Time series analysis via maximum likelihood From theory to practice

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Year





Year



Parameter inference:

Identifiability, maximum likelihood estimates, confidence intervals

Model selection:

objective ranking of models, which hypothesis best explains the data?

Tristan da Cunha (1971) a two-wave flu epidemic



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A simple mechanistic approach



Long-term

immunity

λ = β I/N mass-action
I/ε : mean latent period
I/ν : mean infectious period
I/γ : mean removed period

HI: the virus mutated during the first epidemic-wave (Mut)



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- $\sigma \in [0, 1]$ cross-immunity
- 2-strain history-based model (Rios-Doria & Chowell 2009)

H2: intra-host recrudescence of infection (InH)



α : the probability to clear the viral load



I/T: the mean duration of the window of susceptibility before developing immunity

Likelihood-based inference

For a given time series: $y_{1:T} = (y_1, y_2, ..., y_T)$ and a state space model completely specified by:

$$M: \begin{cases} f(x_t|x_{t-1},\theta) & \text{the conditional transition density} \\ f(y_t|x_t,\theta) & \text{the conditional distribution} \\ & \text{of the observation process} \\ f(x_0|\theta) & \text{the initial density} \end{cases}$$

the likelihood is given by the identity:

$$f(y_{1:T}|\theta) = \prod_{t=1}^{T} f(y_t|y_{1:t-1},\theta)$$

where x_t is the unobserved Markov process, θ is the unknown vector of parameters and f(.|.) is a generic density specified by its arguments

Exploring the likelihood surface



Exploring the likelihood surface with MIF (lonides et al. 2006)

 3 local maxima
 I global maximum

Param 1





Param 2

Exploring the likelihood surface with MIF (lonides et al. 2006)

 3 local maxima
 I global maximum

Param 1



Local trap:
initial θ
MIF parametrization



Log-likelihood profile



Parameter identifiability



Structural non-identifiability (Mutation: HI)



Structural non-identifiability between σ and $\beta_2 \Rightarrow \beta_2 = \beta_1$

Practical non-identifiability (In-Host: H2)



Likelihood-based inference

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the Log-likelihood is given by the identity:

$$\log(f(y_{1:T}|\theta)) = \sum_{t=1}^{T} \log(f(y_t|y_{1:t-1},\theta))$$

where x_t is the unobserved Markov process, θ is the unknown vector of parameters and f(.|.) is a generic density specified by its arguments

$$AIC_c = -2\mathcal{L}(\theta_{MLE}) + 2k + \frac{2k(k+1)}{T-k-1} \text{ with } k = ||\theta||$$

Model	Win	Mut	In-Host
k	9	10	9
Log-Like	-112.52	-115.20	-117.50
ΔAIC _c	0	8.27	9.96



Dynamics comparison



Conclusion

- Maximum likelihood via Iterated Filtering (MIF, Ionides et al. 2006) is a rigorous statistical framework for parameter inference and selection based on AIC for mechanistic stochastic models.
- Identifiability analysis and 95% CI via log-likelihood profile
- Illustrate and compare the dynamics of the different models in a visual and intuitive manner
- (I) Ionides EL, Breto CM, King AA. Inference for nonlinear dynamical systems. *PNAS*. 2006
- (2) Raue A, Kreutz C, Maiwald T, et al. Structural and practical identifiability analysis of partially observed dynamical models by exploiting the profile likelihood. *Bioinformatics (Oxford, England)*. 2009
- (3) Camacho A, Ballesteros S, Graham AL, Carrat F, Cazelles B. Explaining rapid reinfection in multiplewave influenza outbreaks? Tristan da Cunha epidemic as a case study. (*in preparation*)

Likelihood-based inference

For a given time series: $y_{1:T} = (y_1, y_2, ..., y_T)$ and a state space model completely specified by:

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Global estimate:
$$\hat{\theta}^{(n)} = \hat{\theta}^{(n-1)} + V_1^{(n)} \sum_{t=1}^T \frac{\hat{\theta}_t^{(n)} - \hat{\theta}_{t-1}^{(n)}}{V_t^{(n)}}$$

As shown by Ionides *et al.* (2006), under rather mild assumptions,

$$\lim_{\sigma \to 0} \sum_{t=1}^{T} \frac{\hat{\theta}_t - \hat{\theta}_{t-1}}{V_t} = \nabla \log f(y_{1:T} | \theta, \sigma = 0)$$

so that, for a sufficiently small σ_n , the algorithm iteratively updates $\hat{\theta}^{(n)}$ in the direction of increasing likelihood, with a fixed point at a local maximum of the likelihood surface.