### Estimating variability in nonlinear & stochastic epidemiological models

An application to the transition from invasion to long-term persistence

Anton Camacho, Sébastien Ballesteros, Amaury Lambert & Bernard Cazelles

Department of Biology UMR 7625, UPMC-CNRS-ENS <u>camacho@biologie.ens.fr</u>

## A paradigmatic model



Key quantities:  $R_0 = \beta/\nu$  is the fitness of the pathogen  $I/\gamma$  is the mean duration of immunity

## A paradigmatic model



Basic assumptions: Closed population Homogeneous mixing Exponential durations Improve realism: Latent period, seasonal forcing, gamma durations, heterogeneous mixing etc.



Time (day)



Time (day)

### Stochastic extinctions (1/3) Failed invasion



### Stochastic extinction (2/3) Epidemic fade-out



Time (day)

### Stochastic extinction (3/3) Endemic fade-out



# Quantitative characterization of the dynamics (1/2): Monte-Carlo



# Quantitative characterization of the dynamics (1/2): Monte-Carlo





#### Quantitative characterization of the dynamics (2/2): Asymptotic approximations





# Qualitative characterization of the dynamics: basic idea

400

300

200

We use an ODE-based (fast) and automatized (wide application) method to characterize the stochastic dynamics from invasion to long-term persistence



Master equation

$$S + I \xrightarrow{\beta \frac{SI}{\Omega}} 2I \qquad I \xrightarrow{\nu I} R \qquad R \xrightarrow{\gamma R} S$$

 $\frac{d}{dt}P(S, I, R, t) = \beta(S+1)\frac{I-1}{\Omega}P(S+1, I-1, R, t)$ 

Cannot be solved analytically due to nonlinearity of the contact process

$$+\nu(I+1)P(S,I+1,R-1,t)$$
$$+\gamma(R+1)P(S-1,I,R+1,t)$$
$$-\left(\beta S\frac{I}{\Omega}+\nu I+\gamma R\right)P(S,I,R,t)$$

### **Notations** $S + I \xrightarrow{\beta \frac{SI}{\Omega}} 2I$ $I \xrightarrow{\nu I} R$ $R \xrightarrow{\gamma R} S$

Species vector (size: N)  

$$\mathbf{X} := \begin{bmatrix} S \\ I \\ R \end{bmatrix} = \Omega \begin{bmatrix} S/\Omega \\ I/\Omega \\ R/\Omega \end{bmatrix} = \Omega \begin{bmatrix} s \\ i \\ r \end{bmatrix} = \Omega \mathbf{x}$$

Propensity function vector (size: M)

$$\begin{bmatrix} \beta \frac{SI}{\Omega} \\ \nu I \\ \gamma R \end{bmatrix} = \Omega \begin{bmatrix} \beta si \\ \nu i \\ \gamma r \end{bmatrix} = \Omega \mathbf{f}(\mathbf{x}$$

Stoichiometric matrix (size: NxM)

$$\mathbb{S} := \begin{bmatrix} -1 & 0 & +1 \\ +1 & -1 & 0 \\ 0 & +1 & -1 \end{bmatrix}$$

### Linear Noise Approximation



Following a Taylor expansion of the ME in powers of  $I/\sqrt{\Omega}$ , we find at the next to leading order:

$$\frac{d}{dt}\phi = \mathbb{S}\mathbf{f}(\phi) \qquad \xi \hookrightarrow \mathcal{N}(0,\Xi) \qquad \frac{d}{dt}\Xi = \mathbb{A}\Xi + \Xi \mathbb{A}^T + \mathbb{B}$$

with:  $\Xi = \langle \xi \xi^T \rangle \quad \mathbb{A} = \frac{d}{d\phi} \mathbb{S}\mathbf{f}(\phi) \quad \mathbb{B} = \mathbb{S} \ diag(\mathbf{f}(\phi)) \ \mathbb{S}^T$ 

#### «How close are we to 0?»

Since  $I_t = \Omega \phi_2(t) + \sqrt{\Omega} \xi_2$  we have:

$$I_t \hookrightarrow \mathcal{N}\Big(\Omega\phi_2(t), \Omega\Xi_{22}(t)\Big)$$

We define:  $I_t^* = \Omega \phi_2(t) - \mathcal{Q}_p \sqrt{\Omega \Xi_{22}(t)}$  as the lower bound of the *p*-quantile of the distribution. In practice we choose p = 99% and thus  $\mathcal{Q}_{99\%} \approx 2.6$ 

We neglect the extinction probability whenever  $I_t^* > 0$  and simply define the critical population size at time t,  $\Omega_t^*$ , such that  $I_t^* = 0$ 

$$I_t^* = 0 \Leftrightarrow \Omega_t^* = \mathcal{Q}_p^2 \frac{\Xi_{22}(t)}{\phi_2^2(t)}$$

### «How close are we to 0?»



# From invasion to persistence

3 different qualitative outcomes





Duration of immunity



# Improve realism: gamma distributions







# Summary & Outlook

- Linear Noise Approximation: Fast & Automatized for complex models
- Evaluate stochastic fluctuations around the deterministic trajectory (stability analysis)
- Which regions of the parameter space (do not) require stochastic simulations?
- Qualitative & quantitative information on the stochastic dynamics

# Summary & Outlook

- Improving realism: seasonal forcing (but no equilibrium), immigration (how to define extinction), more than 2 R classes (Hopf bifurcation of the endemic equilibrium)
- The linear noise approximation can also be used to derive the power spectral density of the endemic fluctuations