# Inference for Epidemic Data using Diffusion processes with small diffusion coefficient

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# Outline



2 Parametric inference for discretely observed diffusion process

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8 Return to the epidemics and simulations results

#### Plan



2 Parametric inference for discretely observed diffusion process

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# Notations, and model assumptions

#### Notations

- N : population size
- m : initial invectives
- $\lambda$  : transmission rate
- $\gamma$  : recovery rate

 $R_0$ : basic reproduction number

S(t), l(t): numbers of susceptibles, infecteds,  $s(t) = \frac{S(t)}{N}, i(t) = \frac{l(t)}{N}$ : proportion of susceptibles, infecteds

# Assumptions

- Homogenous mixing in closed population
- Discrete observations of S and I on a fixed interval [0, T], with sampling interval Δ (T = nΔ)

$$\begin{array}{c} S \xrightarrow{\lambda I/N} I \xrightarrow{\gamma} R \end{array}$$

# Markov Pure Jump Model

Let 
$$X_0 = (N - m, m)$$
 and  $X_t = (S_t, I_t)$ .

Transitions and holding time

$$\begin{array}{c} (S,l) \xrightarrow{\frac{\lambda}{N}Sl} (S-1,l+1) \\ (S,l) \xrightarrow{\gamma l} (S,l-1) \\ \text{Exponentials holding times} \end{array}$$

Maximum Likelihood Estimators from complete observations (all jumps)

$$\hat{\lambda}_{MLE} = N \frac{N - m - S(T)}{\int_0^T S(t) I(t) dt}, \ \hat{\gamma}_{MLE} = \frac{N - S(T) - I(T)}{\int_0^T I(t) dt}$$

### Asymptotic Normality

$$\begin{split} \sqrt{N} \left( \begin{pmatrix} \hat{\lambda}_{MLE} - \lambda_0 \\ \hat{\gamma}_{MLE} - \gamma_0 \end{pmatrix} \right) & \xrightarrow[N \to \infty]{} \mathcal{N} \left( 0, \begin{pmatrix} var(\lambda_0) & 0 \\ 0 & var(\gamma_0) \end{pmatrix} \right) \\ \text{with } var(\lambda_0) &= \frac{\lambda_0^2}{(1 - \frac{m}{N})(1 - s(T))}, var(\gamma_0) = \frac{\gamma_0^2}{(1 - \frac{m}{N})(1 - \frac{m}{N} - s(T) - i(t))} \end{split}$$

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# ODE Model

Let 
$$x_{\lambda,\gamma}(t) = (s(t), i(t)), \ (s(0), i(0)) = (1 - \frac{m}{N}, \frac{m}{N})$$

Classical ODE System
$\frac{ds}{dt} = -\lambda si$
$rac{di}{dt} = \lambda si - \gamma i$ Do not depend on the population size !

#### Observations

Discrete observations at times  $t_k = k\Delta$ ,  $k = 0, ..., n X_{t_k} = x_{\lambda,\gamma}(t_k) + \epsilon_k$  with  $\epsilon_k \underset{iid}{\sim} \mathcal{N}_2 \left( 0, \begin{pmatrix} \sigma_1^2 & 0 \\ 0 & \sigma_2^2 \end{pmatrix} \right)$ 

# Statistical Inference for ODE

#### Least Square Estimator

$$LSE(\lambda,\gamma) = \sum_{k=0}^{n} (X_{t_{k}} - x_{\lambda,\gamma}(t_{k}))^{2}, (\hat{\lambda}_{LSE}, \hat{\gamma}_{LSE}) = \underset{(\lambda,\gamma)\in\Theta}{\operatorname{argminLSE}} (\lambda,\gamma)$$

#### Asymptotic Normality

$$\sqrt{n} \left( \begin{pmatrix} \hat{\lambda}_{LSE} - \lambda_0 \\ \hat{\gamma}_{LSE} - \gamma_0 \end{pmatrix} \right) \xrightarrow[n \to \infty]{} \mathcal{N} \left( 0, \begin{pmatrix} \sigma_1^2 & 0 \\ 0 & \sigma_2^2 \end{pmatrix} \right)$$

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### Diffusion approximation model

Let  $X_t = (s_t, i_t)$ ,  $B_1, B_2$  two independent Brownians motions,  $(s(0), i(0)) = (1 - \frac{m}{N}, \frac{m}{N})$ 

#### Stochastic Differential Equation

$$ds_t = -\lambda s_t i_t dt + \frac{1}{\sqrt{N}} \sqrt{\lambda s_t i_t} dB_1(t)$$

$$di_t = (\lambda s_t i_t - \gamma i_t) dt - \frac{1}{\sqrt{N}} \sqrt{\lambda s_t i_t} dB_1(t) + \frac{1}{\sqrt{N}} \sqrt{\gamma i_t} dB_2(t)$$

#### Remarks

- Classic Approximation : studies asymptotic properties of Pure Jump process (Ethier and Kurz) or Van Kampen approximation of Master Equation
- MLE untractable when discretely observed
- Multidimensionnal diffusion processes
- Small noise  $\sim \frac{1}{\sqrt{N}}$  in large population
- Parameters  $(\lambda,\gamma)$  both in drift and diffusion coefficient

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#### Classicals SIR epidemics model and diffusion approximation

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# Exemple of trajectory : proportion of infecteds over time



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### Plan



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## Theoretical model and existing results

#### Let $X_t^{\epsilon}$ be the unique strong solution of the SDE

• 
$$dX_t^{\epsilon} = b(\alpha, X_t^{\epsilon})dt + \epsilon \sigma(\beta, X_t^{\epsilon})dB_t, \ X_0 = x_0 \in \mathbb{R}^p$$

- We observe  $X_t^\epsilon$  at times  $t_k = k\Delta$  on a fixed interval [0, T]  $(T = n\Delta)$
- $\sigma(\beta, x) \in M_{P}(\mathbb{R}), b(\alpha, x) \in \mathbb{R}^{p}, \Sigma(\beta, x) = {}^{t}\sigma(\beta, x)\sigma(\beta, x) \in GL_{P}(\mathbb{R})$

#### Existing estimation result for high-frequency data (Gloter and Sorensen (2009))

Under the condition  $\exists \rho > 0, \frac{1}{\epsilon n^{\rho}}$  bounded For a class of contrast processes, associated Minimum Contrast Estimators (MCEs) are consistent and :

$$\begin{pmatrix} \epsilon^{-1}(\hat{\alpha}_{\epsilon,n} - \alpha_0) \\ \sqrt{n}(\hat{\beta}_{\epsilon,n} - \beta_0) \end{pmatrix} \xrightarrow[n \to \infty, \epsilon \to 0]{} N \begin{pmatrix} 0, \begin{pmatrix} I_b^{-1}(\alpha_0, \beta_0) & 0 \\ 0 & I_{\sigma}^{-1}(\alpha_0, \beta_0) \end{pmatrix} \end{pmatrix}$$

 $I_{b}^{-1}(lpha_{0},eta_{0})$  being optimal

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# Main Idea of our inference approach (Generalization of Genon-Catalot(90))

Use of Taylor's Stochastic Expansion formula (Azencott (82))

$$X_t^{\epsilon} = x_{lpha}(t) + \epsilon g_{lpha,eta}(t) + \epsilon^2 R_{lpha,eta}^{\epsilon}(t)$$

where  $x_{\alpha}(t)$  is the deterministic solution  $\frac{dx_{\alpha}(t)}{dt} = b(\alpha, x_{\alpha}(t)), \ x(0) = x_0 \in \mathbb{R}^p$ 

$$dg_{\alpha,\beta}(t) = \frac{\partial b}{\partial x}(\alpha, x_{\alpha}(t))g_{\alpha,\beta}(t)dt + \sigma(\beta, x_{\alpha}(t))dB_{t}, \ g_{\alpha,\beta}(0) = 0_{\mathbb{R}^{p}}$$

where  $R^{\epsilon}_{lpha,eta}$  satisfies :

$$\sup_{\epsilon \in [0,T]} \{ \| \epsilon R^{\epsilon}_{\alpha,\beta}(t) \| \} \underset{\mathbb{P}, \epsilon \to \mathbf{0}}{\longrightarrow} 0$$

Let  $\Phi_{\alpha}$  be the invertible matrix solution of  $\frac{d\Phi_{\alpha}}{dt}(t,t_0) = \frac{\partial b}{\partial x}(\alpha, x_{\alpha}(t))\Phi_{\alpha}(t,t_0), \ \Phi_{\alpha}(t_0,t_0) = I_p$ 

## Properties of $g_{lpha,eta}$

•  $g_{lpha,eta}$  is a gaussian process (and we can obtain is analytic expression)

• 
$$g_{\alpha,\beta}(t_k) = \Phi_{\alpha}(t_k, t_{k-1})g_{\alpha,\beta}(t_{k-1}) + Z_k^{\alpha,\beta}$$

•  $Z_k^{\alpha,\beta}$  independent gaussian variables

# Contrast process derived from $Z_k^{lpha,eta}$

$$U_{\Delta,\epsilon}(\alpha,\beta)) = \sum_{k=1}^{n} \log \left[ \det \left( \Sigma(\beta, X_{t_{k-1}}) \right) \right] \\ + \frac{1}{\epsilon^2 \Delta} \sum_{k=1}^{n} {}^t N_k(\alpha) \Sigma^{-1}(\beta, X_{t_{k-1}}) N_k(\alpha)$$
  
with  $N_k(\alpha) = X_{t_k} - x_\alpha(t_k) - \Phi_\alpha(t_k, t_{k-1}) \left[ X_{t_{k-1}} - x_\alpha(t_{k-1}) \right].$   
 $(\hat{\alpha}_{\epsilon,\Delta}, \hat{\beta}_{\epsilon,\Delta}) = \underset{(\alpha,\beta) \in \Theta}{\operatorname{argmin}} U_{\Delta,\epsilon}(\alpha,\beta)$ 

## Results for high frequency data $(\Delta ightarrow 0)$

Under the condition  $\epsilon^2 n \xrightarrow[\epsilon, \Delta \to 0]{} 0$ 

$$\begin{pmatrix} \epsilon^{-1}(\alpha_{\hat{\epsilon},\Delta} - \alpha_{0}) \\ \sqrt{n}(\hat{\beta_{\epsilon},\Delta} - \beta_{0}) \end{pmatrix} \xrightarrow[n \to \infty, \epsilon \to 0]{} N \left( 0, \begin{pmatrix} I_{b}^{-1}(\alpha_{0}, \beta_{0}) & 0 \\ 0 & I_{\sigma}^{-1}(\alpha_{0}, \beta_{0}) \end{pmatrix} \right)$$

# Results for low frequency data ( $\Delta$ and *n* being fixed)

# n fixed : no asymptotic results for $\hat{eta}_{\epsilon, \Delta}$

#### $\beta$ known

We only consider 
$$\hat{\alpha}_{\epsilon,\Delta}(\beta_0) = \underset{\alpha \in \Theta_a}{\operatorname{argmin}} U_{\Delta,\epsilon}(\alpha, \beta_0)$$
  
and then  $\epsilon^{-1}(\hat{\alpha}_{\epsilon,\Delta}(\beta_0) - \alpha_0) \xrightarrow[\epsilon \to 0]{} \mathcal{N}(0, I_{\Delta}^{-1}(\alpha_0, \beta_0))$   
with  $I_{\Delta}(\alpha_0, \beta_0) \xrightarrow[\Delta \to 0]{} I_b(\alpha_0, \beta_0)$ 

#### $\beta$ unknown

We modify the contrast process in a "conditional least square" contrast :

$$U_{\epsilon}\left(\alpha, (X_{t_k})_{k \in \{1, \dots, n\}}\right) = \frac{1}{\epsilon^2} \sum_{k=1}^{n} {}^{t} N_k(X, \alpha) N_k(X, \alpha)$$
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then 
$$\hat{\alpha}_{\epsilon} = \underset{\alpha \in \Theta_{a}}{\operatorname{argmin}} U_{\epsilon}(\alpha)$$
  
satisfies  $:\epsilon^{-1}(\hat{\alpha}_{\epsilon} - \alpha_{0}) \xrightarrow[\epsilon \to 0]{} \mathcal{N}(0, \tilde{l}_{\Delta}^{-1}(\alpha_{0}, \beta_{0}))$ 

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# Return on the diffusion model

#### Stochastic Differential Equation

$$\begin{aligned} ds_t &= -\lambda s_t i_t dt + \frac{1}{\sqrt{N}} \sqrt{\lambda s_t i_t} dB_1(t), \\ di_t &= (\lambda s_t i_t - \gamma i_t) dt - \frac{1}{\sqrt{N}} \sqrt{\lambda s_t i_t} dB_1(t) + \frac{1}{\sqrt{N}} \sqrt{\gamma i_t} dB_2(t) \end{aligned}$$

#### Simulations

 $\textit{N} \in [1000; 10000], \ \Delta = 1 \ (1 \ {\sf observation}/ \ {\sf day})$ 

• 
$$\epsilon = \frac{1}{\sqrt{N}} << 1$$

- Δ is fixed
- α = (λ, γ) = β ⇒ Special case : Results for known β hold if we replace each β occurence with α.

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# Simulation study (using Matlab)

#### Algorithm

- Exact simulation of an epidemic with Markov Pure Jump process (Gillespie algorithm with choice of  $N, m, \lambda, \gamma$ )
- <sup>(2)</sup> Calculation of  $\hat{\lambda}_{\textit{MLE}}, \hat{\gamma}_{\textit{MLE}}$  (observation of the whole path of the process)
- Observations of discrete data on a fixed interval (1 observation/day) up to extinction time
- Estimation phase for LSE, Gloter and Sorensen contrast, our method for unknown β (Conditionnaly least square contrast), and for α = β, using minimization function of Matlab (fminsearch)

#### Presented results

We repeat 100 times this algorithm to build empiric confidence intervals and avoid early extinction events

#### Remark

Step 4 : (Analytic power) Short time of estimation

# Simulation results $(R_0 = 2)$

For N = 1000, m = 10,  $\lambda$  = 2/3,  $\gamma$  = 1/3, 1 data/day  $\Rightarrow$  40 observations

Method	$\hat{\lambda}$	Cl <sub>95</sub> empiric	Cl <sub>95</sub> theoretical
MLE(all data)	0.657	[0.645; 0.669]	[0.643; 0.671]
LSE	0.643	[0.618; 0.668]	[0.633;0.653]
Gloter Sorensen $(\hat{lpha}_{\epsilon, n})$	0.622	[0.611; 0.634]	[0.622; 0.622]
$\hat{\alpha}_{\epsilon,\Delta}(\alpha=\beta)$	0.656	[0.651; 0.660]	[0.656; 0.657]
$\hat{lpha}_{\epsilon}(eta$ unknown)	0.645	[0.642; 0.649]	[0.644; 0.646]
Method	$\hat{\gamma}$	Cl <sub>95</sub> empiric	Cl <sub>95</sub> theoretical
MLE(all data)	0.336	[0.330; 0.342]	[0.330; 0.342]
LSE	0.329	[0.314; 0.343]	[0.321;0.337]
Gloter Sorensen $(\hat{lpha}_{\epsilon, n})$	0.386	[0.367; 0.404]	[0.386;0.386]
$\hat{\alpha}_{\epsilon,\Delta}(\alpha=\beta)$	0.336	[0.333; 0.338]	[0.335;0.336]
$\hat{\alpha}$ (B unknown)	0 221	[0 330 0 333]	[0 330.0 331]

#### Global remarks

- $\hat{\beta}_{\epsilon,n}$  and  $\hat{\beta}_{\epsilon,\Delta}$  do not provide satisfying results (not shown)
- Red : True value of parameters not in the CI
- Green : best point estimation

# Simulations results $(R_0 = 1.2)$

For N = 10000, m = 100,  $\lambda = 0.4$ ,  $\gamma = 1/3$ , 1 observation/day  $\Rightarrow$  115 observations

Method	$\hat{\lambda}$	empiric Cl <sub>95</sub>	$\hat{\gamma}$	empiric Cl <sub>95</sub>
MLE(all data)	0.397	[0.395; 0.399]	0.337	[0.336; 0.338]
LSE	0.387	[0.377;0.398]	0.328	[0.319; 0.337]
Gloter Sorensen $(\hat{\alpha}_{\epsilon,n})$	0.410	[0.409; 0.411]	0.330	[0.330; 0.331]
$\hat{\alpha}_{\epsilon,\Delta}(\alpha=\beta)$	0.396	[0.396; 0.397]	0.329	[0.329; 0.330]
$\hat{lpha}_{\epsilon}(eta$ unknown)	0.396	[0.396; 0.397]	0.336	[0.336; 0.337]

For N = 1000, m = 10,  $\lambda = 0.4$ ,  $\gamma = 1/3$ , 1 observation/day  $\Rightarrow$  65 observations

Method	$\hat{\lambda}$	empiric Cl <sub>95</sub>	$\hat{\gamma}$	empiric Cl <sub>95</sub>
MLE(all data)	0.387	[0.381; 0.393]	0.362	[0.346; 0.377]
LSE	0.402	[0.364; 0.440]	0.353	[0.322; 0.384]
Gloter Sorensen $(\hat{lpha}_{\epsilon,n})$	0.382	[0.381;0.383]	0.336	[0.334; 0.338]
$\hat{\alpha}_{\epsilon,\Delta}(\alpha=\beta)$	0.396	[0.394; 0.397]	0.357	[0.355; 0.359]
$\hat{lpha}_{\epsilon}(eta$ unknown)	0.392	[0.385; 0.399]	0.363	[0.359; 0.367]

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# Exemple of trajectory for $R_0 = 1.2$ and N=1000



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## Limits and perspectives

#### Limits

- Limits of the SIR model
- The total number of infecteds is assumed observed (instead of incidences, more realistic assumption)
- The two coordinates (s<sub>t</sub>, i<sub>t</sub>) are assumed observed (which is not often the case)

#### Next Directions

- Results hold for any autonomous system (SEIR,...)
- **2** Modifying the diffusion model (observe  $(u_t, v_t)$  with  $u_t = s_t i_t, v_t = i_t$ ) and observe integrated diffusion

Work to do

Thank you!

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