Markov-switching (discrete-time) Hawkes process

S. Robin

joint work with A. Bonnet

LPSM, Sorbonne université

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'Motivation'

Counting process

Overnight recording of bat cries in continuous time



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Can we detect changes in the distribution of events?



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- Can we detect changes in the distribution of events?
- Can we associate each time period with some underlying behavior?



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Modelling. Point process with (latent) Markov switching regime.

Point process

Point process

Reminder.



- $(T_k)_{k \ge 1}$ a random collection of points
- Count process $H(t) = \sum_{k \ge 1} \mathbb{I}\{T_k \le t\}$
- Intensity function $\lambda(t)$: immediate probability of observing an event at time t

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Examples

- Homogeneous Poisson process: $\lambda(t) \equiv \lambda$
- Heterogeneous Poisson process: $\lambda(t) = deterministic function$
- Hawkes process: $\lambda(t)$ = function of the past events = random function

Outline

(Discrete) Hawkes process

Continuous-time Hawkes process

Discrete-time Hawkes process Markovian representation

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Model Identifiability & Inference

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Univariate Hawkes process



(Conditional) intensity function for the Hawkes process [Haw71]:

$$\lambda(t) = \lambda(t \mid \mathcal{H}_t) = \lambda_0 + \sum_{T_k < t} h(t - T_k)$$

• $\lambda_0 = \text{baseline}$

h = kernel = influence of past events

Self-exciting exponential Hawkes process

$$\lambda(t) = \lambda_0 + \sum_{T_k < t} a e^{-b(t - T_k)}$$

Self exciting: Each event increases the probability of observing another event

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Self exciting: Each event increases the probability of observing another event



- Exponential kernel function $h(t) = ae^{-bt}$
- $a \ge 0$ to ensure that λ is non negative
- ▶ a/b < 1 to ensure stationarity</p>
- Applications: sismology, epidemiology, vulcanology, neurosciences, ecology, ...

Cluster representation [H074]



- Immigrants arrive at rate λ_0
- Each immigrant or descendant produces new individuals at rate h(t T)

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Discrete-time Hawkes process

Continuous time exponential Hawkes process

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Discretization [Seo15,Kir16,Kir17]

- $I_k = [\tau_{k-1}; \tau_k]$ with $\tau_k = k\Delta$
- $H_k = H(I_k)$ the number of events on I_k



• Distribution of $(H_k)_{k \ge 1}$?

Decomposition of the count

 H_k = number of events on $I_k = [\tau_{k-1}; \tau_k]$

$$H_k \stackrel{\Delta}{=} B_k + \sum_{\ell \leqslant k-1} \sum_{T \in I_\ell} M_T(I_k) + R_k$$

where

•
$$B_k$$
 = number of immigrants within I_k :

$$B_k \sim \mathcal{P}(\mu)$$

with $\mu = \lambda_0 \Delta$,

• $M_T(I_k)$ = number of descendants of $T < \tau_k$ within I_k :

$$M_{T}(I_{k}) \sim \mathcal{P}\left(\int_{I_{k}} ae^{-b(t-T)} dt\right) = \mathcal{P}\left(\alpha e^{-b(\tau_{k}-T)}\right)$$

with $lpha={\it a}({\it e}^{b\Delta}-1)/{\it b}$,

▶ R_k = number of descendants of points $T \in I_k$ within I_k

Discrete time Hawkes process

When Δ is small:

- $R_k \simeq 0$
- For $T \in I_{\ell}$: $e^{-b(\tau_k T)} \simeq e^{-b(\tau_k \tau_{\ell})} = \beta^{k-\ell}$ with $\beta = e^{-b\Delta}$, so

$$\sum_{\ell \leqslant k-1} \sum_{T \in I_{\ell}} M_{T}(I_{k}) \stackrel{\Delta}{\simeq} \sum_{\ell \leqslant k-1} \sum_{T \in I_{\ell}} \mathcal{P}\left(\alpha \beta^{k-\ell}\right) \stackrel{\Delta}{=} \mathcal{P}\left(\sum_{\ell \leqslant k-1} H_{k-\ell} \alpha \beta^{\ell-1}\right)$$

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Discrete-time Hawkes process $Y = \{Y_k\}_{k \leq 1}$.

$$\mathbf{Y}_{k} \mid (\mathbf{Y}_{\ell})_{\ell \leqslant k-1} \sim \mathcal{P}\left(\mu + \sum_{\ell=1}^{k-1} \alpha \beta^{\ell-1} \mathbf{Y}_{k-\ell}\right)$$

See [Kir16] for the convergence toward a continuous-time Hawkes process.

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 $\rightarrow (Y_k)_{k \ge 1}$ is not a Markov chain (infinite memory).

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Markovian representation.

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• we have for $k \ge 1$ (with $U_0 = Y_0 = 0$)

 $U_k = \alpha Y_{k-1} + \beta U_{k-1}, \qquad Y_k \mid U_k \sim \mathcal{P}(\mu + U_k).$

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 $\rightarrow ((Y_k, U_k))_{k \ge 1}$ forms a Markov Chain.

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Graphical model

Discrete time Hawkes process.

$$(Y_k)_{k \ge 1} \sim \text{Discrete Hawkes}(\mu, \alpha, \beta)$$

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$$U_1 = 0, \qquad \qquad U_k = \alpha Y_{k-1} + \beta U_{k-1}, \qquad \qquad Y_k \sim \mathcal{P}(\mu + U_k)$$

Graphical model:

$$p(U_k, Y_k \mid (U_{\ell}, Y_{\ell})_{\ell \leq k-1}) = p(U_k, Y_k \mid U_{k-1}, Y_{k-1})$$

= $p(U_k \mid U_{k-1}, Y_{k-1}) p(Y_k \mid U_k)$

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• Hidden path: $(Z_k)_{k \ge 1}$ homogeneous Markov chain with Q states, transition matrix π and initial distribution ν :

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• Observed counts: for $k \ge 1$ and

$$(\mathbf{Y}_k \mid (\mathbf{Y}_\ell)_{\ell \leq k-1}, \mathbf{Z}_k = \mathbf{q}) \sim \mathcal{P}\left(\mu_{\mathbf{q}} + \sum_{\ell=1}^{k-1} \alpha \beta^{\ell-1} \mathbf{Y}_{k-\ell}\right)$$

or, for
$$k \ge 1$$
 (with $U_0 = Y_0 = 0$)

$$U_{k} = \alpha Y_{k-1} + \beta U_{k-1}, \qquad \qquad Y_{k} \mid U_{k} \sim \mathcal{P} \left(\mu_{\mathbb{Z}_{k}} + U_{k} \right)$$

Model: Q hidden states

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• Observed counts: for $k \ge 1$ and

$$(Y_k \mid (Y_\ell)_{\ell \leq k-1}, Z_k = q) \sim \mathcal{P}\left(\mu_q + \sum_{\ell=1}^{k-1} \alpha \beta^{\ell-1} Y_{k-\ell}\right)$$

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Assumptions:

- The immigration rate μ varies with the hidden state
- The distribution of the number of offspring (α, β) does not vary with the hidden state

Graphical model:

$$\xrightarrow{\gamma} \begin{array}{c} & & Z_{k-1} \longrightarrow Z_k \longrightarrow Z_{k+1} \longrightarrow Z_{$$

 $(Z_k)_{k \ge 1} = hidden path, \quad (U_k)_{k \ge 1} = memory, \quad (Y_k)_{k \ge 1} = observed process.$

Graphical model:

$$\xrightarrow{\gamma} Z_{k-1} \xrightarrow{Z_k} Z_{k+1} \xrightarrow{Z_{k+1}} Z_{k+1} \xrightarrow$$

 $(Z_k)_{k \geqslant 1} =$ hidden path, $(U_k)_{k \geqslant 1} =$ memory, $(Y_k)_{k \geqslant 1} =$ observed process.

Remarks:

- ▶ The memory of the past is 'stored' in the variable U_k , which can still be computed recursively $(U_k = \alpha Y_{k-1} + \beta U_{k-1})$
- The Markovian property still holds if the influence of the past varies with the hidden state $(\alpha \rightarrow \alpha_q, \beta \rightarrow \beta_q)$.

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Identifiability

Proposition: The model parameter $\theta = (\nu, \pi, (\mu_q)_{1 \leq q \leq Q}, \alpha, \beta)$ is identifiable from the joint distribution $p_{\theta}^{Y_1, Y_2, Y_3}$:

$$\theta' \neq \theta \qquad \Rightarrow \qquad p_{\theta'}^{Y_1,Y_2,Y_3} \neq p_{\theta}^{Y_1,Y_2,Y_3}.$$

¹The generic technique from [AMR09] does not apply here.
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Sketch of proof. Finite Poisson mixtures are identifiable [Tei61], so, because¹

$$p_{\theta}^{Y_1, Y_2, Y_3}(x, y, z) = \sum_{1 \leq q, \ell, m \leq Q} \nu_q \pi_{q\ell} \pi_{\ell m} \mathcal{P}(x; \mu_q) \mathcal{P}(y; \mu_\ell + \alpha x) \mathcal{P}(z; \mu_m + \alpha \beta x + \alpha y),$$

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1. ν and μ can be identified from $p_{\theta}(Y_1)$, [sum over y and z]

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- 2. then α can be identified from $p_{\theta}(Y_2 \mid Y_1 = 1)$, [fix x = 1, sum over z]

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4. then π can be identified from the joint mixture [sum over z]

$$p_{\theta}^{Y_1,Y_2}(x,y) = \sum_{1 \leq q,\ell \leq Q} \nu_q \pi_{q\ell} \mathcal{P}(x;\mu_q) \mathcal{P}(y;\mu_\ell + \alpha x),$$

which is proven identifiable.

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Aim: Infer the parameter θ

$$\widehat{ heta} = rg\max_{ heta} \log p_{ heta}(Y)$$

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EM algorithm for HMM: [DLR77,CMR05]

$$\theta^{(h+1)} = \underset{\theta}{\operatorname{arg\,max}} \underbrace{\mathbb{E}_{\theta^{(h)}}}_{\mathsf{E} \text{ step}} [\log p_{\theta}(Y, Z) \mid Y]$$

► E step: Evaluate $Q(\theta \mid \theta^{(h)}) = \mathbb{E}_{\theta^{(h)}}[\log p_{\theta}(Y, Z) \mid Y]$ (forward-backward recursion)

• M step: Gradient descent, computing $\nabla_{\theta} Q(\theta \mid \theta^{(h)})$ by recursion

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Classification:

Marginal:
$$\widehat{Z}_k = \arg \max_q P_{\widehat{\theta}} \{ Z_k = q \mid Y \},$$
Joint (Viterbi): $\widehat{Z} = \arg \max_z P_{\widehat{\theta}} \{ Z = z \mid Y \}$

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Model selection: Penalized likelihood

$$\begin{aligned} AIC_Q &= \log p_{\hat{\theta}_Q}(Y) - D_Q, \\ BIC_Q &= \log p_{\hat{\theta}_Q}(Y) - D_Q \frac{\log(N)}{2} \end{aligned}$$

with D_Q = number of parameters = 2 + Q^2 and N = number of time bins.

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Simulation design (Q = 3)

▶ Baseline continuous parameters: $m^0 = [10, 200, 1000], a^0 = 40, b = 160$

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Simulated process:

$$(H_t)_{0 \le t \le 1} \sim Heterogeneous Continuous Hawkes(a, b^0, m)$$

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Simulated process:

 $(H_t)_{0 \leq t \leq 1} \sim$ Heterogeneous Continuous Hawkes (a, b^0, m)

• Discretized process: n = H(1)

$$N = c n, \qquad c = 0.5, \ 1, \ 2, \ 4,$$
$$Y_k = H\left(\left[\frac{k-1}{N}; \frac{k}{N}\right]\right), \qquad k = 1, \dots N.$$

 \rightarrow not a discrete-time Hawkes process as defined earlier

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Simulation results ($Q^* = 3$)



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Simulation results ($Q^* = 3$)

Model selection: BIC. Distribution of $BIC_Q - BIC_1$



nb events

nb events

000

20

200

8

2000

8

00

8

2000

000

200

8



2

lam=1.5 c=0.5















2

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lam=1.5 c=2

lam=2 c=2

8

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8

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8

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2

0

8

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99

8

2 3

lam=1.5 c=4

à. à

з

lam=2 c=4

lam=0.5 c=4





Qbic lam=0.5

Qbic lam=1



Qbic lam=1.5



Qbic lam=2



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Simulation results ($Q^* = 3$)

Model selection: AIC. Distribution of $AIC_Q - AIC_1$









lam=1 c=2





















nb events

nb events

nb events



lam=1.5 c=0.5









lam=1.5 c=1









lam=1.5 c=2

lam=2 c=2





з



lam=1.5 c=4

Simulation results ($Q^* = 3$)



Simulation conclusions

- Inference easier when more signal (large λ)!!!
- Inference easier with thinner discretization step (large N) But at the price of a higher computational cost
- BIC does not capture the right number of states Sequences not simulated according to the model
- AIC does, with reasonable signal (λ) and discretization (N) Blind to the simulation shift from the model?

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Practical recommendations.

Take N = 2n and use AIC

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Bat cries

Data set. 1555 overnight recordings all over France

Bat cries

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Poisson vs Hawkes / Homogeneous vs HMM. Best model based on AIC

	Poisson	Hawkes	Total
Homogeneous	34	353	387
Hidden Markov	24	1144	1168
Total	58	1497	1555

- ▶ Memory (95%) and heterogeneity (75%) are present in most sequences
- Hawkes-HMM best fits almost 3 sequences out of 4.

Example



- Poisson-HMM needs many state changes to account for self-excitation
- Hawkes-HMM state changes do not correspond to slope changes

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States and species

The number of bat species was also recorded



The number of states does not match the number of species

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Summary

What we did.

▶ The discretized Hawkes process with exponential kernel is a Markov model

 \Rightarrow The discretized Markov switching Hawkes process with exponential kernel is a hidden Markov model

- The standard EM machinery applies to achieve maximum likelihood inference.
- Not shown: initialization based on existing estimation procedures for homogeneous Hawkes ([Che21],[CL22]) and Poisson HMM.

Discussion

What we did not do.

- Goodness-of-fit: 'Poissonisation' (on-going).
- ▶ Model selection: derive a proper (BIC?) criterion accounting for the discretization step.
- Understand the inferred latent states in terms of animal behavior, biogeography, species, ...

Discussion

What we did not do.

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In parallel. With C. Dion-Blanc, D. Hawat and E. Lebarbier

- Efficient change-point detection ('segmentation') in (marked) Poisson & Hawkes processes.
 → Dynamic programming applies [DBHLR24].
- Segmentation-classification of Poisson processes.

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Discrete HMM

Conversion formulas from continuous to discrete Hawkes

$$\alpha = \frac{a(e^{b\Delta} - 1)}{b}, \qquad \beta = e^{-b\Delta}$$

Discrete HMM

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3-step initialization

- Homogeneous Hawkes for the reproduction parameters α and β (hawkesbow R package [Che21])
- Poisson-HMM for the rates μ_1, \ldots, μ_Q and transition π
- Correction $\mu_k \rightarrow \widetilde{\mu}_k$ to account for reproduction rate

Simulation results ($Q^* = 1$, N = cn, $m^0 = 400$)



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Markov-switching (discrete-time) Hawkes process

Lancaster'25

Simulation results ($Q^* = 1$, N = cn, $m^0 = 400$)



Simulation results ($Q^* = 1$, N = cn, $m^0 = 400$)

Model selection: AIC. Distribution of $AIC_Q - AIC_1$





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Simulation results ($Q^* = 2$, N = cn, $m^0 = [10, 800]$)


Simulation results ($Q^* = 2$, N = cn, $m^0 = [10, 800]$)

Model selection: BIC. Distribution of $BIC_Q - BIC_1$

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Simulation results ($Q^* = 2$, N = cn, $m^0 = [10, 800]$)

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Model selection: AIC. Distribution of $AIC_Q - AIC_1$

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Simulation results ($Q^* = 2$, N = cn, $m^0 = [10, 800]$)



Model comparison for bat cries sequences

Poisson vs Hawkes / Homogeneous vs HMM. Best model based on BIC

	Poisson	Hawkes	Total
Homogeneous	132	775	907
Hidden Markov	21	627	648
Total	153	1402	1555

States and locations



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