Statistics 222, Spatial Statistics.

Outline for the day:

1. Old projects and exam on the course website

http://www.stat.ucla.edu/~frederic/222/F23 .

- 2. Fitting Hawkes models with covariates. fithawkeswithcovariates.r
- 3. Integral term in loglikelihood for Hawkes processes.
- 4. Analytic nonparametric estimation.
- Next week we will do review and Stoyan-Grabarnik statistic and Stoyan-Grabarnik estimation.

3. The integral term in the loglikelihood for Hawkes processes.

loglikelihood = $\sum_{i} \log(\lambda(t_i, x_i, y_i)) - \iiint \lambda(t, x, y) dxdydt.$

The space-time region is $B = [0,T] \times S$.

For a Hawkes process, $\lambda(t,x,y) = \mu \rho(x,y) + K \sum_{i:ti < t} g(t-t_i, x-x_i, y-y_i)$, where ρ and g are densities.

 $\int_0^T \iint \lambda(t,x,y) \, dx dy dt = \int_0^T \iint \mu \, \rho(x,y) \, dx dy dt + \int_0^T \iint K \sum_{i:ti < t} g(t-t_i, x-x_i, y-y_i) \, dx dy dt$

$$= \mu T + \int_0^T \iint K \int_B 1_{\{t' < t\}} g(t-t', x-x', y-y') dN(t', x', y') dxdydt$$

interchanging the integrals

$$= \mu T + K \int_B \int_0^T \iint 1_{\{t' < t\}} g(t-t', x-x', y-y') dxdydt dN(t',x',y')$$

changing coordinates, letting u = t-t', v = x-x', w = y-y',

$$= \mu T + K \int_{B} \int_{0}^{T-t'} \iint_{S-(x',y')} g(u, v, w) dudvdw dN(t', x', y')$$

~ μ T + K $\int_{B}(1) dN(t',x',y')$

 $= \mu T + KN(B).$

This is approximate because typically $\int_0^{\infty} \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} g(u,v,w) \, du \, dv \, dw = 1$, but instead, we have $\int_0^{T-t'} \iint_{S-(x',y')} g(u,v,w) \, du \, dv \, dw$ which is often close to 1.

4. Nonparametric triggering function estimation.

Marsan and Lengliné (2008) assume g is a step function, and estimate steps β_k as parameters.

$$\ell(heta) = \sum_i \log \left(\lambda(au_i, \mathbf{x_i} | \mathcal{H}_{ au_i})
ight) - \int_0^T \int_S \lambda(t, \mathbf{x} | \mathcal{H}_t) \, d\mathbf{x} dt$$

Setting the partial derivatives of this loglikelihood with respect to the steps β_k to zero yields

$$0 = \partial \ell(\theta) / \partial \beta_k = \sum_{(i,j): \tau_i - \tau_j \in U_k} K / \lambda(\tau_i) - Kn |U_k|,$$

where $|U_k|$ is the width of step k, for k = 1, 2, ..., p. This is a system of *p* equations in *p* unknowns. However, the equations are nonlinear. They depend on $1/\lambda(\tau_i)$.

Gradient descent methods: way too slow for large p.

Marsan and Lengliné (2008) find *approximate* maximum likelihood estimates using the E-M method for point processes. You pick initial values of the parameters, then given those, you know the probability event *i* triggered event *j*. Using these, you can weight each pair of points by its probability and re-estimate the parameters, and repeat until convergence. This method works well but is iterative and time-consuming.

Analytic solution.

Set p = n. (p = number of steps in the step function, g, and <math>n = # of observed points.) Setting the derivatives of the loglikelihood to zero we have the p equations

$$0 = \partial \ell(\theta) / \partial \beta_k = \sum_{(i,j):\tau_i - \tau_j \in U_k} K / \lambda(\tau_i) - K n |U_k|,$$

which are *p* linear equations in terms of $1/\lambda(\tau_i)$, for i = 1, 2, ..., n. (!)

So, if p=n, then we can use these equations to solve for $1/\lambda(\tau_i)$,

and if we know $1/\lambda(\tau_i)$, then we know $\lambda(\tau_i)$,

and if we know $\lambda(\tau_i)$, then we can solve for β_i because the def. of a Hawkes process is

$$\lambda(\tau_j) = \mu + K \sum_{i < j} g(\tau_j - \tau_i),$$

which results in *n* linear equations in the *p* unknowns β_1 , β_2 , ..., β_p , when *g* is a step function.

Analytic solution.

We can write the resulting estimator in very condensed form.

Let $\boldsymbol{\lambda} = \{\lambda(\tau_1), \lambda(\tau_2), ..., \lambda(\tau_n)\}.$

Suppose the steps of g have equal widths, $|U_1| = |U_2|$, etc. Call this width U.

Let A_{ij} = the number points τ_k such that $\tau_j - \tau_k$ is in U_i, for *i*,*j* in {1,2,...,*p*}.

Then the loglikelihood derivatives equalling zero can be rewritten

$$0 = KA(1/\lambda) - Kb,$$

where $\mathbf{b} = nU\mathbf{1}$, with $\mathbf{1} = \{1, 1, ..., 1\}$.

This has solution $\underline{1/\lambda} = A^{-1}b$, if A is invertible.

Similarly, the Hawkes equation can be rewritten $\lambda = \mu + KA^T \beta$, whose solution is

$$\hat{\boldsymbol{b}} = (\boldsymbol{K}\boldsymbol{A}^T)^{-1}(\boldsymbol{\lambda}\boldsymbol{-}\boldsymbol{\mu}).$$

Combining these two underlined formulas yields the estimates

$$\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$$

This is very simple, trivial to program, and rapid to compute.

Analytic solution.

There are problems, however.

1. Estimating n=p steps. High variance.

However, if we can assume g is smooth, then we can smooth our estimates for stability.

2. Need to estimate K and μ too.

We can use Marsan and Lengliné's method or take derivatives for these as well.

3. What about spatially-varying steps and unequally sized steps for g?

No problem. The estimation generalizes in a completely obvious way.

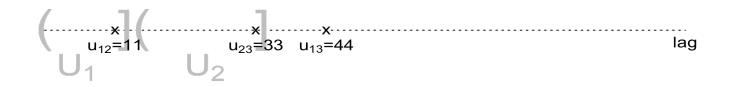
4. A can be singular.

We may need better solutions for this.

I let $u_j = \tau_j - \tau_{j-1}$, sorted the u_j values, and then used $[u_{(1)}, u_{(2)})$, etc. as my binwidths, so each row and column of *A* would have at least one non-zero entry. If it still isn't singular, adding in a few random 1's into *A* often helps.

$$\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$$





4. A can be singular.

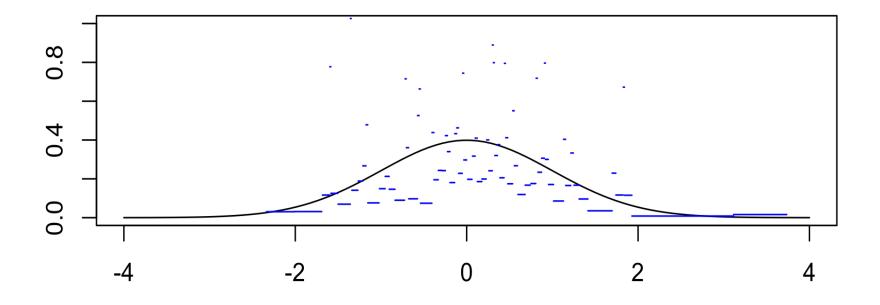
We may need better solutions for this.

I let $u_j = \tau_j - \tau_{j-1}$, sorted the u_j values, and then used $[u_{(1)}, u_{(2)})$, etc. as my binwidths, so each row and column of *A* would have at least one non-zero entry. If it still isn't singular, adding in a few random 1's into *A* often helps.

$$\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$$

Note: take the simple case of a dataset where point i is only influenced by point i-1. This is basically a renewal process, and we are just estimating a renewal density.

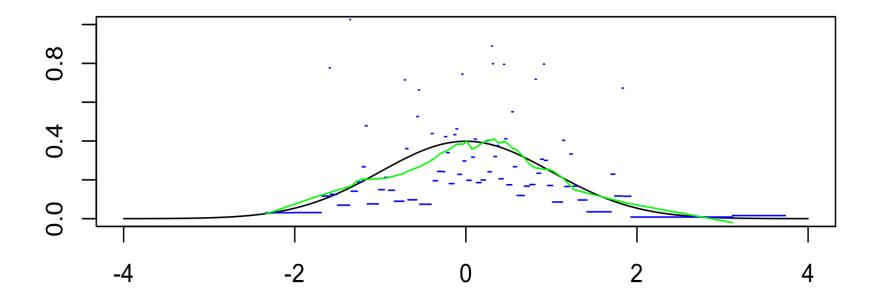
Here A = I, K = 1, and we get the density estimator $1/\{n(x_i - x_{i-1})\}$.



 $\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$

Note: take the simple case of a dataset where point i is only influenced by point i-1. This is basically a renewal process, and we are just estimating a renewal density.

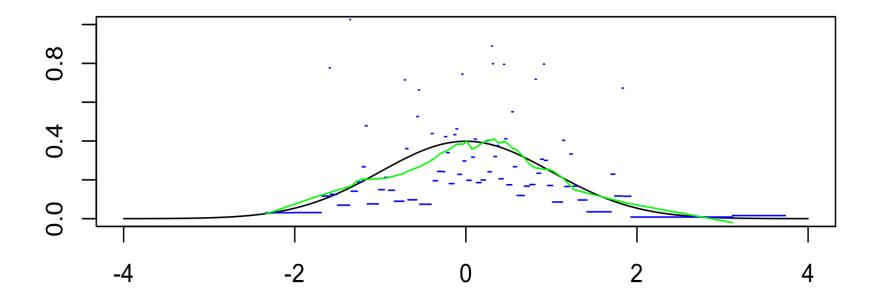
Here A = I, K = 1, and we get the density estimator $1/\{n(x_i - x_{i-1})\}$.



$$\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$$

Note: take the simple case of a dataset where point i is only influenced by point i-1. This is basically a renewal process, and we are just estimating a renewal density.

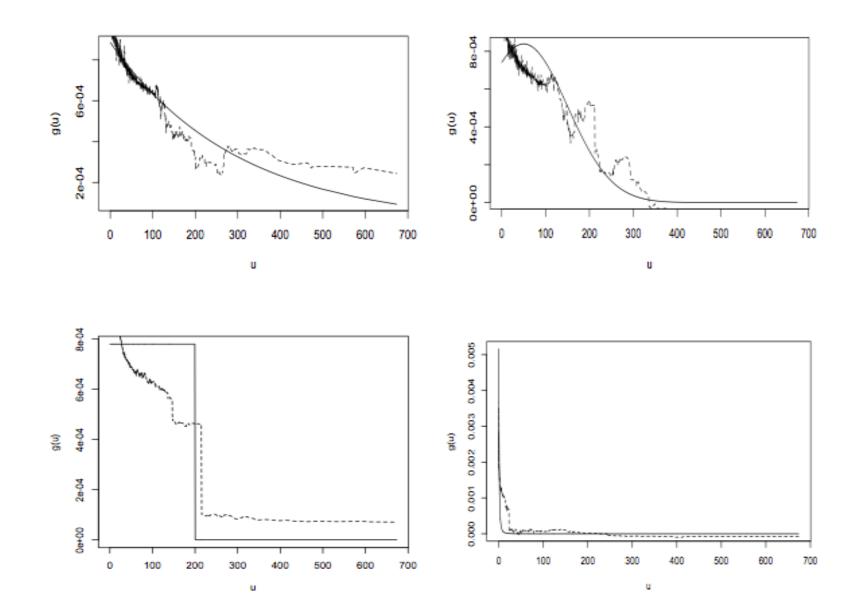
Here A = I, K = 1, and we get the density estimator $1/\{n(x_i - x_{i-1})\}$.



$$\hat{\beta} = (KA^T)^{-1}[1/(A^{-1}b) - \mu]$$

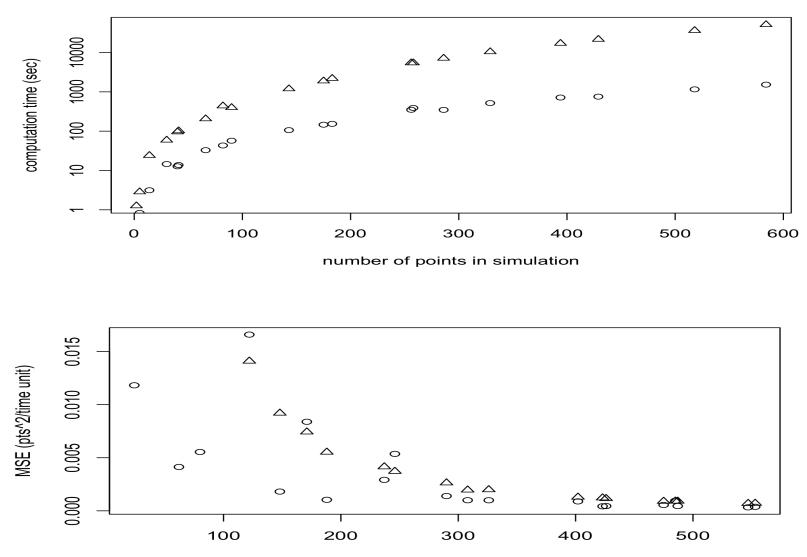
Computation time and performance comparison.

Test of concept. Examples of exponential, truncated normal, uniform, and Pareto g.



Computation time and performance comparison.

Triangles = Marsan and Lengliné (2008) method. Circles = analytic method.



number of points in simulation