattern:









### Expected plot:

- for clustered events,  $\hat{G}(d)$  rises sharply at short distances, and then levels off at larger d-values
- for randomly-spaced events,  $\hat{G}(d)$  rises gradually up to the distance at which most events are spaced, and then increases sharply

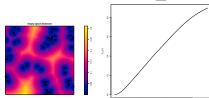


• proportion of point-to-nearest-neighbor distances (i.e., empty space distances)  $\tilde{d}_{min}(s_p)$  no greater than given distance cutoff d, estimated as:

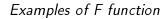
$$\hat{F}(d) = \frac{\#\{\tilde{d}_{min}(\tilde{s}_p) < d, p = 1, \dots, m\}}{m}$$

- alternative definition: cumulative distribution function (CDF) of all
   m point-to-nearest-neighbor distances
- the F function provides information on event proximity to voids
- Examples for previous clustering point pattern:

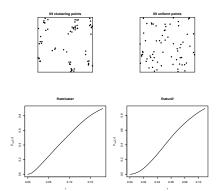




GIST 4302, Point Pattern Analysis





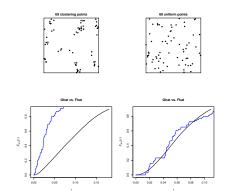


# Expected plot:

- for clustered events,  $\hat{F}(d)$  rises sharply at short distances, and then levels off at larger d-values
- for randomly-spaced events,  $\hat{F}(d)$  rises rapidly up to the distance at which most events are spaced, and then levels off (there are more nearest neighbors at small distances from randomly placed points)



# Comparing G and F functions



# Expected plot:

- for clustered events,  $\hat{G}(d)$  rises faster
- for randomly-spaced events,  $\hat{F}(d)$  tends to be close to  $\hat{G}(d)$





Working with pair-wise distances&looking beyond nearest neighours

### Concept

- 1. construct set of concentric circles (of increasing radius d) around each event
- 2. count number of events in each distance "band"
- 3. cumulative number of events up to radius d around all events becomes the sample K function  $\hat{K}(d)$





Working with pair-wise distances&looking beyond nearest neighours

Formal definition:

$$K(d) = \frac{1}{\lambda} \frac{\#\{d_{ij} \leq d, i, j = 1, \dots, n\}}{n}$$

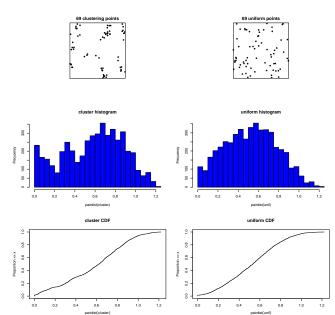
$$= \frac{|A|}{n} \frac{\#\{d_{ij} \leq d, i, j = 1, \dots, n\}}{n}$$

$$= |A| \text{(proportion of event-to-event distance } \leq d\text{)}$$

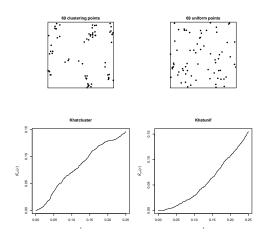
• In other words, the  $\hat{K}(d)$  is the sample cumulative distribution function (CDF) of all  $n^2$  event-to-event distances, scaled by |A|



# Examples of Event-to-Event Distance Histogram and CDFs







• the sample K function  $\hat{K}(d)$  is monotonically increasing and is a scaled (by area |A|) version of the CDF of E2E distances





# Spatial point patterns

• set of *n* point locations with recorded "events"

# Describing the first-order effect

- overal intensity
- local intensity (quadrat count and kernel density estimation)

# Describing the second-order effect

- nearest neighbour distances
  - ► the G function
- empty space distances
  - ► the F function
- pair-wise distances
  - ► the K function





#### Caveats:

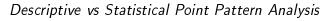
- theoretical G, F, K functions are defined and estimated under the assumption that the point process is stationary (homogeneous)
- these summary functions do not completely characterise the process
- if the process is not stationary, deviations between the empirical and theoretical functions (e.g. K and K) are not necessarily evidence of interpoint interaction, since they may also be attributable to variations in intensity











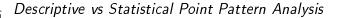


### Descriptive analysis:

- set of quantitative (and graphical) tools for characterizing spatial point patterns
- different tools are appropriate for investigating first- or second-order effects (e.g., kernel density estimation versus sample G function)
- can shed light onto whether points are clustered or evenly distributed in space

#### Limitation:

- no assessment of <u>how</u> clustered or <u>how</u> evenly-spaced is an observed point pattern
- no yardstick against which to compare observed values (or graph) of results





# Statistical analysis:

- assessment of whether an observed point pattern can be regarded as one (out of many) realizations from a particular spatial process
- measures of confidence with which the above assessment can be made (how likely is that the observed pattern is a realization of a particular spatial process)

# Are daisies randomly distributed in your garden?





# Complete Spatial Randomness (CSR)

# Complete Spatial Randomness (CSR)

- yardstick, reference model that observed point patterns could be compared with, i.e., null hypothesis
- = homogeneous (uniform) Poisson point process
- basic properties:
  - ► the number of points falling in any region A has a Poisson distribution with mean  $\lambda |A|$
  - given that there are n points inside region A, the locations of these points are i.i.d. and uniformly distributed inside A
  - $\blacktriangleright$  the contents of two disjoint regions A and B are independent







#### Quadrat counting test

- partition study area A into L sub-regions (quadrats),  $A_1, \ldots, A_L$
- count number of events  $n(A_I)$  in each sub-region  $A_I$
- Under the null hypothesis of CSR, the  $n(A_I)$  are i.i.d. Poisson random variables with the same expected value
- The Pearson  $\chi^2$  goodness-of-fit test can be used
  - ▶ test statistics: Pearson residual  $\sum_{l} \epsilon(A_{l})^{2}$

$$\epsilon(A_I) = \frac{n(A_I) - \mu(A_I)}{\sqrt{\mu(A_I)}},$$

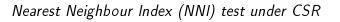
where  $\mu(A_l)$  indicates the expected number of events in  $A_l$ 

 $ightharpoonup \sum_{l=1}^{J} \epsilon(A_l)^2$  is assumed to follow  $\chi^2$  distribution





- three values indicate the number of observations, CSR-expected number of observations, and the Pearson residuals
- p-value = 0.617





# Nearest neighbour index

• Compares the mean of the distance observed between each point and its nearest neighbor  $(\bar{d}_{min})$  and the expected mean distance under CSR  $E(d_{min})$ 

$$NNI = \frac{\bar{d}_{min}}{E(d_{min})}$$

• Under CSR, we have:

$$E(d_{min}) = \frac{1}{2\sqrt{\lambda}}$$

$$\sigma(d_{min}) = \frac{0.26136}{\sqrt{n^2/A}}$$



### Nearest neighbour index test

Test statistics:

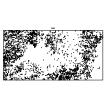
$$z = \frac{\bar{d}_{min} - E(d_{min})}{\sigma(d_{min})},$$

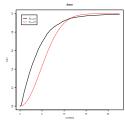
• z is assumed to follow Gaussian distribution, thus, if z<-1,96 or z>1.96, we are 95% confident that the distribution is not randomly distributed



- The G function is a function of nearest-neighbour distances
- For a homogeneous Poisson point process of intensity  $\lambda$ , the nearest-neighbur distance distribution (the G function) is known to be:

$$G(d) = 1 - \exp\{-\lambda \pi d^2\}$$





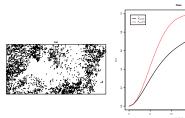


#### The F Function under CSR

- The F function is a function of empty space distances
- For a homogeneous Poisson point process of intensity  $\lambda$ , the empty space distance distribution (the F function)is known to be:

$$F(d) = 1 - \exp\{-\lambda \pi d^2\}$$

- Equivalent to the G function
- Intuitively, because points (events) of the Poisson process are independent of each other, the knowledge that a random point is a event of a point pattern does not affect any other event of the process





#### The K Function under CSR

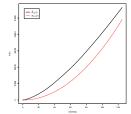
- The K function is a function of pair-wise distances
- For a homogeneous Poisson point process of intensity  $\lambda$ , the pair-wise distance distribution (the K function) is known to be:

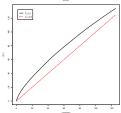
$$K(d) = \pi d^2$$

• A commonly-used transformation of K is the L-function:

$$L(d) = \sqrt{\frac{K(d)}{\pi}} = d$$

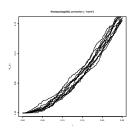








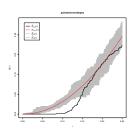
 because of random variability, we will never obtain perfect agreement between sample functions (say the K function) with theoretical functions (the theoretical K functions), even with a completely random pattern



#### Monte Carlo test



- A Monte Carlo test is a test based on simulations from the null hypothesis
- Basic procedures:
  - ▶ generate M independent simulations of CSR inside the study region A
  - rightharpoonup compute the estimated K functions for each of these realisations, say  $K^{(j)}(r)$  for  $j=1,\ldots,M$
  - obtain the pointwise upper and lower envelopes of these simulated curves
  - ► not a confidence interval





# Statitsical analysis of spatial point patterns:

- allows to quantify departure of results obtained via exploratory tools, e.g.,  $\hat{G}(d)$ , from expected such results derived under specific null hypotheses, here CSR hypothesis
- can be used to assess to what extent observed point patterns can be regarded as realizations from a particular spatial process (here CSR)
- Same concepts can be applied for hypothesis of other types of point processes (e.g., Poisson cluster process, Cox process)

# Sampling distribution of a test statistics

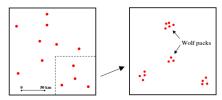
- lies at the heart of any statistical hypothesis testing procedure, and is tied to a particular null hypothesis
- simulation and analytical derivations are two alternative ways of computing such sampling distributions (the latter being increasingly replaced by the former)

# Edge Effects



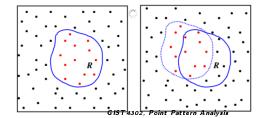
# Scale effects

Wolf pack example



• Nearest neighour distance (NN distance, G,F functions) vs K function

# Edge effects





# Extended into line processes

• Line density

